

APPENDIX A

DATA MANAGEMENT SUMMARY REPORT (ECO-CHEM, 2000)

Data Management Summary Report

Fox River Remedial Investigation/Feasibility Study

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Prepared for:

**Wisconsin Department of Natural Resources
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Madison, Wisconsin 53707**

October 3, 2000

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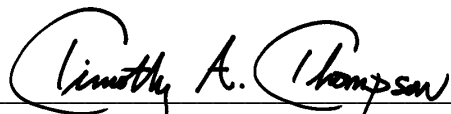
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1 Introduction

This report summarizes the processes and data utilized to create the Fox River Database (FRDB). The FRDB was created to provide data management support to the Lower Fox River Remedial Investigation/Feasibility Study (RI/FS) and Risk Assessment (RA). The data management and data quality assessment have been conducted with two primary goals in mind:

- The identification and incorporation of available electronic data sets for immediate use in the support of RA and RI/FS activities and the assessment of these data sets for overall quality and defensibility.
- The generation of a useable database of Fox River data produced through the identification, acquisition, review (quality assessment or validation), catalog, classification, and archive of all available data (electronic and hardcopy) pertinent to the Fox River RA and RI/FS.

Environmental data generated by numerous sources in support of several different actions on the Fox River were collected and assessed for overall quality and included in the FRDB.

For the purposes of this document the following definitions will apply:

- **Data Set** - an electronic set of data that is associated with or is identified by a unique study name or sampling event. Identified data sets were submitted in many different formats (e.g., spreadsheets, databases, ASCII files, etc.).
- **Sample** – a unique, representative fraction of a matrix of interest (sediment, fish tissue, water, etc.) collected during a discrete time period.
- **Record** – collection of all data associated with a single analytical result in the FRDB (location, qualifiers, comments, etc.).
- **Data Validation (DV)** - data validation is the process of independent data review which provides information pertaining to limitations of data based on specific quality control criteria.

- **Useable Data** - useable data have been thoroughly assessed through review of the analytical data itself and associated quality assurance/quality control (QA/QC) documents. The data are of known and verifiable quality. Useable data is identified as such in the “qa_status” field in the FRDB.
- **Supporting Data** - supporting data have not been subjected to as rigorous an assessment as the useable data. As such, the precise data quality is not known. This is due to insufficient or incomplete QA/QC information available at the present time. In these cases, QA/QC information may or may not exist. The collection and assessment of this information might render the data fully useable. Until a full data validation is conducted, these data should be used for supporting purposes only. Supporting data is identified as such in the “qa_status” field in the FRDB.
- **Indeterminate Data** – status of a data set is described as indeterminate if: it is unknown whether the data set has been validated, and/or, QC data to support validation is not available. Indeterminate data is identified as such in the “qa_status” field in the FRDB.

2 Data Collection

2.1 Electronic Data Collection

The data management process began with the initial collection of electronic data sets from the Wisconsin Department of Natural Resources (WDNR) the week of March 30, 1998. The data collection effort proceeded in two stages, corresponding with the report delivery schedule developed for the RI/FS and RA documents. Data collection for Stage 1 continued through November 30, 1998, and all data were available to support the Draft RI/FS and RA documents published in February of 1999. Stage 2 of data collection began in March of 1999, and continues through the present (May 2000). Data were received in many different formats and were reviewed, standardized, and organized into a database-compatible format. The following table lists the data received and the stage that it was collected.

| Stage 1 | Data Source | Stage 2 |
|---|-------------|--|
| 1989–1990 Fox River Mass Balance Study | | 1997 Demonstration Project Data - Deposit N |
| 1989–1990 Green Bay Mass Balance Study (GLNPO) | | 1997–1998 Demonstration Project Data - SMU 56/57 |
| 1992–1993 BBL Deposit A Sediment Data | | 1998 FRG/BBL Sediment/Tissue Data |
| 1993 Triad Assessment | | 1998–1999 Deposit N Data: |
| 1993 USFWS Tree Swallow Data | | Remediation |
| 1994 GAS/SAIC Sediment Data | | Pre-Dredge |
| 1994 Woodward-Clyde Deposit A Sediment Data | | Post-Dredge |
| 1994–1995 Cormorant Data | | Operational Monitoring |
| 1995 WDNR Sediment Data | | 1998 FRG/Exponent Data |
| 1996 FRG/BBL Sediment/Tissue Data | | 1999 Demonstration Project Data - SMU 56/57 |
| 1995–1996 WDNR Fish Tissue Data | | Ankley and Call Data |
| 1997 USFWS NRDA Waterfowl Tissue Data | | State of Michigan Fish Consumption Advisory Data |
| 1997 WDNR Caged Fish Bioaccumulation Study Data | | Lake Michigan Mass Balance Data |
| 1998 RETEC RI/FS Supplemental Data | | Lake Michigan Tributary Monitoring Data |
| Fox River Fish Consumption Advisory Data | | Minergy Mineralogical Data |
| Lower Fox River Background Metals Assessment | | |
| Stromberg Eagle Data | | |
| 1996–1999 USFWS NRDA Fish Tissue Data | | |
| USGS NAWQA Data | | |
| WDNR Wildlife Tissue Data | | |
| WPDES Permit Influent Data | | |

2.2 Collection of Historical Analytical Data and Supporting Quality Assurance and Quality Control Documents

The goal of the review was to assess previously generated analytical data sets and associated Sampling and Analysis Plans (SAPs), Quality Assurance Project Plans (QAPPs), Laboratory Standard Operating Procedures (SOPs), and other project-specific documents. Historical data (both hardcopy and electronic) and supporting QA documents were collected for review and verification.

3 Data Manipulation and Assessment

3.1 Data Management and Data Validation Overview

Most of the data sets required a substantial amount of manipulation to transform the structure to a common database format. The data were usually obtained from report documents that had undergone extensive formatting. This formatting had to be removed to restore the data set to its most basic state and transform individual data sets into a useable condition.

The formats in which data were received are included in Table 3-1. A brief description of how the data were adapted is provided below.

- **Spreadsheet:** Numerous data tables were provided in spreadsheets, but not necessarily in a database-compatible format. It was often necessary to manually rearrange data within the spreadsheet from a horizontally oriented format (multiple results on a single line) to a vertical format (one individual result per record). Spreadsheet columns were then rearranged into the proper record order as necessary and the file appended to the FRDB.
- **ASCII:** Data were imported into a spreadsheet or database table. The table was then checked to verify that the information was separated into individual fields properly. Information was then rearranged into the proper record order as necessary and the file appended to the FRDB.
- **Database:** Data were provided in multiple database formats. When necessary, the data were exported to FoxPro tables. Field headers were then standardized to match the established database format and the file appended to the FRDB.
- **Hardcopy:** Information was provided in a written report with data tables (one data set only). Information was gathered from the tables provided and the supporting text. The data were hand entered into an empty spreadsheet table with the same record setup as the database. All hand-entered information was proofread by a second party to insure accuracy prior to inclusion in the FRDB.

In addition to reducing the data to a database useable format, the disparate data sets required standardization. This process consists of developing master lists of acceptable entries for pertinent data types (valid value lists) and verifying that all new data sets conform to those master lists. The following items offer examples of the standardization that took place:

- A single analyte list was developed in order to account for different naming conventions reported by multiple laboratories. A cross-reference table was used to update each data set to a standardized list of analytes. For example, all instances of 4,4'-DDT were changed to p,p'-DDT and all PCB congener results were put into the format "PCB Congener XXX." The original analyte name as received in the import file is maintained in the "analyte_old" field of the FRDB.
- Units were standardized to parts per million (mg/L or mg/kg) for inorganic constituents and parts per billion (mg/L or mg/kg) for organic analytes. Two different possibilities exist for unit changes: unit changes that do not require numeric calculations, e.g., ng/ml to mg/L (both represent parts per billion units) and units changes that require numeric calculations e.g., 10 mg/kg changed to 0.01 mg/kg. All original values and units were concatenated and placed in the "result_old" field of the FRDB.
- Qualifiers were standardized to the extent possible. For the most part, this consisted of changing "<" signs to "U," and interpreting laboratory-assigned qualifiers. Where this information is unavailable or has yet to be obtained, original qualifiers have been maintained. In those data sets where multiple qualifiers are available (laboratory qualifiers and validation qualifier), the multiple qualifiers have been merged to a single qualifier (i.e., "U" qualified from laboratory and "UJ" qualified by the validator = "UJ" qualified). When non-standard qualifiers were present in data received, the data provider was contacted and a list of qualifiers and definitions was requested. Qualifiers were standardized accordingly. The original qualifiers received in the import file are maintained in the "qual_old" field of the FRDB.
- All sample dates were standardized to one common data format where possible: mm/dd/yyyy.

- The media field was populated using a standard list of sample matrices: ambient air, pore water-sediment, sediment, tissue, or water.
- The species (common name) was standardized. For example, Northern Pike was also listed as N. Pike, northern pike, and Northern pike. The most accurate descriptor was chosen and all permutations were changed to match.
- The sample type (whole body, surface sediment, fillet skin-on) was standardized.
- Sample depth was standardized to measurement in centimeters. For some sediment samples, the sampling depth was included in the sample identification. This information was moved to the “depthfrom” and “depthto” fields in the database. Units of measurement were placed in the “depthunits” field.

Beyond the standardization process, information was added to delivered data sets in order to provide unique information where required, and to enable grouping of information (by location, analysis type, etc.) in support of the RI/FS or RA.

- Unique sample identifiers (IDs) were generated for samples that did not have a single unique identifier. Tissue samples generated by different researchers often had identical sample IDs. In these cases, a letter in parenthesis was appended to the original sample ID to indicate the researcher [(P) - Patnode data, (S) - Stromberg data, etc.]. In other cases, multiple researchers used an identical counting scheme to identify samples, based on the year and the numerical sample count (i.e., the first sample in 1995 was 95001, the second was 95002). In cases where more than one researcher collected samples in this manner, the samples were identified as 95001a, 95001b, and so forth. Water samples were often analyzed as filtered and unfiltered, or filtered and particulate. When such samples had similar sample IDs, a (U) – unfiltered, (F) – filtered, or (P) – particulate was appended to the sample ID making it unique.
- Individual samples from various data sets were assigned location information to allow for spatial association to other data sets. All samples were assigned one of the following nine designations:

background or reference; Little Lake Butte des Morts; Appleton to Little Rapids; Little Rapids to De Pere; De Pere to Green Bay; Green Bay Zone 2 (2A & 2B); Green Bay Zone 3A; Green Bay Zone 3B; or Green Bay Zone 4. Descriptive location information and coordinate information were used to successfully associate 99.9 percent of the samples with one of the above areas. Where possible, samples collected on the upper stretch of the river were also associated with the deposit from which they were sampled.

- The “northing” and “easting” fields contain specific coordinate information provided by the originator of the data or WDNR based on original site mapping.
- The “lab” and “validator” fields were populated if the information was available.
- The spelling, case, and date format (where applicable) were standardized for the fields titled “Source,” “Methodtype,” “Group,” “Group2,” “Importfile,” and “Timestamp.”
- The following fields were populated if the information was provided: “labid,” “date_recd,” “date_ext,” “detlimit,” “sdg,” “aliquot,” “method,” “blind_id,” “sampler,” “comment,” “loc_description,” and “county.” No standardization was applied to this information.

Tabular results of analysis for all data sets included in the FRDB are provided in Table 3-1.

The quality assessment of the historical data followed a stepwise approach. First, it was determined whether the data had been subjected to an independent third-party data validation. If the data were validated and the validation report or validation worksheets were available, they were reviewed. If the validation was determined to follow basic U.S. EPA quality assurance guidelines (at a minimum), the data were considered to be acceptable for use (useable) in the RI/FS and risk assessment decision-making process.

If the data were not validated or concurrence was not reached with the previous validation (and the QC results were available), a limited review was performed. All available documents were reviewed to determine what quality control measures were included and what data quality objectives (DQOs) were required. The measures of accuracy and precision were evaluated against either

the control limits/DQOs in the QAPP, the method, the laboratory SOPs, or U.S. Environmental Protection Agency (EPA) National Functional Guidelines. QC elements such as sample duplicates, matrix spike/matrix spike duplicates (MS/MSD), laboratory control sample/laboratory control sample duplicate (LCS/LCSD), and field duplicates were acceptable measures of precision. QC elements such as blanks, calibration standards (initial and continuing), surrogates, MS/MSD, LCS/LCSD, and standard reference materials (SRMs) were acceptable measures of accuracy. A determination of the usability of the data was made from the findings of these reviews. The analysis of the available QA/QC elements for each data set are summarized in Table 3-2.

3.2 Data Sets

The reduced and standardized data sets were compiled in a working database for use in support of the ongoing RA and RI/FS. This interim database is essentially a large flat file, currently containing more than 450,000 records from 35 individual data sets. Each data set is discussed in the following subsections of this report.

3.2.1 1989–1990 Fox River Mass Balance Study and 1989–1990 Green Bay Mass Balance Study (GLNPO)

The 1989–1990 Fox River Mass Balance data were collected by WDNR along the length of the river in 1989 and 1990. The sediment and water matrices of this data set were received from WDNR in six spreadsheet files (1989-1.wks, 1989-2.wks, allsed.wks, basic-5.wks, deep-cor.wks, and gravity.wks). These spreadsheets contain polychlorinated biphenyl (PCB) congener and total PCB concentrations, as well as grain size and total organic carbon (TOC) information. Each file exists in a unique format and was transformed into a standard database format. These data represent 1,967 samples and 25,457 analytical records in the FRDB. Data management occurred during Stage 1 of the data collection process.

The Green Bay Mass Balance (GBMB) data are represented in their entirety in the files posted on the Great Lakes National Program Office (GLNPO) website. Several mass balance studies have been conducted by different regulatory agencies and groups. Consequently, there is a significant overlap of data which is considered “common” data within the different studies. Redundant data identified in the collective GLNPO set were segregated and removed prior to inclusion of the GLNPO data into the FRDB (2,069 samples and 201,701 records). Data management occurred during Stage 1 of the data collection process except for the phyto- and zooplankton fractions of the data. These

data were originally omitted from the FRDB. During Stage 2 of the data collection and management process, these data were determined to be required for food chain models and were added to the FRDB.

Samples were analyzed and data were generated by eight different laboratories for the GBMB study. Seven of the laboratories performed PCB analyses; one laboratory performed metals analyses. Each of the seven laboratories analyzing samples for PCBs were required to analyze a series of 10 performance evaluation (PE) samples (of differing concentration levels) prior to analyzing samples for the study. The results of these PE sample analyses were available for review by EcoChem for four laboratories. A wide range of percent recovery (%R) values were reported (60% to 233%).

Prior to the study, each laboratory was given a copy of the document, *Quality Assurance Plan Green Bay Mass Balance Study - PCBs and Dieldrin*, which outlined general guidelines and data quality objectives. According to this document, data sets generated for the GBMB Study were reviewed and approved by the Green Bay Quality Assurance Coordinator (QAC) prior to the release of data. EcoChem, Inc. interviewed the GBMB QAC at the University of Minnesota in September 1998 regarding the data review procedures. It was determined from that meeting that the data were not fully validated. The review of the data consisted of verification of laboratory-generated QA/QC forms prior to data release. A formal comparison to any specific project DQOs was not made, thus no validation qualifiers were assigned to the data.

One participating laboratory, the Wisconsin State Laboratory of Hygiene (SLOH), was visited by EcoChem personnel who interviewed analysts and managers. Sample handling, preparation, and analysis systems were reviewed. In-depth discussions occurred concerning peak identification and quantitation. All hardcopy and electronic data are available and could be validated if requested. The disposition of the data and supporting information for the other labs is not known. Thus, it was determined that, in general, the data from the GBMB Study should be used as supporting data only. Refer to 2.2.18 for a discussion of the review of more recent data generated by SLOH.

3.2.2 1992–1993 BBL Deposit A Sediment Data

Sediment and water samples were collected in late 1992 and early 1993 by Blasland, Bouck and Lee (BBL) at Deposit A. The samples were analyzed for volatiles, semivolatiles, PCB Aroclor, pesticides, metals, and wet chemistry tests. Aroclor™ data was received during Stage 1 of the data management

process, the other analyses during Stage 2. These data represent 117 samples in the FRDB and accounts for 1,094 data records.

EcoChem, Inc. conducted a full data validation of these data in 1999 (Stage 2). The samples were analyzed by Hazleton Environmental Services, Inc. in Madison, Wisconsin. Analytical data were reviewed using quality control criteria documented in the analytical method, National Functional Guidelines, and the project QAPP. Validation was performed on volatile, semivolatile, PCB as Aroclor™, pesticide, and metals data. Accuracy and precision were generally acceptable. Qualifiers were assigned by EcoChem due to blank contamination, calibration outliers, secondary column confirmation precision outliers, laboratory control sample outliers, MS/MSD outliers, surrogate outliers, laboratory duplicate results, and graphite furnace post-digestion spike recovery results. Data, as qualified by EcoChem, are acceptable for use. The Data Validation Report is included as Appendix A of this report.

3.2.3 1993 Triad Assessment

The Triad data were collected by WDNR from several sites and analyzed in 1992 and 1993. The data were received from WDNR in 11 spreadsheet files (joint.wb2, orgpest.wb2, rtrben.wb2, tables.wb2, toxicity.wb2, triad92.wb2, triad92b.wb2, triad93.wb2, triaddat.wb2, triadhis.wb2, and foxriver.wq1) during Stage 1 data collection. All data were represented in files triad92b and triad93, and were redundant in the rest of the files. These spreadsheets contain polynuclear aromatic hydrocarbon (PAH), metals, Aroclor™, chlorinated pesticide, invertebrate, and benthos data. These data represent 27 samples and 631 analytical records in the FRDB. The original Triad data were modified to create unique sample IDs. A designation of “(Tr)” was appended to the existing sample IDs to ensure uniqueness. Data management occurred during Stage 1 of the data collection process.

Samples collected for the Triad Study were submitted to several different laboratories for physical and chemical characterization. These laboratories include University of Wisconsin-Extension’s Soil and Plant Analysis (particle size and soil texture analyses); the State Laboratory of Hygiene (bulk sediment chemistry); and Hazleton Laboratory (PAHs collected in 1993). Quality control data were not available for review; however, full data validation on SLOH data could be conducted if requested. As these data have not undergone full validation, these data should be used as supporting data only.

3.2.4 1994 GAS/SAIC Sediment Data

The Graef, Anhalt, Schloemer & Associates/Science Application International Corporation (GAS/SAIC) data were collected during late 1994 for the Fox River Coalition. This data set includes sediment data collected at several deposits above the De Pere dam. Samples were analyzed for PCB Aroclors™, chlorinated pesticides, volatile organics, semivolatile organics, metals, and dioxins. These data were delivered by WDNR in six files (clp_data.xls, cnv_data.xls, dxn_data.xls, hg_data.xls, pcb_data.xls, and frgrnsiz.xls). The GAS/SAIC data set consists of 253 samples that comprise 5,654 records in the FRDB. Data management occurred during Stage 1 of the data collection process.

Approximately 20 percent of the GAS/SAIC data was fully validated by SAIC. The remainder of the data underwent a cursory review that excluded verification of compound identifications and raw data calculation checks. This evaluation followed specified methods described in the November 1994 *Final Report Sampling and Analysis Plan, Fox River Remedial Investigation*. The data validation reports do not specifically address chain of custody records associated with the samples.

In the process of reviewing the initial PCB and pesticide data reported by the initial laboratory involved, SAIC found incorrect PCB quantitations, inconsistent pesticide identifications, consistently poor surrogate recoveries, retention time shifts, and overall poor quality of work associated with the pesticides/PCB data. Based on EcoChem's review, these data should be used as supporting data only.

PCB-only analyses (from archived samples) and dioxin analyses were performed later by Analytical Resources, Inc. and Triangle Laboratories. In general, precision and accuracy for these analyses were judged acceptable by SAIC. PCB results were qualified as estimated by SAIC due to continuing calibration verification percent difference exceedances and poor surrogate recoveries. The dioxin results received minor qualifications due to blank contamination and elevated matrix spike recovery values. These data, as qualified by SAIC, are acceptable for use.

3.2.5 1995 WDNR Sediment Data

The 1995 sediment collection was conducted by WDNR and consists of sediment data collected from below the De Pere dam. Samples were analyzed for PCB Aroclors™ and metals. These data were provided by WDNR in eight files (corelocs.xls, convdata.xls, 95sedata.xls, metals.xls, metals2.xls,

pcbdata.xls, pcbdata2.xls, and sumdata.xls). The data set consists of 488 samples comprising 6,433 records. Data management occurred during Stage 1 of the data collection process.

Data validation was conducted by the M. A. Kuehl Company on approximately 20 percent of the 1995 De Pere data. The data validation reports were reviewed by EcoChem. Based on this evaluation, it was determined that the laboratory followed the specified methods described in the September 1995 *Quality Assurance Project Plan for Assessment of PCBs in Sediment of the Lower Fox River from De Pere to Green Bay*. Chain of custody records were reviewed, and they indicated that samples were received in good condition. These data, as qualified by M. A. Kuehl, are acceptable for use.

3.2.6 1996 FRG/BBL Sediment/Tissue Data

The 1996 BBL data set consists of 25 sediment and fish tissue samples collected for the Fox River Group (FRG). These samples were analyzed for PCB congeners and TOC. These data were provided by WDNR in six spreadsheet files (02771543.wq1, 02671543.wq1, 02571543.wq1, 03071543.wq1, 03171543.wq1, and 03271543.wq1) and comprise 2,771 records in the FRDB. Data management occurred during Stage 1 of the data collection process.

These data were validated by BBL to ensure that they met method quality control criteria and the project data quality objectives. No formal SAP or QAPP was issued prior to implementation of sample collection or analysis; however, BBL stated they used collection and analytical procedures that had been approved by U.S. EPA Region 5 for other projects. Samples were submitted to Inchcape Testing Services Laboratory of Vermont for chemical analysis. PCB results were not surrogate-corrected.

The memorandum written by BBL dated April 4, 1998, indicates that PCB and TOC data for sediment samples and PCB data for biota were reviewed. Chain of custody procedures were not documented by BBL in this *Data Quality Assessment Memorandum*. Qualifiers were applied to sediment and biota data because of quantitative confirmation differences, blank contamination, and surrogate and matrix spike outlier values. The data, as qualified by BBL, are acceptable for use.

3.2.7 1995–1996 WDNR Fish Tissue Data

The WDNR collected fish tissue samples along the length of the river in 1996. These data were provided by WDNR in a single, multiple-page spreadsheet

(all_fish.wb1). Samples were analyzed for PCB Aroclors™ and TOC. This data set comprises 1,673 records in the FRDB and consists of 200 samples. Data management occurred during Stage 1 of the data collection process.

Data validation was performed by the M. A. Kuehl Company on 20 fish tissue samples collected by the WDNR in 1996. The data validation report for SDG-1 was reviewed by EcoChem. This data validation was performed using the specified methods described in the April 1996 *Addendum to the Quality Assurance Project Plan for Assessment of PCBs in Sediment of the Lower Fox River from De Pere to Green Bay for PCB Analysis of Fish Tissue*. Chain of custody records were reviewed and they indicated that samples were received in good condition. Precision and accuracy were judged to be acceptable by the M. A. Kuehl Company. PCB results were qualified because they were detected above the MDL but below the PQL. The data, as qualified by the M. A. Kuehl Company, are acceptable for use.

3.2.8 1996–1999 USFWS NRDA Fish Tissue Data

As part of the Natural Resource Damage Assessment (NRDA) investigation, the U.S. Fish & Wildlife Service (USFWS) collected and analyzed 376 tissue samples in 1996. Samples were collected below De Pere and in Green Bay. The samples were analyzed for PCB congeners or PCB Aroclors™ and TOC. The USFWS NRDA data represents 16,017 records in the FRDB and was provided by the USFWS in a single file (pcbsecd.dbf.) Data management occurred during Stage 1 of the data collection process.

A full data validation was conducted by EcoChem on 376 tissue samples analyzed for the Green Bay NRDA project. This data validation was performed based on the specified method criteria described in the Battelle Laboratory SOP, *Identification and Quantitation of Polychlorinated Biphenyls (by Congener and Aroclor™) and Chlorinated Pesticides by Gas Chromatography/Electron Capture Detection*. Accuracy and precision were generally acceptable. Qualifiers were assigned by EcoChem due to blank contamination, continuing calibration verification percent difference outliers, blank spike results, surrogate outliers, laboratory duplicate results, reference material recovery results, and chromatographic interferences. Data, as qualified by EcoChem, are acceptable for use.

3.2.9 1997 USFWS NRDA Waterfowl Tissue Data, 1994–1995 Cormorant Data, and 1993 USFWS Tree Swallow Data

Results from waterfowl tissue sample analyses were provided by USFWS in two files (tcuster2.mdb and tcuster2.wpd). The samples were analyzed for chlorinated pesticides. This data set consists of 70 samples and 1,680 analytical data points.

Results from cormorant tissue sample analyses were provided by USFWS in two files (tcuster1.mdb and tcuster1.wpd). The samples were analyzed for PCB Aroclors™, chlorinated pesticides, and dioxins. This data set consists of 193 samples and 6,178 analytical data points.

Results from tree swallow tissue sample analyses were provided by the USFWS in two files (ccuster.mdb and ccuster.wpd). The samples were analyzed for PCB congeners, chlorinated pesticides, and dioxins. This data set consists of 200 samples and 5,429 analytical data points. Data management for all data types occurred during Stage 1 of the data collection process.

Three electronic text files were reviewed by EcoChem for data validation information regarding these data sets. Files reviewed include 1997 waterfowl data from Green Bay and Lake Michigan (tcuster1.wpd), 1994 through 1995 double-crested cormorants data from Green Bay (tcuster2.wpd), and Fox River and Green Bay 1993 through 1995 Tree Swallow Study (ccuster.wpd). Of these three documents, one (tcuster1.wpd) gives a brief synopsis of field sampling and chemical analysis procedures used to collect and analyze the samples. The information provided did not specifically address chain of custody records associated with the samples. No qualifiers were assigned based on this review although the statement “concentrations of PCB 118 may be overestimated because of coelution with PCB 106” may be considered a qualification. With regards to quality assurance and quality control approval, a reference is made to the Patuxent Analytical Control Facility (Patuxent) of USFWS, Laurel, Maryland. It is not clear from this statement if Patuxent established the quality control criteria, approved the method of analysis, or reviewed the results of the study. For these reasons the data should be used only as supporting data.

3.2.10 Fox River Fish Consumption Advisory Data

The initial fish contaminant data in the FRDB represents tissue samples collected by WDNR in the Fox River and Green Bay between 1971 and 1996 were addressed as part of the Stage 1 effort. These samples were analyzed for

PCB congeners, PCB Aroclors™, metals, chlorinated pesticides, and dioxins. The FRDB contains 1,766 samples from the fish contaminant study comprising 11,620 records. This data set is primarily tissue data with a small number of sediment samples. Data management occurred during Stage 1 of the data collection process. A second delivery of 1998 fish contaminant data (tissue) was received during Stage 2 data collection. These data represent 130 samples and 777 data records in the FRDB and was conducted during Stage 2 of the data management process.

In 1995, the M. A. Kuehl Company conducted a laboratory audit at the Wisconsin SLOH. The purpose of this audit was to assess the laboratory capability to analyze tissue and sediment samples for PCB, TOC, and metals. Although she made a few observations and had a few findings, Ms. Kuehl found the laboratory to be capable of performing the requested analyses. The Wisconsin SLOH was also visited by EcoChem personnel, and analysts and managers were interviewed. Sample handling, preparation, and analysis systems were reviewed. In-depth discussions occurred concerning peak identification and quantitation. All hardcopy and electronic data are available, and could be fully validated if requested. As these data have not undergone full validation, these data should be used as supporting data only. Refer to Section 2.2.1 for further discussion of data generated by SLOH and refer to 2.2.18 for a discussion of the review of more recent data generated by SLOH.

3.2.11 WDNR Wildlife Tissue Data

This data set is a collection of wildlife tissue sample data collected by WDNR during the time period from 1984 to 1996 and collated in three files (all.db, geese.db, and ducks.db). The data set represents bird and mammal tissue samples analyzed for chlorinated pesticides. This data set contains 417 samples and 2,532 analytical data points. Data management occurred during Stage 1 of the data collection process.

Quality control information was not available, therefore these data should be used as supporting data only.

3.2.12 Lake Michigan Tributary Monitoring Data

The Lake Michigan Tributary Monitoring samples from the Fox River were collected by the U.S. Geological Survey (USGS) in support of the Lake Michigan Mass Balance Study, administered by the U.S. EPA's GLNPO. These water samples were analyzed for PCB congeners, chlorinated pesticides, and mercury. This data set consists of 88 samples and 5,722 analytical data points. Data management occurred during Stage 1 of the data collection process.

These data were validated by the M. A. Kuehl Company, and these data are considered useable, as qualified.

3.2.13 Stromberg Eagle Data

Eagle samples were collected for the USFWS under the direction of Ken Stromberg between 1991 and 1996. The data were provided by the USFWS in a text file report (strmbrg.wpd) and required manual extraction point by point. The samples were analyzed for PCB congeners, chlorinated pesticides, and dioxins. This data set contains 31 samples and 954 analytical data points. Data management occurred during Stage 1 of the data collection process.

Quality control information was not available, therefore these data should be used as supporting data only.

3.2.14 USGS NAWQA Data

The National Ambient Water Quality Assessment Program (NAWQA) data represent samples collected by the USGS between 1992 and 1997. There are 441 samples of sediment, water, and tissue. These samples were analyzed for an extensive list of chlorinated pesticides and herbicides, organophosphorus pesticides, semivolatile, and metallic analytes. These data were provided by the USGS in 21 files with additional information obtained on the NAWQA website. These sample analyses represent 11,879 records in the FRDB, approximately 90 percent of which is from waterways other than the Fox River and is noted as “reference.” Data management occurred during Stage 1 of the data collection process.

Of the 441 environmental samples collected between 1992 and 1997, approximately 15 percent were quality control samples collected concurrently during field sampling activities. Types of quality control samples collected include field blanks and trip blanks for surface water and groundwater matrices, and field replicates and splits for all matrices. Surface water and groundwater samples were spiked to assess precision and accuracy of the volatile and pesticide methods. Surrogates were added to all environmental samples undergoing pesticide, volatile, and other trace organic analyses.

The results of the quality control samples were reviewed by the USGS NAWQA group and were reported in the USGS Water-Resources Investigations Report 97-4148, *Results of Quality-Control Sampling of Water, Bed Sediment, and Tissue in the Western Lake Michigan Drainages Study Unit of the National Water-Quality Assessment Program*. All results were found to be

acceptable by NAWQA. Accuracy was generally acceptable, as demonstrated by the percent recovery values of the surrogate and matrix spike values. Precision was generally acceptable, as demonstrated by the relative percent difference values of the sample duplicates. While thorough investigations, and in some cases corrective actions, were performed to explain quality control anomalies (e.g., blank contamination, occasional poor spike recovery values, and possible interferences causing bias), no qualifiers were applied directly to the analytical results. In summary, the data user should refer to this report when using these data to gain a complete understanding of its limitations. As the content of the data packages is not known, the data may or may not be amenable to independent validation. For the reasons mentioned above, the NAWQA data should be used as supporting data only.

3.2.15 1994 Woodward-Clyde Deposit A Sediment Data

Sediment samples were collected by Woodward-Clyde in 1994 at Deposit A. These samples were analyzed for PCB Aroclors™ and TOC. They were provided by WDNR in 12 files, only one of which contained analytical data (pcb_to~1.xls). This data set contains 66 samples and represents 585 records in the FRDB. Data management occurred during Stage 1 of the data collection process.

A limited data validation was conducted by EcoChem (September 1998) on these data for the Little Lake Butte des Morts (LLBdM) Deposit A project. This data validation was performed using the specified methods described in the August 1994 *Quality Assurance Project Plan (QAPP) for the Pre-Design Study on Little Lake Butte des Morts*. It should be noted that the specific procedures to be used for data validation (Sections 2 and 9 of the QAPP) were slightly modified to account for differences in laboratory deliverables. For instance, holding times could not be assessed since chain of custody forms were not provided and a case narrative describing any deviations from proposed analysis was not provided. Accuracy was generally acceptable, as demonstrated by the percent recovery values of the surrogate, and matrix/blank spikes. Precision was generally acceptable, as demonstrated by the relative percent difference values of the sample and laboratory duplicates. Qualifiers were assigned by EcoChem due to poor matrix spike recovery values. Based on this limited review, all data, as qualified by EcoChem, are acceptable for use.

3.2.16 WPDES Permit Influent Data

Influent water samples along the Fox River were collected by various entities (commercial and governmental) as part of the Wisconsin Pollutant Discharge Elimination System (WPDES) regulatory program, then analyzed for various

fractions by WDNR-certified laboratories. These data were provided by WDNR in a spreadsheet and consist of samples collected in 1993 and 1997. These data do not adhere to a regular sampling schedule and were provided as supplemental water quality data. These data do not have associated QA/QC data, as the samples were not collected for an RI/FS-type activity. This data set consisted of eight samples and 847 records. Data management occurred during Stage 1 of the data collection process.

As QC information was not available, these data should be used only as supporting data.

3.2.17 Lower Fox River Background Metals Assessment

These data were collected from 1991 to 1993 and consist of 14 samples and 78 records in the FRDB. Data management occurred during Stage 1 of the data collection process.

Raw data and accompanying quality control information were not available for review. The data should be used only as supporting data.

3.2.18 1997 WDNR Caged Fish Bioaccumulation Study Data

WDNR placed caged fish near the demonstration projects conducted at Deposit N and SMU 56/57 prior to the initiation of the projects. The fish and collocated sediment samples were collected and analyzed for PCB congeners by the Wisconsin SLOH (for more discussion of SLOH, see Section 2.2.1). This data set consists of 25 samples and 1,672 records in the FRDB. Data management occurred during Stage 1 of the data collection process.

At the request of WDNR, select sediment and fish tissue data from this study were reviewed to show the quality of the older data (e.g., Green Bay Mass Balance) was consistent with that of the new data sets. The data packages from the laboratory consisted of strip charts containing the chromatograms and associated instrument printouts of the standards, QC sample results, and field sample results. Data packages summarizing calibration and other ancillary QC results (as provided under the EPA Contract Laboratory Program) were not available from the laboratory. The samples were analyzed using the protocol outlined in the *Quality Assurance Plan (QAP), Green Bay Mass Balance Study* (March 11, 1988). The data were reviewed using the criteria listed in the QAP and the *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994).

Overall, these sets of data met the QC criteria as specified in the QAP. Although not assigned in this review, qualifiers could be assigned due to surrogate and matrix spike outliers indicating the potential for high bias. It is unlikely that any data would be rejected.

As determined by this review, these data should be used as supporting data. Refer to Sections 2.2.1 and 2.2.10 for further discussion of data generated by SLOH.

3.2.19 1997 Demonstration Project Data – Deposit N

Sediment, water, and wipe samples were collected by Foth & Van Dyke from Deposit N. The environmental samples were analyzed for PCB Aroclors™, mercury, and TOC. This data set contains 10 samples and represents 83 records in the FRDB. Data management occurred during Stage 2 of the data collection process.

Full data validation was conducted by the M. A. Kuehl Company on approximately 10 percent of the 1997 Fox River Deposit N data (PCBs and mercury). A limited data review was conducted on the remainder of the data (PCBs, mercury, and TOC). Results of this evaluation indicate that the laboratory followed the specified methods described in the October 1997 *Fox River Deposit N Removal Project Pre-Design Phase Quality Assurance Project Plan*. Chain of custody documentation, although not referred to directly by M. A. Kuehl's December 26, 1997 *Technical Memorandum - Data Validation for Fox River Deposit N*, was acceptable (report mentions discrepancies only). PCB data were qualified due to holding time exceedances and poor matrix spike recovery. No qualifiers were assigned to the TOC and mercury data. Matrix spike and lab duplicates were not performed on water samples submitted for PCB analysis due to insufficient sample volumes. No action was taken because the laboratory performed alternative QC measures (control spikes) with acceptable recoveries. The data, as qualified by M. A. Kuehl, are acceptable for use.

3.2.20 1997–1998 Demonstration Project Data – SMU 56/57

Sediment samples were collected in late 1997 and early 1998. Montgomery Watson and Harrington Engineering & Construction implemented a sediment removal demonstration project at SMU 56/57 on behalf of the WDNR. The environmental samples were analyzed for a full suite of parameters that included PCB Aroclors™, mercury, and TOC. This data set contains 295 samples and represents 3,114 records in the FRDB. Data management occurred during Stage 2 of the data collection process.

Data validation was performed by Montgomery Watson on over 100 analytical batches of data collected at SMU 56/57 in 1997 and 1998. Full data validation was performed on sediment PCB and mercury data and a limited data review was conducted on all other analytical parameters. The full data validation and limited review were performed using the specified methods described in the *Field Sampling Plan Pre-Design Investigation Sediment Management Unit 56/57 Sediment Removal Demonstration Project* and accompanying *Quality Assurance Project Plan* (May 1998) and *U.S. EPA Contract Laboratory Program National Functional Guidelines for Organic Analysis Review* (February 1994). Chain of custody documentation was not covered in the data validation or the review. Precision and accuracy were judged to be acceptable by Montgomery Watson. PCB results were qualified as estimated by Montgomery Watson because PCBs were analyzed beyond holding times. Mercury results were qualified as estimated because matrix spike percent recovery values exceeded the control limit criteria. Results from other analytical methods were qualified for holding time exceedances (total Kjeldahl nitrogen results) and blank contamination (variety of conventionals analyses). Only the QC elements for the PCB and mercury sediment results were summarized in Table 3-2 due to the number of analytical tests performed on the effluent samples. Based on Montgomery Watson's limited review, the data are considered usable.

3.2.21 1998 RETEC RI/FS Supplemental Data

Supplemental sediment samples were collected from the Lower Fox River in June of 1998 by Remediation Technologies, Inc. (RETEC) for the WDNR. Samples were collected according to procedures outlined in the *Sampling and Analysis Plan and Quality Assurance Project Plan for Supplemental Data Collection, Fox River RI/FS*. This data set consists of 252 samples and 10,781 records in the FRDB. Data management occurred during Stage 1 of the data collection process.

A full data validation was conducted by EcoChem, Inc. (1998). Analytical data were reviewed using quality control criteria documented in the analytical method, National Functional Guidelines, and the project QAPP. Validation was performed on PCB, semivolatile, pesticide, metals, and conventional (TOC and total solids) data packages. Accuracy and precision were generally acceptable. Qualifiers were assigned by EcoChem due to holding time exceedances, blank contamination, continuing calibration verification percent difference outliers, lack of secondary column confirmation, blank and matrix spike outliers, surrogate outliers, laboratory duplicate results, and reference material recovery results. Data, as qualified by EcoChem, are acceptable for use.

3.2.22 Lake Michigan Mass Balance Data

The Lake Michigan Mass Balance samples were collected in 1994 and 1995. Sediment, water, tissue, and air samples were collected and were analyzed for PCB congeners, volatiles, pesticides/herbicides, metals, and wet chemistry tests. Electronic data were received on compact disc (CD) for 21 focus groups. This data set contains 6,987 samples and represents 91,621 records in the FRDB. Data management occurred during Stage 2 of the data collection process.

EcoChem, Inc. performed a review of the Lake Michigan Mass Balance (LMMB) Study QA program and assessed the quality of the data generated for the study. This evaluation of the quality assurance program included a review of the measurement quality objectives (MQOs), the *Lake Michigan Mass Balance (LMMB) Study QA and Data Management Workgroups Peer Review Meeting Briefing Book* (April 29–30, 1999), and the *Lake Michigan Mass Budget/Mass Balance Work Plan* (October 14, 1993). To clarify the QA process followed in this study, telephone interviews with several LMMB Study participants were conducted. Third-party review of the data was not performed, nor were raw data available for this review. Thus, the quality of the data was judged on the assumption that the QA program and the MQOs were met. Although the data were not reviewed by an independent third-party, sufficient information was available about the QA program to render a judgment on the probable usability of the data. The samples were analyzed for PCB congeners, pesticides, metals, atrazine, nutrients, conventionals, various biological measurements, lead 210 and cesium 137.

The samples were analyzed by reputable commercial and academic/research laboratories that were audited prior to sample analysis and again during sample analysis by the program QA personnel and by the U.S. EPA. The MQOs that were followed by the academic/research laboratories were different than those employed under the U.S. EPA Contract Laboratory Program (CLP); the U.S. EPA *Test Methods for Evaluating Solid Waste, Physical/Chemical Methods, SW-846, 3rd Edition* (as updated); or the U.S. EPA *National Functional Guidelines (NFG) for Organic Data Review* (February 1994) and *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994). For instance, the acceptability of the initial calibration, as specified by NFG, is measured by a correlation coefficient (r). The correlation coefficient must be greater than or equal to 0.995 (or $r^2 \geq 0.990$). For the congener analyses of the samples in this study, the criterion for several laboratories was that r^2 must be greater than or equal to 0.95. The criteria for this study used by each laboratory were approved by the U.S. EPA. However, because the QC criteria are different

from NFG, the precision and accuracy may differ from that of the data sets collected using NFG. Because of this, the data should be considered as supporting data only. Although it is likely that some data would be estimated if the data were reviewed by an independent third party using the U.S. EPA NFG criteria, it is unlikely that any data would be rejected.

3.2.23 Minergy Mineralogical Data

The Minergy data are comprised of results from the analysis of 15 sediment samples for 11 different mineral oxides, sulfur, chloride, and two different loss on ignition (LOI) procedures. Two hundred nineteen (219) analytical records were generated. Data management occurred during Stage 2 of the data collection process. The Mineral Lab analyzed the samples for mineral oxides, sulfur, and chloride. Badger Laboratories & Engineering performed the loss on ignition procedure.

EcoChem, Inc. performed a review of the Minergy site data generated for the study. The evaluation of the quality control elements with these analyses included telephone interviews with personnel at each laboratory. Third-party review of the data was not performed, nor were raw data available for this review. Thus, the quality of the data was judged solely on the information obtained during the telephone interviews. Although the data were not reviewed by an independent third party, sufficient information was available about the QA program to render a judgment on the probable usability of the data.

Based on the information received during the telephone interview with Badger Laboratories and Engineering, the LOI data are usable as reported.

Based on the information received during the telephone interview with The Mineral Lab, the mineral oxide, sulfur, and chloride data should be considered as estimated. The data users should be aware that these data may be potentially biased. The mineral oxide, sulfur, and chloride data should be considered as supporting data only; it is unlikely that any data would be rejected during a full validation.

3.2.24 1998 FRG/Exponent Data

Exponent collected tissue samples in the summer of 1998 for the Fox River Group (FRG). Samples were collected from Little Lake Butte des Morts to Green Bay Zone 3 and were analyzed for PCB congeners and PCB Aroclors™, pesticides/herbicides, metals, and wet chemistry tests. The data set contains

225 samples that account for 17,708 records in the FRDB. Data management occurred during Stage 2 of the data collection process.

EcoChem performed a review of the FRG 1998 data validation reports authored by Exponent, Inc. EcoChem evaluated the validation reports for completeness and technical agreement. To clarify some of the findings, raw data were reviewed. The samples were analyzed by U.S. EPA SW-846 methodology and other miscellaneous EPA methods. The data were validated by BBL using the *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994); *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994), and *Lower Fox River System NRDA Quality Assurance Project Plan* (December 1998).

Overall, the data are of acceptable quality. The samples were analyzed and validated as specified in the QAPP. A more detailed review of the data would result in additional qualifiers being assigned. As determined by this review, the data, as qualified, are usable for the intended purpose.

3.2.25 1998 FRG/BBL Sediment/Tissue Data

BBL collected tissue, sediment and water samples in 1998 for the FRG. Samples were analyzed for semivolatiles, PCB congeners and PCB Aroclors™, pesticides/herbicides, radchem, metals, and wet chemistry tests. The data set contains 1,315 samples that account for 18,824 records in the FRDB. Data management occurred during Stage 2 of the data collection process.

EcoChem performed a review of the FRG 1998 data validation reports authored by BBL. EcoChem evaluated the validation reports for completeness and technical agreement. To clarify some of the findings, raw data were reviewed. The samples were analyzed by U.S. EPA SW-846 methodology and other miscellaneous EPA methods. The data were validated by BBL using the *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994), *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994), and *Lower Fox River System NRDA Quality Assurance Project Plan* (December 1998).

Overall, the data are of acceptable quality. The samples were analyzed and validated as specified in the QAPP. In some cases, criteria from NFG, rather than the analytical method criteria, were used to evaluate the data. A more detailed review of the data would result in additional qualifiers being assigned. It is unlikely that any more data would be rejected. As determined by this review, the data, as qualified, are usable for the intended purpose.

3.2.26 1998–1999 Deposit N Data: Remediation/Pre-Dredge/Post-Dredge/Operational Monitoring

Data for the Deposit N pilot remediation project was received in four sections: pre-dredge data, post-dredge data, operational monitoring data, and sediment remediation (environmental monitoring) data. Collectively, sediment, tissue, and water samples were collected and analyzed for PCB Aroclors™, PCB congeners, metals, and wet chemistry tests. The Deposit N pilot remediation data represents 305 samples and accounts for 12,514 records in the FRDB. Data management occurred during Stage 2 of the data collection process.

EcoChem performed a review of the data validation reports authored by the M. A. Kuehl Company. EcoChem evaluated the validation reports for completeness and technical agreement. To clarify some of the findings, raw data were reviewed.

The samples were analyzed by U.S. EPA SW-846 methodology. The data were validated using the Region 5 Modifications to *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994), *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994), and the *Fox River Group Deposit N Demonstration Project Quality Assurance Project Plan* (1998).

Overall, the data are of acceptable quality. The samples were analyzed and validated as specified in the QAPP. A more detailed review of the data would result in additional qualifiers being assigned in some cases and qualifiers being removed in others. It is unlikely that any more data would be rejected. As determined by this review, the data, as qualified, are usable for the intended purpose.

3.2.27 Ankley and Call Data

EcoChem conducted a data entry process on data presented in the *Sediment Quality Evaluation in the Lower Fox River and Southern Green Bay of Lake Michigan Report*. A second party verified the data entry. These data represent 62 individual samples and comprises 1,607 records in the FRDB. Data management occurred during Stage 2 of the data collection process.

EcoChem did not conduct any data quality assessment on these data. The quality of the data is therefore indeterminate.

3.2.28 State of Michigan Fish Consumption Advisory Data

The State of Michigan Fish Consumption Advisory data included in the FRDB are the results of fish tissue samples collected between 1983 and 1999. The samples were from Green Bay zones 3A and 4, as well as from tributaries flowing into Green Bay. The samples were analyzed for PCB Aroclors™, pesticides/herbicides, dioxins, metals, and wet chemistry tests. The data represents 434 samples and accounts for 6,979 records in the FRDB. Data management occurred during Stage 2 of the data collection process.

At the request of the WDNR, EcoChem performed a review of the FRG 1998 data validation reports authored by Exponent, Inc. See Table 3-1 for a listing of reports and samples. EcoChem was to evaluate the validation reports for completeness and technical agreement. To clarify some of the findings, raw data were reviewed.

The samples were analyzed by U.S. EPA SW-846 methodology and other miscellaneous EPA methods. The data were validated by BBL using the *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994), *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994), and *Lower Fox River System NRDA Quality Assurance Project Plan* (December 1998).

Overall, the data are of acceptable quality. The samples were analyzed and validated as specified in the QAPP. A more detailed review of the data would result in additional qualifiers being assigned.

As determined by this review, the data are usable for the intended purpose.

3.2.29 1999 Demonstration Project Data – SMU 56/57

These data are in the process of being appended to the database.

At the request of the WDNR, EcoChem performed a review of the FRG data validation reports for the 1999 SMU 56/57 and Deposit N demonstration projects authored by the M. A. Kuehl Company.

The samples were analyzed according to U.S. EPA SW-846 methodology. The data were validated using *U.S. EPA Region 5 Standard Operating Procedure for Validation of CLP Organic Data* (February 1997), *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994), *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994), *Draft Quality Assurance Project Plan Environmental Monitoring of SMU 56/57 Demonstration*

Project – Mass Balance Approach, Revision I (August 1999), and the Draft Quality Assurance Project Plan Monitoring of Deposit N Demonstration Project – Mass Balance Approach (December 1998).

Overall, the data are of acceptable quality. The samples were analyzed and validated as specified in the QAPP. A more detailed review of the data would result in additional qualifiers being assigned in some cases. It is unlikely that any more data would be rejected. As determined by this review, the data are usable for the intended purpose. No further review is recommended at this time.

3.3 Data Usability

3.3.1 Fully Validated Data

The following data sets have been validated by an independent party and are considered useable, as qualified:

- 1994 GAS/SAIC Sediment Data;
- 1994 Woodward-Clyde Deposit A Sediment Data;
- 1995 WDNR Sediment Data;
- 1996–1999 USFWS NRDA Fish Tissue Data;
- 1995–1996 WDNR Fish Tissue Data;
- 1997–1998 Demonstration Project Data – SMU 56/57;
- 1998 RETEC RI/FS Supplemental Data;
- 1996 FRG/BBL Sediment/Tissue Data;
- 1997 Demonstration Project Data – Deposit N;
- 1992–1993 BBL Deposit A Sediment Data;
- 1998 FRG/Exponent Data;
- 1998 FRG/BBL Sediment/Tissue Data;

- 1998–1999 Deposit N Data: Remediation/Pre-Dredge/Post-Dredge/Operational Monitoring;
- 1999 Demonstration Project Data – SMU 56/57;
- State of Michigan Fish Consumption Advisory Data; and
- Lake Michigan Tributary Monitoring Data.

Although the data sets (listed above) were found to be validated and useable, it must be stressed that there were individual data points that were rejected. These rejected data points have not been used in support of the RI/FS or RA.

3.3.2 Supporting Data

The following data sets have not been validated and, in general, should be used only as supporting data. The data have been collected within different programs and with different data quality objectives therefore, varying degrees of supporting documentation may be available.

- 1989–1990 Fox River Mass Balance Study,
- 1989–1990 Green Bay Mass Balance Study (GLNPO),
- 1993 Triad Assessment,
- 1993 USFWS Tree Swallow Data,
- 1994–1995 Cormorant Data,
- 1997 USFWS NRDA Waterfowl Tissue Data,
- 1997 WDNR Caged Fish Bioaccumulation Study Data,
- Fox River Fish Consumption Advisory Data,
- Stromberg Eagle Data,
- USGS NAWQA Data,
- WDNR Wildlife Tissue Data,
- WPDES Permit Influent Data,
- Lake Michigan Mass Balance Data,
- Minergy Mineralogical Data, and
- Lower Fox River Background Metals Assessment.

3.3.3 Indeterminate Data

The following data sets have not been validated and have not been subjected to a data quality review. This is due to complete lack of supporting QA/QC documentation; or, the hardcopy data and documents were not received by

EcoChem by the date of this report. At this time, the overall quality of these data sets is unknown and the data should be used with that fact in mind.

- Ankley and Call Data

Table 3-1 Data Set Analysis

| Data Source | Number of Samples | Matrices ¹ | Analyses Conducted ² | Number of Records | Number of Files in Delivery | File Type | Report Section | Earliest Year of Collection | Latest Year of Collection |
|---|-------------------|-----------------------|--|-------------------|-----------------------------|-------------|----------------|-----------------------------|---------------------------|
| 1989–1990 Fox River Mass Balance Study | 1,967 | S, W | PCB-A, PCB-C, W | 25,457 | 6 | Spreadsheet | 2.2.01 | 1989 | 1990 |
| 1989–1990 Green Bay Mass Balance Study (GLNPO) | 2,069 | S, T, W | B, PCB-C, W | 201,701 | 92 | Database | 2.2.01 | 1987 | 1990 |
| 1992–1993 BBL Deposit A Sediment Data | 117 | S, W | M, P/H, PCB-A, SVOA, V, W | 1,094 | 1 | Spreadsheet | 2.2.02 | 1992 | 1993 |
| 1993 Triad Assessment | 27 | S | B, M, P/H, PCB-A, SVOA, W | 631 | 11 | Spreadsheet | 2.2.03 | 1992 | 1993 |
| 1993 USFWS Tree Swallow Data | 200 | T | B, DXN, P/H, V, W | 5,429 | 2 | Database | 2.2.09 | 1993 | 1993 |
| 1994 GAS/SAIC Sediment Data | 253 | S | DXN, M, P/H, PCB-A, SVOA, V, W | 5,654 | 6 | Spreadsheet | 2.2.04 | 1994 | 1994 |
| 1994 Woodward-Clyde Deposit A Sediment Data | 66 | S | PCB-A, W | 585 | 12 | Spreadsheet | 2.2.15 | 1994 | 1994 |
| 1994–1995 Cormorant Data | 193 | T | B, DXN, P/H, PCB-C, W | 6,178 | 2 | Database | 2.2.09 | 1994 | 1995 |
| 1995 WDNR Sediment Data | 488 | S | M, PCB-A, W | 6,433 | 8 | Spreadsheet | 2.2.05 | 1995 | 1995 |
| 1996 FRG/BBL Sediment/Tissue Data | 25 | S, T | B, PCB-C, W | 2,771 | 6 | Spreadsheet | 2.2.06 | 1996 | 1996 |
| 1995–1996 WDNR Fish Tissue Data | 200 | T | B, PCB-A, W | 1,673 | 1 | Spreadsheet | 2.2.07 | 1995 | 1996 |
| 1997 Demonstration Project Data - Deposit N | 10 | S | M, PCB, W | 83 | 1 | Spreadsheet | 2.2.19 | 1997 | 1997 |
| 1997–1998 Demonstration Project Data - SMU 56/57 | 295 | S, W | DXN, M, P/H, PCB-A, SVOA, V, W | 3,114 | 12 | Spreadsheet | 2.2.20 | 1997 | 1998 |
| 1997 USFWS NRDA Waterfowl Tissue Data | 70 | T | B, P/H, PCB, V, W | 1,680 | 2 | Database | 2.2.09 | 1997 | 1997 |
| 1997 WDNR Caged Fish Bioaccumulation Study Data | 25 | S, T | B, PCB-C, W | 1,672 | 2 | Spreadsheet | 2.2.18 | 1997 | 1997 |
| 1998 FRG/BBL Sediment/Tissue Data | 1,315 | S, T, W | B, M, P/H, PCB-A, PCB-C, RAD, SVOA, W | 18,824 | 1 | Database | 2.2.25 | 1998 | 1998 |
| 1998–1999 Deposit N Data: Post-Dredge | 43 | S | PCB-A, PCB-C, W | 690 | 8 | Spreadsheet | 2.2.26 | 1999 | 1999 |
| 1998–1999 Deposit N Data: Pre-Dredge | 53 | S | PCB-A, PCB-C, W | 1,437 | 6 | Spreadsheet | 2.2.26 | 1998 | 1998 |
| 1998 FRG/Exponent Data | 225 | T | B, M, P/H, PCB-A, PCB-C, W | 17,708 | 3 | Database | 2.2.24 | 1998 | 1998 |
| 1998 RETEC RI/FS Supplemental Data | 252 | S, T | B, DXN, M, P/H, PCB-A, PCB-C, SVOA, V, W | 10,781 | 1 | ASCII | 2.2.21 | 1998 | 1998 |
| Fox River Fish Consumption Advisory Data: 1998 WDNR Fish Consumption Data | 130 | T | B, M, PCB-A, W | 777 | 1 | ASCII | 2.2.10 | 1998 | 1998 |
| 1998–1999 Deposit N Data: Remediation Data | 197 | T, W | PCB-C, W | 10,264 | 1 | Spreadsheet | 2.2.26 | 1998 | 1999 |
| Ankley and Call Data | 62 | PW, S, T, W | DXN, M, P/H, PCB, SVOA, W | 1,607 | 0 | Hardcopy | 2.2.27 | 1989 | 1989 |
| 1998–1999 Deposit N Data: Operational Monitoring Data | 12 | S | M, PCB-A, W | 123 | 1 | Spreadsheet | 2.2.26 | 1998 | 1998 |
| Fox River Fish Consumption Advisory Data | 1,766 | S, T | B, DXN, M, P/H, PCB-A, PCB-C, SVOA, V, W | 11,620 | 2 | ASCII | 2.2.10 | 1971 | 1996 |
| State of Michigan Fish Consumption Advisory Data | 434 | T | B, DXN, M, P/H, PCB-A, W | 6,979 | 1 | Database | 2.2.28 | 1983 | 1999 |
| Lake Michigan Mass Balance Data | 6,987 | A, S, T, W | M, P/H, PCB-C, V, W | 91,621 | 211 | Database | 2.2.22 | 1993 | 1996 |
| Lake Michigan Tributary Monitoring Data | 88 | W | M, P/H, PCB-C, V | 5,722 | 5 | Spreadsheet | 2.2.12 | 1994 | 1995 |
| Lower Fox River Background Metals Assessment | 14 | W | M | 78 | 1 | Spreadsheet | 2.2.17 | 1991 | 1993 |
| Minergy Mineralogical Data | 15 | S | W | 219 | 1 | Spreadsheet | 2.2.23 | 1995 | 1999 |
| Stromberg Eagle Data | 31 | T | B, DXN, P/H, PCB-A, PCB-C, SVOA, V, W | 954 | 1 | ASCII | 2.2.13 | 1991 | 1996 |
| 1996–1999 USFWS NRDA Fish Tissue Data | 376 | T | DXN, P/H, PCB-A, PCB-C, W | 16,017 | 5 | Spreadsheet | 2.2.08 | 1996 | 1999 |
| USGS NAWQA Data | 441 | S, T, W | B, M, P/H, PCB, SVOA, V, W | 11,879 | 21 | Spreadsheet | 2.2.14 | 1992 | 1997 |
| WDNR Wildlife Tissue Data | 417 | T | B, M, P/H, PCB-A | 2,532 | 3 | Database | 2.2.11 | 1984 | 1996 |
| WPDES Permit Influent Data | 8 | W | B, DXN, M, P/H, PCB-A, RAD, SVOA, V, W | 847 | 1 | Spreadsheet | 2.2.16 | 1993 | 1997 |
| Total: 35 Data Sets | 18,871 | | | 474,834 | 438 | | | | |

¹ Matrices:

A - Ambient Air
PW - Sediment Pore Water
S - Sediment
T - Tissue
W - Water

² Analyses:

B - Biological
DXN - Dioxins
M - Metals
PCB - Total PCBs only
PCB-A - PCB Aroclor

PCB-C - PCB Congener
P/H - Pesticides/Herbicides
SVOA - Semivolatiles
V - Volatiles
W - Wet Chemistry (including all physical and conventional data)

Table 3-2 QC Elements for Data Sets Supporting the Fox River RI/FS and RA

| Parameters: Requirements | 1989–1990 Green Bay Mass Balance Study (GLNPO) | 1995–1996 WDNR Fish Tissue Data | 1996 USFWS/ Hagler Bailly Data | 1995 WDNR Sediment Data | | |
|---|---|---|---|--|---|-----------------------------|
| | PCBs Sediment | PCB Fish Tissue | PCB Fish Tissue | PCBs Sediment | TOC Sediment | Metals Sediment |
| SDG #s | University of Minnesota - Data groups: IN0042, IN0047, IN0052, IN0057, IN0061, IN0070, IN0076, IN0078, IN0037, and IN0041 | SLOH Fish SDG-1 | Battelle Laboratory Multiple SDGs | Hazleton SDG #s TBD2, 10, 1 & 20 | Hazleton SDG #s TBD2, 10, 1 & 20 | Hazleton SDG #s TBD2 & 20 |
| Data Review 1) Third-party Validation Performed | Verification Only Deborah Swackhamer, Ph.D. | M. A. Kuehl Co. | EcoChem | Y - M. A. Kuehl | Y - M. A. Kuehl | Y - M. A. Kuehl |
| Deliverables 1) Electronic Deliverables | Y | Y | Y | Y | Y | Y |
| 2) Hardcopy | Some - Not sure if this is a complete set | Y | Y | Some | Some | Some |
| Data Review Details 1) Package Completeness | Not determined | Y | Y | Y | Y | Y |
| 2) Chain of Custody Procedures | Not determined | Not determined | Y - Minor issues | Not determined | Not determined | Not determined |
| 3) Holding Times | Not summarized on the QA/QC Summary Report Sheet | Y | Y | Y | Y | Y |
| 4) Initial Calibration | Not summarized on the QA/QC Summary Report Sheet | Y (25%) | Y (35%) | 25% | Y | Y |
| Curve (# of standards) | Not summarized on the QA/QC Summary Report Sheet | 5 pt | 5 pt | 5 pt | Daily 1 pt | 1 pt/6 pt for Hg |
| 5) Calibration Verification | Not summarized on the QA/QC Summary Report Sheet | 15 %D | Varies between GC/EC/D & GC/MS, <25% for 75% analytes | 15% | 20% | 10% for metals & 20% for Hg |
| Secondary Column | Not summarized on the QA/QC Summary Report Sheet | 25 %D | Y - Data not used | 25 %D for CC on 2 nd column | NA | NA |
| 6) Laboratory Blanks | Not clear | Y | Y | Y | Y | Y |
| 7) Surrogate Recoveries (# required) | Y - 50%–120% | Y - 70%–120% | Y - 50%–125% | 60%–150% | NA | NA |
| 8) Matrix Spike (# required) | Y - 50%–120% | Y - 65%–125% | Y - 50%–125% tri- & deca- 30%–125% for mono- & dichloro- | 65%–125% | 75%–125% | 75%–125% |
| 9) Lab Duplicate | Y - Not clear what limits are | Y - 26% limit | Y - 50% | 26% | 20% | 20% |
| Lab Control Sample (SRM results?) | None/QAPP says that series of blindly-coded QA samples will be analyzed | N | SRM NRC %D Carp-1 <35% | NA | NA | Y - EPA |
| 10) Gel Permeation/Forisil Cleanup | Not provided | Y | Not mentioned | Y | NA | NA |
| 11) Detection Limit | Not provided | 50 µg/kg | Results reported to 0 | 50 ppb | NA | CRDL |
| 12) Calc and Transposition Verification (Qualitative verification?) | Not able to determine if this was done | Y - Recalc. | Y - Recalc. & verification | Y - Recalc. performed >10% frequency | NA | 10% |
| 13) Field QC Results | Not apparent | NA | None | None | None | None |
| 14) Usability Usable/ Supporting Qualifiers | Y Qualifiers mentioned but not defined | Usable Y - Minor J quals due to detections below PQL | Usable Y - Qualls due to CCV %D outliers, BS results, surr. outliers, lab dups., SRM results & interferences | Usable Y - Minor J flags due to low surr. recovery or below PQL and above MDL | Usable Y - Minor J flags due to poor lab RPD | Usable None |
| 15) Other IC Samples | NA NA | | | NA | NA | 20% |
| SAP | N - Study Plan | | N | Y | | |
| QAPP | Y | | Y - Tech Memo | Y | | |
| Lab QAM | Answer Pending/U of M SOPs? | Y | Y - Tech Memo | Y - Hazleton SOPs | | |

Table 3-2 QC Elements for Data Sets Supporting the Fox River RI/FS and RA

| Parameters: Requirements | Lake Michigan Mass Balance Data | 1998 Fox River NRDA | | | | |
|---|--|---|--|---|---|---|
| | Asst. Convs., Pest/PCB, Hg, Atrazine, DEA, DIA Water (open lake, tributary), Air, Sediment, Phytoplankton | PCB Fish Tissue | PCB Congener Fish Tissue | PCB Congener Fish Tissue | Pesticide Fish Tissue | Mercury Fish Tissue |
| SDG #s | BALN, GPLN, GRAN, GRLN, IUAA, IUAP, LHLL, LHLM, LHTN, LHPT, MDLH, MIAH, MNPH, RUAP, RULA, RUTA, SSSP, USTN, WSAW, WWTH, WWTN | Enchem Multiple SDGs | Michigan State University | Quanterra | Enchem Multiple SDGs | Enchem Multiple SDGs |
| Data Review 1) Third-party Validation Performed | N - Data reviewed by QC Coordinators | Exponent | Exponent | Exponent | Exponent | Exponent |
| Deliverables 1) Electronic Deliverables | Y | Y | Y | Y | Y | Y |
| 2) Hardcopy | Unknown | Y | Y | Y | Y | Y |
| Data Review Details 1) Package Completeness | Not addressed | Y | Y | Y | Y | Y |
| 2) Chain of Custody Procedures | Not addressed | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable |
| 3) Holding Times | No DV reports provided | Y | Some exceedances samples J/UJ | Y | Some exceedances samples J/UJ | Y |
| 4) Initial Calibration | No DV reports provided | Y | Y | Y | Y | Y |
| Curve (# of standards) | No DV reports provided | Y | Y | Y | Y | Y |
| 5) Calibration Verification | No DV reports provided | 20% | 20% | 20% | 20% | 10% |
| Secondary Column | No DV reports provided | Y | Y | Y | Y | NA |
| 6) Laboratory Blanks | No DV reports provided | Y | Y - U based on BC | Y | Y | Y |
| 7) Surrogate Recoveries (# required) | No DV reports provided | Y | Y | Y | Y | Y |
| 8) Matrix Spike (# required) | No DV reports provided | Y - No quals. for %R outliers | Y - No quals. for %R outliers | Y - No quals. for %R outliers | Y | Y |
| 9) Lab Duplicate | No DV reports provided | Y - MS/MSD | Y - MS/MSD | Y - MS/MSD | Y - MS/MSD | Y |
| Lab Control Sample (SRM results?) | No DV reports provided | Y | Y | Y | Y | Y |
| 10) Gel Permeation/Forisil Cleanup | No DV reports provided | Not mentioned | Not mentioned | Not mentioned | Not mentioned | NA |
| 11) Detection Limit | No DV reports provided | NA | NA | NA | NA | NA |
| 12) Calc and Transposition Verification (Qualitative verification?) | No recalculations were provided unable to determine if transcription checks were done | No recalcs. provided, unable to determine if transcription checks were done | No recalcs. provided, unable to determine if transcription checks were done | No recalcs. provided, unable to determine if transcription checks were done | No recalcs. provided, unable to determine if transcription checks were done | No recalcs. provided, unable to determine if transcription checks were done |
| 13) Field QC Results | Not addressed | None identified | None identified | None identified | None identified | None identified |
| 14) Usability Usable/ Supporting Qualifiers | Supporting Y - Specific LLMB 3-character qual. codes | Usable Y - HT, surr. %R, LCS %R | Usable - Some results rejected for low surr. %R Y - Surr. %R, BC, U, coplanars, J/UJ diff between GC & HRGCMS, interference, coelutions | Usable Y - Coelutions >calibration range | Usable Y - HT, MS/MSD %R, surr. %R, PCB interference, all +J | Usable Y - Dup RPD |
| 15) Other IC Samples | | | | | | |
| SAP | | | | | | |
| QAPP | | | | | | |
| Lab QAM | | | | | | |

Table 3-2 QC Elements for Data Sets Supporting the Fox River RI/FS and RA

| Parameters: Requirements | 1994 GAS/SAIC Sediment Data | | | | | | | |
|---|--|--|--|--|---|--|--|---|
| | PCBs Sediment | PCBs Sediment | PCBs Sediment | PCBs Sediment | PCBs Sediment | PCBs Sediment | PCBs Sediment | PCBs Sediment |
| SDG #s | ARI M172 | ARI M174 | ARI M176 | ARI M177 | ARI M178/ M179/M364 | ARI M365 | ARI M367/M368 | ARI M370 |
| Data Review 1) Third-party Validation Performed | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC |
| Deliverables 1) Electronic Deliverables | Y | Y | Y | Y | Y | Y | Y | Y |
| 2) Hardcopy | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed |
| Data Review Details 1) Package Completeness | Y | Y | Y | Y | Y | Y | Y | Y |
| 2) Chain of Custody Procedures | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined |
| 3) Holding Times | Y (frozen) | Y - Some exceedances | Y | Y | Y - Some exceedances, 1 sample qual. J for gross exceedances (M178) | Y - Exceedances, several samples qual. J for gross exceedances (M365) | Y - Minor violations | Y - Minor violations |
| 4) Initial Calibration Curve (# of standards) | Y 3-5 pt | Y 3-5 pt | Y 5 pt | Y 5 pt | Y 5 pt | Y 5 pt | Y 5 pt | Y 5 pt |
| 5) Calibration Verification Secondary Column | 15 %D but avg. was higher, results flagged (J/UJ) Not indicated | 15 %D but avg. was higher, results flagged (J/UJ) Not indicated | 15 %D but avg. was higher, results flagged (J/UJ) Not indicated | 15 %D but avg. was higher, results flagged (J/UJ) Not indicated | 15 %D but avg. was higher, results flagged (J/UJ) Not indicated | 15 %D but avg. was higher, results flagged (J/UJ) Not indicated | 15 %D but avg. was higher, results flagged (J/UJ) Not indicated | 15% Not indicated |
| 6) Laboratory Blanks | Y | Y | Y | Y | Y | Y | Y | Y |
| 7) Surrogate Recoveries (# required) | TCMX 55%-115%/DCB 70%-125% | TCMX 55%-115%/DCB 70%-125% | TCMX 55%-115%/DCB 70%-125% | TCMX 55%-115%/DCB 70%-125% | TCMX 55%-115%/DCB 70%-125% | TCMX 55%-115%/DCB 70%-125% | TCMX 55%-115%/DCB 70%-125% | TCMX 55%-115%/DCB 70%-125% |
| 8) Matrix Spike (# required) | 35% min-130% max | 35% min-130% max | 35% min-130% max | 35% min-130% max | 35% min-130% max | 35% min-130% max | 35 min%-130% max | 35 min%-130% max |
| 9) Lab Duplicate Lab Control Sample (SRM results?) | N Y | Not mentioned Y | Not mentioned Y | Not mentioned Y | Not mentioned Y | Not mentioned Y | Not mentioned Y | Not mentioned Y |
| 10) Gel Permeation/Forisil Cleanup | Y - If necessary | Y - If necessary | Not sure | Not sure | Not sure | Not sure | Not sure | Not sure |
| 11) Detection Limit | 50 ppb wet wt | NA | NA | NA | NA | NA | NA | NA |
| 12) Calc and Transposition Verification (Qualitative verification?) | Y - 10%? | N - No chros | ID & quants. could not be verified, raw data not provided | ID & quants. could not be verified, raw data not provided | ID & quants. could not be verified, raw data not provided | Data verified | N | Not verified |
| 13) Field QC Results | None | None | None | Not identified | Not identified | Not identified | Not identified | Not identified |
| 14) Usability Usable/ Supporting Qualifiers | Usable Y - Minor quals. assigned due to CCV (J/UJ) | Usable Y - Minor quals. assigned due to CCV (J/UJ) | Usable Y - Minor quals. assigned due to CCV, surr. recoveries J/UJ | Usable Y - Minor quals. assigned due to CCV, surr. recoveries J/UJ | Usable Y - Minor quals. assigned due to CCV, surr. recoveries J/UJ | Usable Y - Minor quals. assigned due to CCV, surr. recoveries J/UJ | Usable Y - Minor quals. assigned due to CCV, surr. recoveries J/UJ | Usable Y - Minor quals. assigned due to surr. recoveries J/UJ |
| 15) Other IC Samples | | | | | | | | |
| SAP | Y | | | | | | | |
| QAPP | Y | | | | | | | |
| Lab QAM | | | | | | | | |

Table 3-2 QC Elements for Data Sets Supporting the Fox River RI/FS and RA

| Parameters: Requirements | 1994 GAS/SAIC Sediment Data (Continued) | | | | | | |
|---|---|--|---|---|---|--|--------------------------------|
| | Dioxins Sediment | CLP Pest/PCBs Sediment | CLP SVOCs Sediment | CLP Metals Sediment | TCLP Metals Sediment | Mercury Sediment | Mercury Sediment |
| SDG #s | Triangle Lab SDG #35589 | Swanson/SDG 948521 | Swanson/SDG 948521 | Swanson/SDGs 12718, 12724, 12745, 12806, 12816, 12941 | Swanson/SDGs 12718, 12724, 12730, 12827, 12718, 12802, 12833, 12844 | Swanson WL12941 | Swanson WL12745 |
| Data Review 1) Third-party Validation Performed | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC |
| Deliverables 1) Electronic Deliverables | Y | Y | Y | Y | Y | Y | Y |
| 2) Hardcopy | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed |
| Data Review Details 1) Package Completeness | Y | Y | N - Forms 1 not supplied by lab | Y | Y | N - Forms 1 not supplied by lab | Y |
| 2) Chain of Custody Procedures | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined |
| 3) Holding Times | Y - Minor violations | N - Samples sent to TL 10 days after collection | N - All samples exceeded HT & are qual. as estimated (J/U) | Y - Hg results are flagged for exceeding HT by 27-42 days (J/U) | Y | N - All samples exceeded HT & are qual. as estimated (J/U) | Y |
| 4) Initial Calibration Curve (# of standards) | Y 5 pt | Y - Not consistent with CLP protocol 5 pt | Y - Not consistent with CLP protocol 5 pt | Y (validator recal. Hg results) Lin Reg | Y Lin Reg | Y - Exceedance 5 pt | Y - Exceedance 5 pt |
| 5) Calibration Verification Secondary Column | 20 %RSD NA | N - Correct concentration not used, certain analytes outside RT window Not indicated | 15 %D - Some exceedances qual. samples as estimated J/U Not indicated | 10 %D NA | 10 %D NA | Y - 15% NA | Y - 15% NA |
| 6) Laboratory Blanks | Y | Y | Y | Y | Y | Y | Y |
| 7) Surrogate Recoveries (# required) | TCFD 25%-150%/TCDD 25%-150% | TCMX 55%-115%/DCB 70%-125% | 8 required, 18% min-137% max | NA | NA | NA | NA |
| 8) Matrix Spike (# required) | TCDD/TCDF 54-162 | 18/9 required, 29 min-152 max | 11 required, 11% min-142% max | 75%-125% | 75%-125% | 75%-125% | 75%-125% |
| 9) Lab Duplicate Lab Control Sample (SRM results?) | Not mentioned Y | Not mentioned Y | Not mentioned Y - Acenaphthene fell outside @ 53% | Y - 20%, Some exceedances qual. J/U Y | Y Y | Y Y | Y Y |
| 10) Gel Permeation/Forisil Cleanup | Not sure | Not sure | Not sure | NA | NA | NA | NA |
| 11) Detection Limit | Elevated in some samples due to BC & noise | Elevated in some samples due to BC & noise | NA | NA | NA | NA | NA |
| 12) Calc and Transposition Verification (Qualitative verification?) | Y - Sample IDs, sample quant. not reviewed | Not verifiable | Y | Y - Some calc. errors | Y | N | N |
| 13) Field QC Results | Not identified | Not identified | Not identified | None | N | Y - FD | N |
| 14) Usability Usable/ Supporting Qualifiers | Usable Y - Due to BC & elevated MSR sample results may be biased positive (J+) | Third-party validation considers it unusable Y - Major issues about overall quality of data, assoc. with RT drift, quality of work poor | Usable Y - Minor quals. due to HT exceedances & low surr. & spike recoveries (J/U) | Usable - 1 data point rejected for Zn Y - Minor & major quals. due poor spike recoveries (J/U) & (R) on Zn | Usable No quals. | Usable Y - Minor J flags | Usable Y - Minor UJ/J flags |
| 15) Other IC Samples | | | | | | | |
| SAP | | | | | | | |
| QAPP | | | | | | | |
| Lab QAM | | | | | | | |

Table 3-2 QC Elements for Data Sets Supporting the Fox River RI/FS and RA

| Parameters: Requirements | 1994 GAS/SAIC Sediment Data (Continued) | | | | | 1998 Fox River Group | | |
|---|---|--|---|---|---|---|---|---|
| | Mercury Sediment | Mercury Sediment | Mercury Sediment | Mercury Sediment | Mercury Sediment | PCB Surface Water | Conventional Surface Water | PCB Sediment |
| SDG #s | Swanson WL12806 | Swanson WL12812/ 12724/12718 | Swanson WL12816/12882/ 12929/12922/ 12853/12852/12851 | Swanson WL12688/ 12725/12783/ 12777 | Swanson WL12693 | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs |
| Data Review 1) Third-party Validation Performed | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Y - SAIC | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee |
| Deliverables 1) Electronic Deliverables | Y | Y | Y | Y | Y | Y | Y | Y |
| 2) Hardcopy | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y - but not easily accessed | Y | Y | Y |
| Data Review Details 1) Package Completeness | Y | Y | Y | Y | Y | Y | Y | Y |
| 2) Chain of Custody Procedures | Not determined | Not determined | Not determined | Not determined | Not determined | Acceptable | Acceptable | Acceptable |
| 3) Holding Times | Y | Y | N - Quals. J/UJ | Y | Y | Y | Y - TSS samples flagged | Y - Dilutions done out of HT, diluted Aroclors J |
| 4) Initial Calibration Curve (# of standards) | Y - Exceedance 5 pt | Y (validator recalc. results) 5 pt | Y (validator recal. results) 5 pt | Y (validator recalc. results) 5 pt | Y (validator recal. results) 5 pt | Y | Y | Y |
| 5) Calibration Verification Secondary Column | Y - 15% NA | Y - 15% NA | Y - 15% NA | Y - 15% NA | Y - 15% NA | 20% 20% qualitative only | 10% NA | 20% 20% qualitative only |
| 6) Laboratory Blanks | Y | Y | Y | Y | Y | Y | Y | Y |
| 7) Surrogate Recoveries (# required) | NA | NA | NA | NA | NA | Y - Control limits not provided | Y - Control limits not provided | Y/Control limits not provided |
| 8) Matrix Spike (# required) | 75%-125% | 75%-125% | 75%-125% | 75%-125% | 75%-125% | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided |
| 9) Lab Duplicate Lab Control Sample (SRM results?) | Y Y | Used MS/MSD Y (not always performed) - CLs were 75%-125% | Y - Occ. used MS/MSD SDG 12922 >35% Used MS/MSD (75%-125%) | Y - Used MS/MSD Used MS/MSD (80%-120%) | Y Y | Y - MS/MSD control limits not provided Y | Y - Control limits not provided Y | Y - MS/MSD control limits not provided Y - Not addressed |
| 10) Gel Permeation/Forisil Cleanup | NA | NA | NA | NA | NA | Not mentioned | NA | Not mentioned |
| 11) Detection Limit | NA | NA | NA | NA | NA | NA | NA | NA |
| 12) Calc and Transposition Verification (Qualitative verification?) | N | Y | Y - Recalc. | Y - Recalc. | Y - Recalc. | No recalcs. provided; unable to determine if transcription checks were done | No recalcs. provided; unable to determine if transcription checks were done | No recalcs. provided; unable to determine if transcription checks were done |
| 13) Field QC Results | N | Y - OK on rinsate, FD (12812) failed No Action | Y - OK on rinsate, <35% on FD | Y - OK on rinsate, <20% on FD | Y - OK on rinsate, OK on FD | FDs - OK, rinsates had cont. | FDs - OK, rinsates had cont. | FDs - OK |
| 14) Usability Usable/ Supporting Qualifiers | Usable Y - Minor U/J/ flags | Usable Y - Minor quals. due to incorrect ICB calc. | Usable Y - Minor J/UJ flags due to HT exceedances, SDG 12853 also qualified on poor FD values | Usable No quals. | Usable Not apparent if no or some minor quals. | Usable Y - Aroclor 1242 ND based on rinsate cont., UJ extraction errors, J/UJ low surr. %R | Usable - Except some TOC/DOC rejected Y - TOC/DOC R DOC > TOC, all parameters U rinsate, TSS J HT | Usable Y - Aroclor 1242 & 1254 J spectral overlap, J dilutions out of HT, minor CCAL %D |
| 15) Other IC Samples | | | | | | | | |
| SAP | | | | | | | | |
| QAPP | | | | | | | | |
| Lab QAM | | | | | | | | |

Table 3-2 QC Elements for Data Sets Supporting the Fox River RI/FS and RA

| Parameters: Requirements | 1998 Fox River Group (Continued) | | | | | |
|---|---|---|--|---|---|---|
| | PCB Congeners Sediment | Pesticides Sediment | SVOC Sediment | Metals Sediment | TOC/Ammonia Sediment | PCB Fish Tissue |
| SDG #s | Enchem Multiple SDGs | Quanterra Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs |
| Data Review 1) Third-party Validation Performed | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee |
| Deliverables 1) Electronic Deliverables | Y | Y | Y | Y | Y | Y |
| 2) Hardcopy | Y | Y | Y | Y | Y | Y |
| Data Review Details 1) Package Completeness | Y | Y | Y | Y | Y | Y |
| 2) Chain of Custody Procedures | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable |
| 3) Holding Times | Y | Y | Y - 1 missed HT sample J/UJ | Y | Y - Some TOC & ammonia samples J | Y |
| 4) Initial Calibration Curve (# of standards) | Y NA | Y NA | Y NA | Y NA | Y NA | Y NA |
| 5) Calibration Verification Secondary Column | 30% target analytes, 40% internal stds. NA | 20% 20% qualitative only | 20% NA | 10% NA | 10% NA | 20% 20% qualitative only |
| 6) Laboratory Blanks | Y | Y | Y | Y | Y | Y |
| 7) Surrogate Recoveries (# required) | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided |
| 8) Matrix Spike (# required) | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided | Y - Control limits not provided |
| 9) Lab Duplicate Lab Control Sample (SRM results?) | Y - MS/MSD control limits not provided Y | Y - MS/MSD control limits not provided Y | Y - MS/MSD control limits not provided Y | Y - Control limits not provided Y | Y - Control limits not provided Y | Y - MS/MSD control limits not provided Y |
| 10) Gel Permeation/Forisil Cleanup | Not mentioned | Not mentioned | Not mentioned | NA | NA | Not mentioned |
| 11) Detection Limit | NA | NA | NA | NA | NA | NA |
| 12) Calc and Transposition Verification (Qualitative verification?) | No recalcs. provided; unable to determine if transcription checks were done | No recalcs. provided; unable to determine if transcription checks were done | No recalcs. provided; unable to determine if transcription checks were done | No recalcs. provided; unable to determine if transcription checks were done | No recalcs. provided; unable to determine if transcription checks were done | No recalcs. provided; unable to determine if transcription checks were done |
| 13) Field QC Results | None identified | FDs - OK | FDs - OK | FDs - OK | FDs - OK | None identified |
| 14) Usability Usable/ Supporting Qualifiers | Usable Y - 1 compound J/UJ CCAL D, MS/MSD/LCS low %R, poor peak resolution | Usable N | Usable - Except hexachlorocyclopentadiene rejected Y - HCCP R 0% MS/MSD, minor CCAL %D, low surr. %R, & missed HT | Usable Y - BC, low MS %R, RPD | Usable Y - HT | Usable Y - Aroclor 1242 & 1254 J spectral overlap, J/UJ due to extraction error |
| 15) Other IC Samples | | | | | | |
| SAP | | | | | | |
| QAPP | | | | | | |
| Lab QAM | | | | | | |

Table 3-2 QC Elements for Data Sets Supporting the Fox River RI/FS and RA

| Parameters: Requirements | 1992–1993 BBL Deposit A Sediment Data | | | | | 1998–1999 Deposit N Data | | |
|---|--|--|--|---|---|--|--|--|
| | VOA Soil | SVOC Soil | PCB Soil | Pesticides Soil | Metals/CN Soil | PCB Slurry, Soil, Liquid | PCB Congener Slurry, Soil, Liquid | TOC/DOC/TSS Slurry, Soil, Liquid |
| SDG #s | Hazleton 104116 203257 | Hazleton 104116 203242 | Hazleton SDG-1, SDG-2, SDG-3, SDG-4, SDG-5 | Hazleton 104135 203256 | Hazleton BASD34 SD01 BASD08 | Severn Trent VT. Fox9, Fox10, Fox11, Fox12, Fox13, Fox14, Fox16 | Severn Trent VT. Fox9, Fox10, Fox11, Fox12, Fox13, Fox14, Fox16 | WSLH |
| Data Review 1) Third-party Validation Performed | EcoChem | EcoChem | EcoChem | EcoChem | EcoChem | M. A. Kuehl Co. | M. A. Kuehl Co. | M. A. Kuehl Co. |
| Deliverables 1) Electronic Deliverables | Y | Y | Y | Y | Y | Y | Y | Y |
| 2) Hardcopy | Y | Y | Y | Y | Y | Y | Y | Y |
| Data Review Details 1) Package Completeness | Y | Y | Y | Y | Y | Y | Y | Y |
| 2) Chain of Custody Procedures | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable |
| 3) Holding Times | Y | Y | Y | Y | Y | Y - Some exceedances | Y - Some results J/U, some results rejected (> 14 days) | Y - Some exceedances |
| 4) Initial Calibration | Y | Y | Y | Y | Y | Y | Y | Y |
| Curve (# of standards) | Y - As required by method | Y - As required by method | Y - As required by method | Y - As required by method | Y - As required by method | NA | NA | NA |
| 5) Calibration Verification | 20% | 20% | 20% | 20% | 10% | 15% | Y | Y |
| Secondary Column | NA | NA | Y | Y | NA | Y - Some %D exceedances | Y | NA |
| 6) Laboratory Blanks | Y - Tics rejected due to cont. | Y - Tics rejected due to cont. | Y | Y | Y | Y | Y - Some results U based on MB cont. | Y |
| 7) Surrogate Recoveries (# required) | Y | Y | Y | Y | Y | Y | Y | Y |
| 8) Matrix Spike (# required) | Y - No MS/MSD for SDG 203257 J/UJ | Y - No MS/MSD for SDG 203242 J/UJ | Y | Y | Y | Y | Y | Y |
| 9) Lab Duplicate | Y - No MS/MSD for SDG 203257 J/UJ | Y - No MS/MSD for SDG 203242 J/UJ | Y | Y | Y | Y | Y | Y |
| Lab Control Sample (SRM results?) | Y - No LCS for SDG 203257 J/UJ | Y - No LCS for SDG 203242 J/UJ | Y | Y | Y | Y - Some %R outliers | Y - Some %R outliers | Y |
| 10) Gel Permeation/Forisil Cleanup | NA | NA | NA | NA | NA | Not addressed | Not addressed | NA |
| 11) Detection Limit | NA | NA | NA | NA | NA | NA | NA | NA |
| 12) Calc and Transposition Verification (Qualitative verification?) | Y | Y | Y | Y | Y | Y | Y | Y |
| 13) Field QC Results | None identified | None identified | Y | Y | None identified | Y | Y - Some outliers, no quals. assigned | Y - DOC RPD outlier |
| 14) Usability Usable/ Supporting | Usable - Tics rejected due to cont. | Usable - Tics rejected due to cont. | Usable | Usable | Usable | Usable - Some results rejected due to possible cross cont. | Usable - Some results rejected due to exceeded HT | Usable |
| Qualifiers | Y - BC U, Ical RSD, CCAL %D, no LCS MS/MSD TICs rejected due to BC | Y - BC, CCAL %D, Internal std. %R, NO LCS MS/MSD, TICs rejected due to BC | Y - Surr. %R, LCS %R, FD RPD 1242 | Y - RPD between main & confirmation columns NJ | Y - BC, ICV %R CN, MS %R, GFAA post-spike %R | Y - Cooler temps., CCAL %D, HT, LCS %R, dual column %D | Y - HT, cooler temps., CCAL %D, MB cont., LCS %R, over cal | Y - HT, cooler temps., FD RPD, DOC>TOC |
| 15) Other IC Samples | | | | | | | | |
| SAP | | | | | | | | |
| QAPP | | | | | | | | |
| Lab QAM | | | | | | | | |

Table 3-2 QC Elements for Data Sets Supporting the Fox River RI/FS and RA

| Parameters: Requirements | 1998–1999 Deposit N Data (Continued) | | | | | |
|---|--------------------------------------|---|-------------------------------|--|-----------------------|---------------------------|
| | PCB Sludge | PCB Congener Sludge | TOC Sludge | PCB Congener Surface Water | PCB Fish | PCB Congener Minnow |
| SDG #s | Sewern Trent VT. Fox17, Fox18 | Sewern Trent VT. Fox17, Fox18 | Sewern Trent VT. Fox17, Fox18 | WSLH | Sewern Trent VT. Fox7 | WSLH |
| Data Review 1) Third-party Validation Performed | M. A. Kuehl Co. | M. A. Kuehl Co. | M. A. Kuehl Co. | M. A. Kuehl Co. | M. A. Kuehl Co. | M. A. Kuehl Co. |
| Deliverables 1) Electronic Deliverables | Y | Y | Y | Y | Y | Y |
| 2) Hardcopy | Y | Y | Y | Y | Y | Y |
| Data Review Details 1) Package Completeness | Y | Y | Y | Y | Y | Y |
| 2) Chain of Custody Procedures | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable |
| 3) Holding Times | Y | Y | Y | Y | Y | Y |
| 4) Initial Calibration | Y | Y | Y | Y | Y | Y |
| Curve (# of standards) | NA | NA | Y | Y | Y | Y |
| 5) Calibration Verification | Y | Y | Y | Y | Y | Y |
| Secondary Column | Y - %D outliers | Y | NA | Y | Y | Y |
| 6) Laboratory Blanks | Y | Y | Y | Y - Some results U because of MB cont. | Y | Y |
| 7) Surrogate Recoveries (# required) | Y | Y | Y | Y | Y | Y |
| 8) Matrix Spike (# required) | Y | Y - Some %R & RPD outliers | Y | N - Not enough sample | N | Y |
| 9) Lab Duplicate | Y | Y | Y - Some RPD outliers | Y | Y | Y |
| Lab Control Sample (SRM results?) | Y - Some %R outliers | Y | Y - 1 outlier | Y | Y | Y |
| 10) Gel Permeation/Forisil Cleanup | Not addressed | Not addressed | NA | Not addressed | Not addressed | Not addressed |
| 11) Detection Limit | NA | NA | NA | NA | NA | NA |
| 12) Calc and Transposition Verification (Qualitative verification?) | Y | Y | Y | Y | Y | Y |
| 13) Field QC Results | Y | Y - Some outliers, no quals. assigned | Y - Some RPD outliers | Y - Some outliers, no quals. assigned | Y | Y |
| 14) Usability Usable/ Supporting | Usable | Usable | Usable | Usable | Usable | Usable |
| Qualifiers | Y - Dual column %D outliers | Y - CCAL, %D outliers, MS/MSD %R & RPD outliers, LCS %R, over cal | Y - LCS %R, dup. RPD, FD RPD | Y - BC, results <LOQ | N | Y - Reported results <LOQ |
| 15) Other IC Samples | | | | | | |
| SAP | | | | | | |
| QAPP | | | | | | |
| Lab QAM | | | | | | |

4 Analytical and Archive Databases

Electronic data have undergone reduction and standardization and currently reside in both a working database (designed for the internal support of the ongoing RA and RI/FS processes) and the FRDB, complete with user interface.

The development of the FRDB required the data management and manipulation of the source data as described previously. Data were acquired prior to design and development of an appropriate and complete underlying data structure. An outline of the data structure is included in Attachment 1.

The FRDB, designed in Microsoft Access[®], includes available environmental analytical data as well as capacity to store bibliographical information for available reports, research studies, and other documents compiled on the Fox River. The basic structure of the database includes several tables that store the actual data and bibliographical information along with several other “lookup” tables (Attachment 2) and indices that will allow flexibility in searching for information included in the database. The basic table structure and relationships are depicted in Attachment 3. A summary of each table’s function within the database is described as follows:

- **Analytical Table.** This table stores all of the analytical information including fields such as analyte, result, qualifier, etc. This is the core of the analytical data processed and validated by EcoChem. Searches of the database can run on several of the fields contained in this table. This table has relationships with the Analysis Type and Qualifier lookup tables.
- **Data Dictionary Table.** This table contains definitions of the fields used in the Fox River database.
- **Data Set Table.** This table, along with the QA Status Lookup Table listed below, is used to store information regarding the quality assurance or validation level of each of the overall data sets that encompass a sample grouping. A relationship exists with the Document Archive Table that enables reference to a document that exclusively describes a data set.

- **Document Archive Table.** This table contains document and bibliographical information related to Fox River sample data. This table includes information such as the main author's name, additional author names, year of publication or release, subject, title, publication type, keywords and, when available, an abstract of the document and/or a hyperlink to online or electronic copies of the document and associated analytical data. Complete bibliographies from several sources (some not directly related to this project) have been added to this table creating a reference library of over 2,000 sources.
- **Sample Attribute Table.** Information regarding each unique sample is stored in this table. This table has relationships with Data Set and Analytical tables, in addition to six lookup tables. The Deposit, Location, Matrix, Sample Area, Sample Type, and Species lookup tables enable fast and efficient searches of sample attributes.
- **Analysis Type Lookup Table.** This table contains the key data on the type of each analyte in the Analytical Table.
- **Deposit Lookup Table.** This table contains the key data on the named deposit from which a sample was extracted, if a deposit exists for a particular sample.
- **Location Lookup Table.** This table contains the key data on the general location of a sample's origin.
- **Matrix Lookup Table.** This table holds the key data for the matrix type of each sample.
- **QA Status Lookup Table.** The key data on the quality assurance level of each data set contained in the Data Set Table is stored in this table.
- **Qualifier Lookup Table.** This table holds key data on the data qualifier assigned to each analyte in the Analytical Table.
- **Sample Area Lookup Table.** This table contains the key data on more specific locations for sample origins than the Location Table.

- **Sample Type Lookup Table.** This table contains key data on the type or form of each sample that is more specific than that contained in the Matrix Table.
- **Species Lookup Table.** This table contains key data on the common or specific name for a sample and the risk pathway that the sample is associated with. For example, a sample originating from the fish carp is listed under benthic fish for an ecological risk pathway and under food fish for a human health risk pathway.

The FRDB has been customized to include various user interfaces and search capabilities that enable access to the stored data by those who are not familiar with retrieving data from a database application. Help capability and integral database definitions are included. In addition, the database is available via a web server, thus allowing access to the data contained in the database by anyone with Internet capability and a web browser.

Finally, the FRDB is designed with a basic relational structure that will allow data addition in the future as well as the easy migration of the data to other relational database systems. Instructions for importing additional data are included in Attachment 4.

Appendix A

Data Validation Report

Attachment 1

Data Structure Outline

| Table | Fox River Database Field | EcoChem Field | Data Type | Length | Index |
|-------------------------------|--------------------------|-------------------------------------|--------------|--------|--------------|
| <i>Data Set Table</i> | DataSet_ID | Primary key | autonumber | --- | yes, no dups |
| | DataSet | DATASET | text | 50 | yes, no dups |
| | Description | to be added | text | 100 | |
| | QA_Status_ID | foreign key from QA STATUS lookup | long integer | --- | yes |
| | Validator | VALIDATOR | text | 20 | yes |
| <i>QA Status Lookup</i> | QA_Status_ID | Primary key | autonumber | --- | yes, no dups |
| | QA_Status | QASTATUS | text | 15 | yes, no dups |
| | Description | to be added | text | 100 | |
| <i>Sample Attribute Table</i> | SampleAttribute_ID | Primary key | autonumber | --- | yes, no dups |
| | Sample_ID | SAMPID | text | 30 | yes |
| | DataSet_ID | foreign key from DATASET table | long integer | --- | yes |
| | Location_ID | foreign key from LOCATION table | long integer | --- | yes |
| | Deposit_ID | foreign key from DEPOSIT table | long integer | --- | yes |
| | SampleArea_ID | foreign key from SAMPLEAREA table | long integer | --- | yes |
| | BlindID | BLIND_ID | text | 12 | |
| | Depth | DEPTH | text | 14 | |
| | StartDepth | DEPTHFROM | text | 10 | yes |
| | EndDepth | DEPTHTO | text | 10 | yes |
| | DepthUnits | DEPTHUNITS | text | 5 | |
| | CoreGrab | CORE_GRAB | text | 20 | yes |
| | Northing | NORTHING | text | 15 | yes |
| | Easting | EASTING | text | 15 | yes |
| | County | COUNTY | text | 20 | yes |
| | SampleDate | SAMPDATE | text | 10 | yes |
| | SampledBy | SAMPLER | text | 10 | yes |
| | CollectionCompany | COMPANY | text | 30 | yes |
| | DateLabReceived | DATE_RCV | text | 10 | |
| | DateLabExtracted | DATE_EXT | text | 10 | |
| | Matrix_ID | foreign key from MATRIX lookup | long integer | --- | yes |
| | SampleType_ID | foreign key from SAMPLE TYPE lookup | long integer | --- | yes |
| | Species_ID | foreign key from SPECIES lookup | long integer | --- | yes |
| DBTimeStamp | TIMESTAMP | date/time | --- | | |
| <i>Sample Area Lookup</i> | SampleArea_ID | Primary key | autonumber | --- | yes, no dups |
| | SampleArea | LOC_DESC | text | 100 | yes, no dups |

| Table | Fox River Database Field | EcoChem Field | Data Type | Length | Index |
|---------------------------|---------------------------------|----------------------|------------------|---------------|--------------|
| <i>Location Lookup</i> | Location_ID | Primary key | autonumber | --- | yes, no dups |
| | Location | LOCATION | text | 50 | yes, no dups |
| | Description | to be added | text | 100 | |
| <i>Deposit Lookup</i> | Deposit_ID | Primary key | autonumber | --- | yes, no dups |
| | Deposit | DEPOSIT | text | 15 | yes, no dups |
| | Description | to be added | text | 100 | |
| <i>Matrix Lookup</i> | Matrix_ID | Primary key | autonumber | --- | yes, no dups |
| | Matrix | MEDIA | text | 25 | yes, no dups |
| | Description | to be added | text | 50 | |
| <i>Sample Type Lookup</i> | SampleType_ID | Primary key | autonumber | --- | yes, no dups |
| | SampleType | SAMPLETYPE | text | 30 | yes, no dups |
| | Description | to be added | text | 50 | |
| <i>Species Lookup</i> | Species_ID | Primary key | autonumber | --- | yes |
| | CommonName | SPECIES | text | 30 | yes, no dups |
| | EcoRisk | GROUP | text | 20 | same index |
| | HHRisk | GROUP2 | text | 20 | same index |
| | Species | TRUESPECIES | text | 20 | |

| Table | Fox River Database Field | EcoChem Field | Data Type | Length | Index |
|-------------------------|--------------------------|---|--------------|--------|--------------|
| <i>Analytical Table</i> | Analytical_ID | Primary key | autonumber | --- | yes |
| | SampleAttribute_ID | foreign key from SAMPLE ATTRIBUTE table | text | 30 | yes |
| | Analyte | ANALYTE | text | 50 | yes |
| | Result | RESULT | text | 15 | yes |
| | Qualifier | foreign key from QUALIFIER lookup | text | 6 | yes |
| | Units | UNITS | text | 15 | |
| | AnalysisType_ID | foreign key from ANALYSIS TYPE table | long integer | --- | yes |
| | ReportingBasis | BASIS | text | 20 | |
| | SDG | SDG | text | 10 | |
| | DetectionLimit | DETLIMIT | text | 15 | |
| | Aliquot | ALQUOT | text | 10 | |
| | Method | METHOD | text | 20 | yes |
| | LabID | LABID | text | 15 | |
| | AnalyteOld | ANALYTEOLD | text | 50 | |
| | ResultOld | RESULTOLD | text | 50 | |
| | QualifierOld | QUALOLD | text | 6 | |
| | Comments | COMMENT | text | 110 | |
| | Lab | LAB | text | 20 | yes |
| ImportFile | IMPORTFILE | text | 15 | | |
| Source | SOURCE | text | 100 | yes | |
| <i>Qualifier Lookup</i> | Qualifier | QUAL (primary key) | text | 6 | yes, no dups |
| | Description | to be added | text | 50 | |

| Table | Fox River Database Field | EcoChem Field | Data Type | Length | Index |
|-----------------------------|--------------------------|--------------------------------|--------------|--------|--------------|
| <i>Document Archive</i> | Document_ID | Primary key | autonumber | --- | yes, no dups |
| | DataSet_ID | foreign key from DATASET table | long integer | --- | yes, no dups |
| | Author | | text | 200 | |
| | Year | | text | 4 | |
| | Title | | text | 255 | |
| | SecondaryTitle | | text | 150 | |
| | Journal | | text | 75 | |
| | Volume | | text | 3 | |
| | Issue | | text | 10 | |
| | Pages | | text | 10 | |
| | AlternateJournal | | text | 75 | |
| | CallNumber | | text | 25 | |
| | Label | | text | 20 | |
| | Keywords | | text | 225 | |
| | Abstract | | memo | --- | |
| | Notes | | text | 40 | |
| | City | | text | 20 | |
| | Institution | | text | 75 | |
| | Date | | text | 20 | |
| | Publisher | | text | 50 | |
| | SeriesEditor | | text | 35 | |
| | SeriesTitle | | text | 100 | |
| | Edition | | text | 5 | |
| | Newspaper | | text | 75 | |
| | ConferenceLocation | | text | 50 | |
| | ConferenceYear | | text | 4 | |
| | ConferenceName | | text | 50 | |
| | AcademicDepartment | | text | 50 | |
| | University | | text | 30 | |
| | Programmer | | text | 40 | |
| Cartographer | | text | 40 | | |
| Scale | | text | 20 | | |
| AccessYear | | text | 4 | | |
| AccessDate | | text | 25 | | |
| <i>Analysis Type Lookup</i> | AnalysisType_ID | Primary key | autonumber | --- | yes, no dups |
| | AnalysisType | METHODTYPE | text | 15 | yes, no dups |
| <i>Data Dictionary</i> | Field | Primary key | text | 30 | yes, no dups |
| | Description | to be added | text | 150 | |

Attachment 2

Lookup Tables

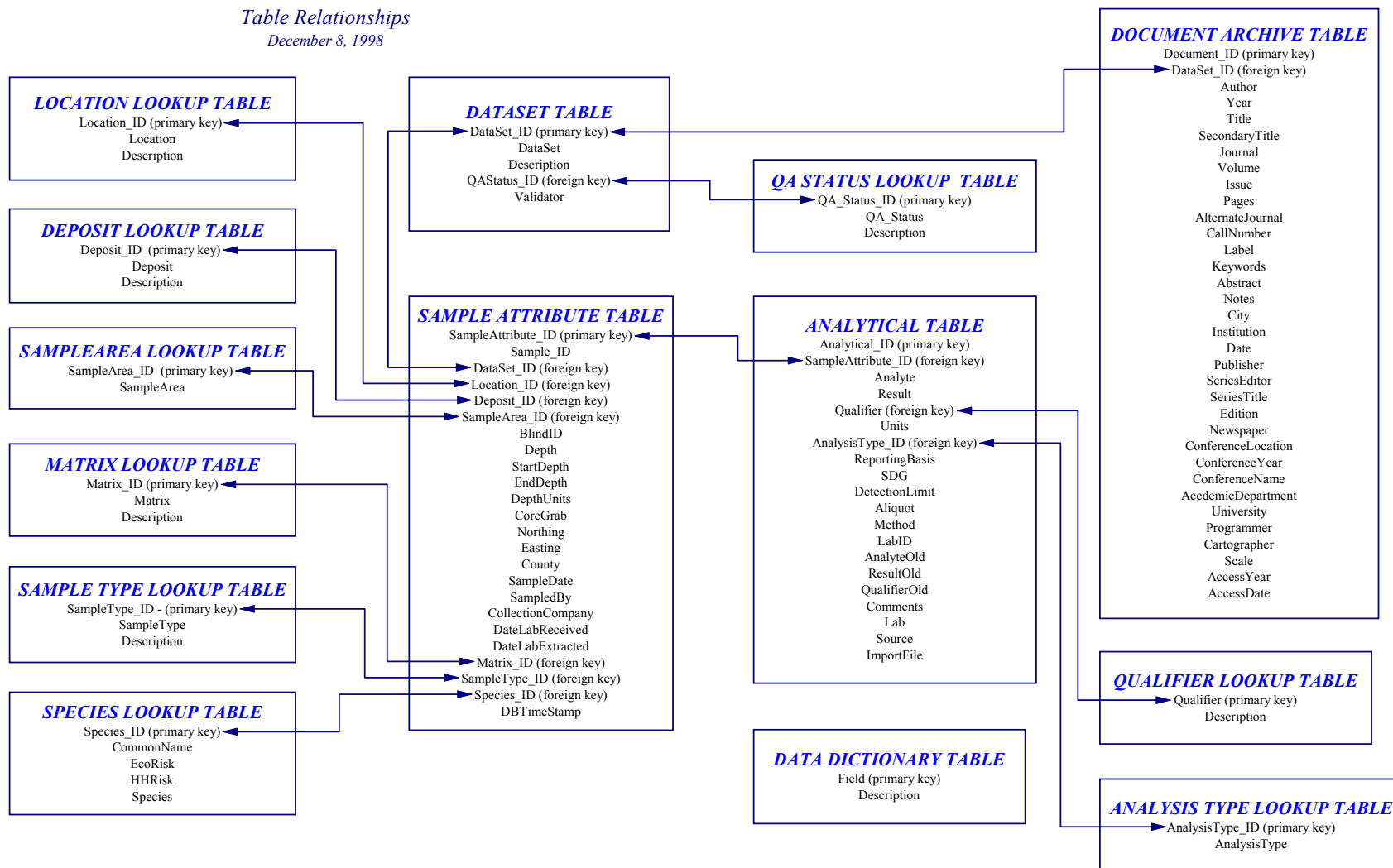
Local Lookup Tables and Queries for Fox River Database Forms.mdb File (Table 1)

| Table Name | Query to Populate the Table | Forms Using the Table |
|---------------------------------------|--|--|
| tblLookup_CriteriaForLists | None – static table (DO NOT ALTER) | frmDataList |
| tblLookup_FieldsForLists | None – static table (DO NOT ALTER) | frmDataList |
| tblLookup_SortFieldsForSearches | None – static table (DO NOT ALTER) | frmDataSearch |
| tblLookup_Unique_AnalysisType | Append tblLookup_Unique_AnalysisType | frmDataList |
| tblLookup_Unique_Analyte | Append tblLookup_Unique_Analyte | frmDataList, frmDataSearch, frmStatistic |
| tblLookup_Unique_CollectionCompany | Append tblLookup_Unique_CollectionCompany | frmDataList |
| tblLookup_Unique_CommonName | Append tblLookup_Unique_CommonName | frmDataList |
| tblLookup_Unique_CoreGrab | Append tblLookup_Unique_CoreGrab | frmDataList |
| tblLookup_Unique_County | Append tblLookup_Unique_County | frmDataList |
| tblLookup_Unique_DataSet | Append tblLookup_Unique_DataSet | frmDataSearch |
| tblLookup_Unique_Deposit | Append tblLookup_Unique_Deposit | frmDataList |
| tblLookup_Unique_EcoRisk | Append tblLookup_Unique_EcoRisk | frmDataList |
| tblLookup_Unique_EcoRiskAndCommonName | Append tblLookup_Unique_EcoRiskAndCommonName | frmDataSearch |
| tblLookup_Unique_HHRisk | Append tblLookup_Unique_HHRisk | frmDataList |
| tblLookup_Unique_HHRiskAndCommonName | Append tblLookup_Unique_HHRiskAndCommonName | frmDataSearch |
| tblLookup_Unique_Lab | Append tblLookup_Unique_Lab | frmDataList |
| tblLookup_Unique_Location | Append tblLookup_Unique_Location | frmDataList |
| tblLookup_Unique_LocationAndDeposit | Append tblLookup_Unique_LocationAndDeposit | frmDataSearch |
| tblLookup_Unique_Matrix | Append tblLookup_Unique_Matrix | frmDataList |
| tblLookup_Unique_MatrixAndSampleType | Append tblLookup_Unique_MatrixAndSampleType | frmDataSearch |
| tblLookup_Unique_Method | Append tblLookup_Unique_Method | frmDataList |
| tblLookup_Unique_QAStatus | Append tblLookup_Unique_QAStatus | frmDataList |
| tblLookup_Unique_Qualifier | Append tblLookup_Unique_Qualifier | frmDataSearch |
| tblLookup_Unique_SampledBy | Append tblLookup_Unique_SampledBy | frmDataList |
| tblLookup_Unique_SampleID | Append tblLookup_Unique_SampleID | frmDataList |
| tblLookup_Unique_SampleType | Append tblLookup_Unique_SampleType | frmDataList |
| tblLookup_Unique_Source | Append tblLookup_Unique_Source | frmDataList |
| tblLookup_Unique_StatisticsChoices | Append tblLookup_Unique_StatisticsChoices | frmStatistic |
| tblLookup_Unique_Validator | Append tblLookup_Unique_Validator | frmDataList |

FOX RIVER DATABASE

Table Relationships

December 8, 1998

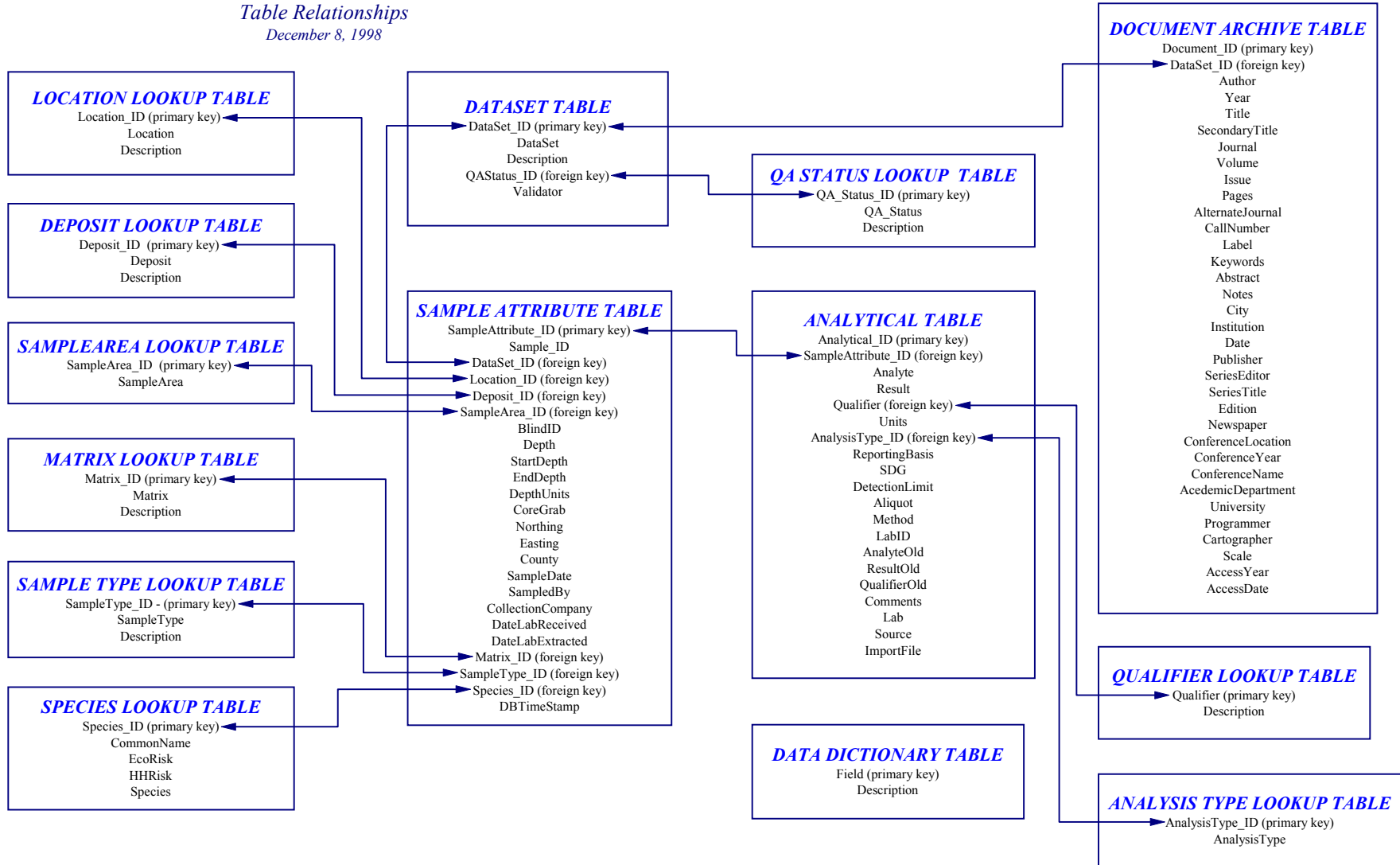


Attachment 3

Table Structure and Relationships

FOX RIVER DATABASE

Table Relationships
December 8, 1998



Attachment 4

Data Importing Instructions

I. Importing Data to the Fox River Database for the First Time (empty database):

Steps for the FoxRiverData.mdb Database File:

1. Import raw data to a new table called SAMPLES in the Fox River Data Tables database. Fields in this import table should be named as below (names in parentheses are the actual database field names). All fields should be of text data type except for TIMESTAMP, which should be of date/time type. TIMESTAMP should be left blank in the import file because a date/time value is added when the data is entered into the database.

- | | | | |
|----|--------------------------------|-----|-----------------------------|
| a. | SAMPID (Sample_ID) | x. | SPECIES (CommonName) |
| b. | ANALYTE (Analyte) | y. | ALQUOT (Aliquot) |
| c. | RESULT (Result) | z. | METHODTYPE (AnalysisType) |
| d. | QUAL (Qualifier) | aa. | METHOD (Method) |
| e. | UNITS (Units) | bb. | BLIND_ID (BlindID) |
| f. | SAMPDATE (SampleDate) | cc. | SAMPLER (SampledBy) |
| g. | MEDIA (Matrix) | dd. | COMMENT (Comments) |
| h. | LABID (LabID) | ee. | DEPOSIT (Deposit) |
| i. | DATE_RCV (DateLabReceived) | ff. | NORTHING (Northing) |
| j. | DATE_EXT (DateLabExtracted) | gg. | EASTING (Easting) |
| k. | DETLIMIT (DetectionLimit) | hh. | GROUP (EcoRisk) |
| l. | SDG (SDG) | ii. | GROUP2 (HHRisk) |
| m. | IMPORTFILE (ImportFile) | jj. | COREGRAB (CoreGrab) |
| n. | SOURCE (Source) | kk. | ANALYTEOLD (AnalyteOld) |
| o. | DATASET (DataSet) | ll. | LOC_DESC (SampleArea) |
| p. | LAB (Lab) | mm. | SAMPLETYPE (SampleType) |
| q. | VALIDATOR (Validator) | nn. | COUNTY (County) |
| r. | QASTATUS (QA_Status) | oo. | RESULTOLD (ResultOld) |
| s. | LOCATION (Location) | pp. | QUALOLD (QualifierOld) |
| t. | DEPTH (Depth) | qq. | TRUESPECIES (Species) |
| u. | DEPTHFROM (StartDepth) | rr. | COMPANY (CollectionCompany) |
| v. | DEPTHTO (EndDepth) | ss. | BASIS (ReportingBasis) |
| w. | DEPTHUNITS (DepthUnits) | tt. | TIMESTAMP (DBTimeStamp) |

2. Run qryTimeStamp_ImportFile to date/time stamp the entry of new samples into the database. This allows for easier importing of new samples in the future as well as keeping a record of when samples were first entered into the database.
3. Populate lookup tables by running the these queries in the exact order listed below:
 - a. qryPopulate_Unique_AnalysisType
 - b. qryPopulate_Unique_QAStatus
 - c. qryPopulate_Unique_DataSet
 - d. qryPopulate_Unique_Deposit
 - e. qryPopulate_Unique_Location
 - f. qryPopulate_Unique_Matrix
 - g. qryPopulate_Unique_Qualifier
 - h. qryPopulate_Unique_SampleArea
 - i. qryPopulate_Unique_SampleType
 - j. qryPopulate_Unique_Species
4. Run qryPopulate_Unique_SampleAttribute to populate tblSampleAttribute.
5. Run qryPopulate_Unique_Analytical to populate tblAnalytical.
6. Run qryPopulate_tblDocumentArchive_WithDataSets to populate DataSet_ID field in tblDocumentArchive with DataSet_IDs from tblDataSet.

Steps for the Fox River Database Forms.mdb Database File:

- I. Run the queries listed in Table 1 to populate the local lookup tables. The queries must be run in the order that they are listed in Table 1. The first three database tables listed in Table 1 are static tables and should never be altered.
- II. **Subsequent Importing of Data to the Fox River Database (populated database):**
 1. To import additional data to the Fox River Database after the database has been filled initially, follow the same steps as outlined above for entering data into the FoxRiverData.mdb file. The lookup tables have indexed fields to prevent entry of duplicate data. When the lookup queries are run and you are trying to enter duplicate data, Access® will show an error message that some data will not be added due to key violations. Choose the option to run the query anyway, and only the new data will be added to the database.

2. After the new data has been added, you must change the lookup tables in the Fox River Database Forms.mdb file. Open the database lookup tables listed in Table 1 and delete all records in each table. After all data has been deleted from all lookup tables, run the Table 1 queries in the order listed to repopulate the lookup tables with the updated database data.
3. The updated Fox River Database Forms.mdb must then be distributed to all users. Replace the old copy of the file with the updated version.

III. Populating the Fox River Web Database File (Fox River Web DB.mdb):

1. For first time populating of data to the web database file (empty database), import the following tables from the respective Access® database files created above:

FoxRiverData.mdb: tblAnalysisType
 tblAnalytical
 tblDataDictionary
 tblDataSet
 tblDeposit
 tblDocumentArchive
 tblLocation
 tblMatrix
 tblQA_Status
 tblQualifier
 tblSampleArea
 tblSampleAttribute
 tblSampleType
 tblSpecies

Fox River Database Forms.mdb: tblLookup_CriteriaForLists
 tblLookup_FieldsForLists
 tblLookup_SortFieldsForSearches
 tblLookup_Unique_AnalysisType
 tblLookup_Unique_Analyte
 tblLookup_Unique_CollectionCompany
 tblLookup_Unique_CommonName
 tblLookup_Unique_CoreGrab
 tblLookup_Unique_County
 tblLookup_Unique_DataSet
 tblLookup_Unique_Deposit

tblLookup_Unique_EcoRisk
tblLookup_Unique_EcoRiskAndCommonName
tblLookup_Unique_HHRisk
tblLookup_Unique_HHRiskAndCommonName
tblLookup_Unique_Lab
tblLookup_Unique_Location
tblLookup_Unique_LocationAndDeposit
tblLookup_Unique_Matrix
tblLookup_Unique_MatrixAndSampleType
tblLookup_Unique_Method
tblLookup_Unique_QAStatus
tblLookup_Unique_Qualifier
tblLookup_Unique_SampledBy
tblLookup_Unique_SampleID
tblLookup_Unique_SampleType
tblLookup_Unique_Source
tblLookup_Unique_StatisticsChoices
tblLookup_Unique_Validator

2. When new data is imported into the Access database as above, you must repopulate the web database file to reflect the new data. To do this, delete all tables in the Fox River Web DB.mdb file except for the static tables listed below. After the tables have been deleted, compact the database file to clear the deleted tables file space. Then, import all tables as described in Step 1 above.

tblLookup_CriteriaForLists
tblLookup_FieldsForLists
tblLookup_SortFieldsForSearches

**Addendum 1 to the Data Management Summary Report
(EcoChem, 2002)**

DATA MANAGEMENT SUMMARY REPORT, ADDENDUM 1
FOX RIVER REMEDIAL INVESTIGATION/FEASIBILITY STUDY

November 25, 2002

ADDENDUM 1 TO THE DATA MANAGEMENT SUMMARY REPORT

Note: As data are collected, reviewed (or validated), and appended to the Fox River Database (FRDB), the Data Management Summary Report will also be appended. A description of the data set, along with results of data review/validation and determination of usability will be discussed in consecutively numbered sections.

As supporting tables (Table 3-1: Data Set Analysis and Table 3-2: QC Elements for Data Sets Supporting the Fox River RI/FS and RA) are appended, the tables will be resubmitted (with each Addendum) in their entirety.

3.2.29 1999 DEMONSTRATION PROJECT DATA - SMU 56/57

This data set has now been appended to the Fox River Database (FRDB) and has been included in Tables 3-1, Data Set Analysis. All previous discussion remains valid, as presented in the DMR, October, 2000.

3.2.30 2000/2001 FRG/CH₂M HILL SEDIMENT & WOOD CHIP DATA

CH₂M Hill collected soil/sediment (and one set of wood chip) samples in 2000 and 2001 for the Fox River Group (FRG). The samples were collected from the Little Lake Butte des Morts area. Samples were analyzed for polychlorinated biphenyl (PCB) Aroclors, metals, volatile organics, semivolatile organics, gasoline- and diesel-range organics, and cyanide. The data set consisted of 428 samples.

EcoChem performed a review of the FRG 2000 and 2001 data validation conducted by CH₂M Hill. EcoChem evaluated the validation results for completeness and technical agreement. The samples were analyzed by United States Environmental Protection Agency (EPA) SW-846 methodology and other miscellaneous EPA methods. The gasoline- and diesel-range analyses were conducted using the Wisconsin GRO and DRO methods. The validation protocols used by CH₂M Hill were not specified.

Overall the data are of acceptable quality. The samples appear to have analyzed as per the cited methods, and the validation worksheets generally follow the guidelines specified in *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994) and *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994). No validation reports were provided. The information reviewed consisted of data validation worksheets and annotated sample result summary forms. The validation worksheets were often not complete. However, there is sufficient information in the notes made by the validator (in the worksheet comments section) to indicate that the data were reviewed, and the issue is one of incomplete documentation, rather than an incomplete review. Most of the worksheets do not include the date that the validation was performed, or the name of the validator. Some of the sample result summary forms were also not dated.

Many of the data qualifiers issued by CH₂M Hill were due to interference caused by the natural overlap of some of the Aroclors (such as Aroclors 1242 and 1254). It is not

possible to evaluate these findings without reviewing the raw data. A more detailed review of the data may result in the removal of some of these qualifiers. For the semivolatile analyses in data package 913426, the qualifiers on the sample result summary forms do not match those discussed in the validation worksheet. A more detailed review of the data for this package would result in additional qualifiers (estimated data). However, the above changes would not significantly impact the reported data. As determined by this review, the data, as qualified, are usable for the intended purpose.

3.2.31 2000 FRG/BBL SUPPLEMENTAL MONITORING PROGRAM DATA: SURFACE WATER

Blasland Bouck & Lee (BBL) collected surface water, particulate, and XAD filter samples in 2000 for the FRG. The samples were collected as part of the Supplemental Monitoring Program – Surface Water. Samples were analyzed for PCB Aroclors, PCB congeners, total suspended solids (TSS), total volatile suspended solids (TVSS), and total organic carbon (TOC). The data set consisted of 205 samples. Not all samples were analyzed for all tests.

EcoChem performed a review of the FRG 2000 data validation conducted by BBL. EcoChem evaluated the validation worksheets and reports for completeness and technical agreement. The samples were analyzed by EPA SW-846 methodology and other miscellaneous EPA methods. The validation report states that the qualifiers are “in accordance with National Functional Guidelines.” The date of the version of Functional Guidelines used is not provided. The validation worksheets do not provide the name(s) of the validator(s), or the date that the validation was performed. The sample result summary forms are usually not initialed and dated.

The samples appear to have analyzed as per the cited methods, and the validation worksheets generally follow the guidelines specified in *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994) and *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994).

For one PCB congener data package, when qualifiers were recommended (in the validation worksheet) based on blank contamination, the sample result summary forms were not qualified. Rather, the reporting limits were elevated, but no “U” qualifier was added to the summary form. During a more detailed review, EcoChem would add the qualifiers. Although surrogate and laboratory control sample (LCS) recovery outliers were noted, no action was taken. A more detailed review of the data would most likely result in additional qualifiers (estimated data). Overall the data are of acceptable quality. The data, as qualified, are usable for the intended purpose.

3.2.32 2000/2001 FRG/BBL SUPPLEMENTAL MONITORING PROGRAM DATA: SEDIMENT DATA

BBL collected sediment samples in 2000 and 2001 for the FRG. The samples were collected as part of the Supplemental Monitoring Program. Samples were analyzed for PCB congeners (one data set), PCB Aroclors, TOC, and grain size. The data set consisted of 158 samples.

EcoChem performed a review of the FRG 2001 data validation conducted by BBL. EcoChem evaluated the validation worksheets and reports for completeness and technical agreement. The samples were analyzed by EPA SW-846 methodology and other miscellaneous EPA methods. The validation report states that the qualifiers are “in accordance with National Functional Guidelines.” The date of the version of Functional Guidelines used is not provided. The validation worksheets do not provide the name(s) of the validator(s), or the date that the validation was performed. The sample result summary forms are usually not initialed and dated.

The samples appear to have analyzed as per the cited methods, and the validation worksheets generally follow the guidelines specified in *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994) and *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994). Only sample results were provided for the grain size analyses, so these were not validated.

Overall the data are of acceptable quality. Qualifiers were issued based on a matrix spike recovery outlier. However, the associated matrix spike duplicate and LCS were acceptable. A more detailed review of the data would most likely result in removal of the qualifiers. With this change, no data would be qualified. The data are usable for the intended purpose.

3.2.33 2001 FRG/BBL GREEN BAY SEDIMENT SAMPLING DATA

BBL collected sediment samples in 2001 for the FRG. The samples were collected as part of the Green Bay Sediment Sampling event. Samples were analyzed for PCB Aroclors, TOC, and grain size. The data set consisted of 30 samples.

EcoChem performed a review of the FRG 2001 data validation conducted by BBL. EcoChem evaluated the validation worksheets and reports for completeness and technical agreement. The samples were analyzed by EPA SW-846 methodology and other miscellaneous EPA methods. The validation report states that the qualifiers are “in accordance with National Functional Guidelines.” The date of the version of Functional Guidelines used is not provided. The validation worksheets do not provide the name(s) of the validator(s), or the date that the validation was performed. The sample result summary forms are usually not initialed and dated.

Overall the data are of acceptable quality. The samples appear to have analyzed as per the cited methods, and the validation worksheets generally follow the guidelines specified

in *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994) and *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994). Only sample results were provided for the grain size analyses, so these were not validated.

In addition to evaluating the validation reports and worksheets, EcoChem also performed a full validation of the data packages. The results of the validation by EcoChem were compared to the validation performed by BBL. The two validations were mostly in agreement; however, BBL estimated a few TOC results and EcoChem did not. The changes would not significantly impact the reported data. As determined by this review, the data, as qualified, is usable for the intended purpose.

3.2.34 2001 FRG/BBL WATER COLUMN-HIGH FLOW DATA

BBL collected surface water, particulate, and XAD filter samples in 2001 for the FRG. The samples were collected as part of the Fox River 2001 Water Column – High Flow study. Samples were analyzed for PCB Aroclors, PCB congeners, TSS, TVSS, and TOC. The data set consisted of 615 samples. Not all samples were analyzed for all tests.

EcoChem performed a review of the FRG 2001 data validation conducted by BBL. EcoChem evaluated the validation worksheets and reports for completeness and technical agreement. The samples were analyzed by EPA SW-846 methodology and other miscellaneous EPA methods. The validation report states that the qualifiers are “in accordance with National Functional Guidelines.” The date of the version of Functional Guidelines used is not provided. The validation worksheets do not provide the name(s) of the validator(s), or the date that the validation was performed. The sample result summary forms are usually not initialed and dated.

The samples appear to have analyzed as per the cited methods, and the validation worksheets generally follow the guidelines specified in *U.S. EPA National Functional Guidelines for Organic Data Review* (February 1994) and *U.S. EPA National Functional Guidelines for Inorganic Data Review* (February 1994).

Many of the surrogate recovery values were less than the acceptance limit and less than 10 percent for the PCB Aroclor analyses. The validation reports state that this was caused by the Florisil cleanup. The reports further state that the Florisil had a negative impact on select peaks (typically Aroclor 1242), and that the results for the affected Aroclors were recalculated using non-impacted peaks. On the sample result summary forms, the reported value was lined out and a revised (elevated) concentration was hand entered.

It is not possible to evaluate the revisions without the raw data. Also, none of the calculations were provided, and so cannot be verified. During a more detailed review of the data, EcoChem would most likely estimate the data. If revised concentrations were appropriate, EcoChem would request that the laboratory recalculate the concentrations and issue a revised sample result summary form.

For the PCB congener analyses, no changes or additional qualifiers are recommended by EcoChem. However, when qualifiers were issued based on blank contamination, the sample result summary forms were not qualified as recommended. Rather, the reporting limits were elevated, but no “U” qualifier was added to the summary form. During a more detailed review, EcoChem would add the qualifiers. For the general chemistry parameters (TSS, TVSS, and TOC), no changes or additional qualifiers are recommended by EcoChem. A more detailed review of the data would most likely not result in additional qualifiers. The data, as qualified, are usable for the intended purpose.

3.3 DATA USABILITY

3.3.1 FULLY VALIDATED DATA

The following data sets have been validated by an independent party and are considered useable, as qualified:

- 1994 GAS/SAIC Sediment Collection
- 1994 Woodward-Clyde Deposit A Sediment Collection
- 1995 WDNR Sediment Data Collection
- 1996 USFWS NRDA Fish Tissue Data Collection
- 1996 WDNR Fish Tissue Data Collection
- 1998 Demonstration Project Data - SMU 56/57
- 1998 RETEC RI/FS Supplemental Data Collection
- 1996 FRG/BBL Sediment/Tissue Data Collection
- 1997 Demonstration Project Data - Deposit N
- 1992/93 BBL Deposit A Sediment Data Collection
- 1998 FRG/Exponent Data Collection
- 1998 FRG/Blasland, Bouck, and Lee, Inc. Sediment/Tissue Data Collection
- 1998 Deposit N Pilot Remediation-Pre-Dredge, Post-Dredge, Operation Monitoring, and Environmental Monitoring Data
- 1999 Demonstration Project Data- SMU 56/57
- State of Michigan Fish Consumption Advisory Data
- Lake Michigan Tributary Monitoring Data
- 1999 Demonstration Project Data - SMU 56/57
- Minergy EPA SITE Program Data
- 2000/2001 FRG/CH2M Hill Sediment & Wood Chip Data;

- 2000 FRG/BBL Supplemental Monitoring Program Data: Surface Water;
- 2000/2001 FRG/BBL Supplemental Monitoring Program Data: Sediment Data;
- 2001 FRG/BBL Green Bay Sediment Sampling Data; and
- 2001 FRG/BBL Water Column-High Flow Data.

Although the data sets (listed above) were found to be validated and useable, it must be stressed that there were individual data points that were rejected. These rejected data points have not been used in support of the RI/FS or RA.

3.3.2 SUPPORTING DATA

The following data sets have not been validated and, in general, should be used only as supporting data. The data have been collected within different programs and with different data quality objectives therefore, varying degrees of supporting documentation may be available.

- 1989/90 Fox River Mass Balance Study
- 1989/90 Green Bay Mass Balance Study (GLNPO)
- 1993 Triad Assessment
- 1993-1996 USFWS Tree Swallow Data Collection
- 1994-1995 Cormorant Data Collection
- 1997 USFWS NRDA Waterfowl Tissue Data Collection
- 1997 WDNR Caged Fish Bioaccumulation Study Data
- Fox River Fish Consumption Advisory Data
- Stromberg Eagle Data Collection
- USGS NAWQA Data
- WDNR Wildlife Tissue Data
- WPDES Permit Influent Data
- Lake Michigan Mass Balance Data
- Minergy Mineralogical Data
- Lower Fox River Background Metals Assessment
- FoxView Data

3.3.3 INDETERMINATE DATA

The following data sets have not been validated and have not been subjected to a data quality review. This is due to complete lack of supporting QA/QC documentation; or,

EcoChem did not receive the hardcopy data and documents by the date of this report. At this time the overall quality of these data sets is unknown and the data should be used with that fact in mind.

- Ankley and Call

**Table 3-1
Data Set Analysis**

| Data Source | Number of Samples | Matrices ¹ | Analyses Conducted ² | Number of Records | Number of Files in Delivery | File Type | Report Section | Earliest Year of Collection | Latest Year of Collection |
|---|-------------------|-----------------------|--|-------------------|-----------------------------|-------------|----------------|-----------------------------|---------------------------|
| 1989 - 1990 Fox River Mass Balance Study | 1967 | S,W | PCB-A, PCB-C, W | 25457 | 6 | Spreadsheet | 3.2.01 | 1989 | 1990 |
| 1989 - 1990 Green Bay Mass Balance Study (GLNPO) | 2069 | S,T,W | B, PCB-C, W | 201701 | 92 | Database | 3.2.01 | 1987 | 1990 |
| 1992 - 1993 BBL Deposit A Sediment Data | 117 | S,W | M, P/H, PCB-A, SVOA, V, W | 1094 | 1 | Spreadsheet | 3.2.02 | 1992 | 1993 |
| 1993 Triad Assessment | 27 | S | B, M, P/H, PCB-A, SVOA, W | 631 | 11 | Spreadsheet | 3.2.03 | 1992 | 1993 |
| 1994 GAS/SAIC Sediment Collection | 253 | S | DXN, M, P/H, PCB-A, SVOA, V, W | 5654 | 6 | Spreadsheet | 3.2.04 | 1994 | 1994 |
| 1995 WDNR Sediment Data | 488 | S | M, PCB-A, W | 6433 | 8 | Spreadsheet | 3.2.05 | 1995 | 1995 |
| 1996 FRG/BBL Sediment/Tissue Data | 25 | S,T | B, PCB-C, W | 2771 | 6 | Spreadsheet | 3.2.06 | 1996 | 1996 |
| 1995 - 1996 WDNR Tissue Data | 200 | T | B, PCB-A, W | 1673 | 1 | Spreadsheet | 3.2.07 | 1995 | 1996 |
| 1996 - USFWS NRDA Tissue Data | 376 | T | DXN, P/H, PCB-A, PCB-C, W | 16017 | 5 | Spreadsheet | 3.2.08 | 1996 | 1999 |
| 1993-1996 Tree Swallow Data | 200 | T | B, DXN, P/H, V, W | 5429 | 2 | Database | 3.2.09 | 1993 | 1993 |
| 1994-1995 Cormorant Data | 193 | T | B, DXN, P/H, PCB-C, W | 6178 | 2 | Database | 3.2.09 | 1994 | 1995 |
| 1997 USFWS NRDA Waterfowl Tissue Data | 70 | T | B, P/H, PCB, V, W | 1680 | 2 | Database | 3.2.09 | 1997 | 1997 |
| Fox River Fish Consumption Advisory Data: 1998 WDNR Fish Consumption Data | 130 | T | B,M, PCB-A, W | 777 | 1 | ASCII | 3.2.10 | 1998 | 1998 |
| Fox River Fish Consumption Advisory Data | 1766 | S,T | B, DXN, M, P/H, PCB-A, PCB-C, SVOA, V, W | 11620 | 2 | ASCII | 3.2.10 | 1971 | 1996 |
| WDNR Wildlife Tissue Data | 417 | T | B, M, P/H, PCB-A | 2532 | 3 | Database | 3.2.11 | 1984 | 1996 |
| Lake Michigan Tributary Monitoring Data | 88 | W | M, P/H, PCB-C, V | 5722 | 5 | Spreadsheet | 3.2.12 | 1994 | 1995 |
| Stromberg Eagle Data | 31 | T | B, DXN, P/H, PCB-A, PCB-C, SVOA, V, W | 954 | 1 | ASCII | 3.2.13 | 1991 | 1996 |
| USGS NAWQA Data | 441 | S,T,W | B, M, P/H, PCB, SVOA, V, W | 11879 | 21 | Spreadsheet | 3.2.14 | 1992 | 1997 |
| 1994 Woodward-Clyde Deposit A Sediment Data | 66 | S | PCB-A, W | 585 | 12 | Spreadsheet | 3.2.15 | 1994 | 1994 |
| WPDES Permit Influent Data | 8 | W | B, DXN, M, P/H, PCB-A, RAD, SVOA, V, W | 847 | 1 | Spreadsheet | 3.2.16 | 1993 | 1997 |
| Lower Fox River Background Metals Assessment Data | 14 | W | M | 78 | 1 | Spreadsheet | 3.2.17 | 1991 | 1993 |
| 1997 WDNR Caged Fish Bioaccumulation Study Data | 25 | S,T | B, PCB-C, W | 1672 | 2 | Spreadsheet | 3.2.18 | 1997 | 1997 |
| 1997 Demonstration Project Data - Deposit N | 10 | S | M, PCB, W | 83 | 1 | Spreadsheet | 3.2.19 | 1997 | 1997 |
| 1997 Demonstration Project Data - SMU 56/57 | 295 | S,W | DXN, M, P/H, PCB-A, SVOA, V, W | 3114 | 12 | Spreadsheet | 3.2.20 | 1997 | 1998 |
| 1998 RETEC RI/FS Supplemental Data | 252 | S,T | B, DXN, M, P/H, PCB-A, PCB-C, SVOA, V, W | 10781 | 1 | ASCII | 3.2.21 | 1998 | 1998 |
| Lake Michigan Mass Balance Data | 6987 | A,S,T,W | M, P/H, PCB-C, V, W | 91621 | 211 | Database | 3.2.22 | 1993 | 1996 |
| Minergy Mineralogical Data | 15 | S | W | 219 | 1 | Spreadsheet | 3.2.23 | 1995 | 1999 |
| 1998 FRG/Exponent Data | 225 | T | B, M, P/H, PCB-A, PCB-C, W | 17708 | 3 | Database | 3.2.24 | 1998 | 1998 |

**Table 3-1
Data Set Analysis**

| Data Source | Number of Samples | Matrices ¹ | Analyses Conducted ² | Number of Records | Number of Files in Delivery | File Type | Report Section | Earliest Year of Collection | Latest Year of Collection |
|---|-------------------|-----------------------|---------------------------------------|-------------------|-----------------------------|--------------------------|----------------|-----------------------------|---------------------------|
| 1998 FRG/BBL Sediment/Tissue Data | 1315 | S,T,W | B, M, P/H, PCB-A, PCB-C, RAD, SVOA, W | 18824 | 1 | Database | 3.2.25 | 1998 | 1998 |
| 1998 - 1999 Deposit N Data: Post-Dredge | 43 | S | PCB-A, PCB-C, W | 690 | 8 | Spreadsheet | 3.2.26 | 1999 | 1999 |
| 1998 - Deposit N Data: Pre-Dredge | 53 | S | PCB-A, PCB-C, W | 1437 | 6 | Spreadsheet | 3.2.26 | 1998 | 1998 |
| 1998/1999 Deposit N Data: Remediation | 197 | T,W | PCB-C, W | 10264 | 1 | Spreadsheet | 3.2.26 | 1998 | 1999 |
| 1998 - 1999 Deposit N Data: Operational Monitoring | 12 | S | M, PCB-A, W | 123 | 1 | Spreadsheet | 3.2.26 | 1998 | 1998 |
| Ankley and Call Data | 62 | PW,S,T,W | DXN, M, P/H, PCB, SVOA, W | 1607 | 0 | Hardcopy | 3.2.27 | 1989 | 1989 |
| State of Michigan Fish Consumption Advisory Data | 434 | T | B, DXN, M, P/H, PCB-A, W | 6979 | 1 | Database | 3.2.28 | 1983 | 1999 |
| 1999 FRG Demonstration Project Data - Deposit N & SMU 56/57 | 2408 | A,O,S,W | PCB-A, PCB-C, M, W, V, SVOA, P/H, DXN | 46389 | 28 | Database/ Spreadsheet | 3.2.29 | 1999 | 1999 |
| 2000 - 2001 FRG/CH2M Hill Sediment/Wood Chip Data | 428 ^a | S,WC | PCB-A, GRO, DRO, M, V, SVOA, CN | 6428 | 1 | Database | 3.2.30 | 2000 | 2001 |
| 2000 FRG/BBL Supplemental Monitoring Program Data: Surface Water ^b | 205 | W, XAD | PCB-A, PCB-C, W | | | | 3.2.31 | 2000 | 2000 |
| 2000 - 2001 FRG/BBL Supplemental Monitoring Program Data: Sediment ^b | 158 | S | PCB-A, PCB-C, W | | | | 3.2.32 | 2000 | 2001 |
| 2001 FRG/BBL Green Bay Sediment Sampling Data ^b | 30 | S | PCB-A, W | | | | 3.2.33 | 2001 | 2001 |
| 2001 FRG/BBL Water Column - High Flow Data ^b | 615 | W, XAD | PCB-A, PCB-C, W | | | | 3.2.34 | 2001 | 2001 |
| Minergy EPA SITE Data | 90 | A,O,S,W | PCB-C, M, W, V, SVOA, DXN | 8053 | 5 | Spreadsheet | na | 2001 | 2001 |
| Total: 41 Data Sets | 22377 | | | 535704 | 472 | | | | |

¹Matrices

S = Sediment
T = Tissue
W = Water
PW = Sediment Pore Water
A = Ambient Air
WC = Wood Chip
XAD = filters

²Analyses

PCB-A = PCB Aroclor
PCB_C = PCB Congener
PCB = Total PCB only
M = Metals
W = Wet Chemistry (including all Physical and Conventional data)
GRO = gas range organics
DRO = diesel range organics

V = Volatiles
SVOA = Semi-volatiles
P/H = Pesticides/Herbicides
DXN = Dioxins
B = Biological
CN = Cyanide

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1989 - 1990 GREEN BAY MASS BALANCE STUDY | 1995 - 1996 WDNR FISH TISSUE | 1996 USFWS/ HAGLER BAILLY DATA | 1995 WDNR BELOW DEPERE | | |
|--|---|--|---|--|--|---------------------------------------|----------------------------------|
| Types | Parameters: | PCBs | PCB | PCB | PCBs | TOC | Metals |
| | Requirements | Sediment | Fish Tissue | Fish Tissue | Sediment | Sediment | Sediment |
| SDG#s | | University of Minnesota - Data groups; IN0042, IN0047, IN0052, IN0057, IN0061, IN0070, IN0076, IN0078, IN0037, and IN0041 | SLOH Fish SDG-1 | Battelle Laboratory Multiple SDGs | Hazleton SDG #'s TBD2,10, 1 and 20 | Hazleton SDG #'s TBD2,10, 1 and 20 | Hazleton SDG #'s TBD2, and 20 |
| Data Review | 1) Third Party Validation Performed | Verification Only Deborah Swackhamer, Ph. D. | MA Kuehl Co | EcoChem | Y/MAKuehl | Y/MAKuehl | Y/MAKuehl |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | Yes |
| | 2) Hard copy | Some - Not sure if this is a complete set | Yes | Yes | Some | Some | Some |
| Data Review Details | 1) Package Completeness | Not determined | Yes | Yes | Yes | Yes | Yes |
| | 2) Chain of Custody Procedures | Not determined | Not determined | Yes/Minor issues | Not determined | Not determined | Not determined |
| | 3) Holding Times | Not summarized on the QA/QC Summary report Sheet | Yes | Yes | Yes | Yes | Yes |
| | 4) Initial Calibration | Not summarized on the QA/QC Summary report Sheet | Y (25%) | Y (35%) | 25% | Yes | Yes |
| | Curve - # of standards | Not summarized on the QA/QC Summary report Sheet | 5pt | 5pt | 5pt | Daily One Pt | 1point/6 point for Hg |
| | 5) Calibration Verification | Not summarized on the QA/QC Summary report Sheet | 15% D | Varies between GC/ECD and GC/MS. <25% for 75% analytes | 15% | 20% | 10% for metals & 20% for Hg |
| | Secondary Column | Not summarized on the QA/QC Summary report Sheet | 25% D | Y, data not used | 25% D for CC on 2nd column | NA | NA |
| | 6) Laboratory Blanks | Not clear. | Yes | Yes | Yes | Yes | Yes |
| | 7) Surrogate Recoveries, # required | Y - 50-120% | Y - 70-120% | Y - 50-125% | 60-150% | NA | NA |
| | 8) Matrix Spike, # required | Y - 50-120% | Y - 65-125% | Y- 50-125% tri and deca 30- 125% for mono and dichloro | 65-125% | 75-125% | 75-125% |
| | 9) Lab Duplicate | Yes/Not clear what limits are. | Y/26% Limit | Y/50% | 26% | 20% | 20% |
| | Lab Control Sample (SRM results?) | None/QAPP says that a series of blindly coded QA samples will be analyzed. | N | SRM NRC %D Carp-1 <35% | NA | NA | Y/EPA |
| | 10) Gel Permeation/Forisil Cleanup | Not provided | Y | Not mentioned | Y | NA | NA |
| | 11) Detection Limit | Not provided | 50 ug/kg | Results reported to zero | 50 ppb | NA | CRDL |
| 12) Calc and transposition verification. Qualitative verification? | Not able to determine if this was done. | Y/Recalc | Y/Recalc and Verification | Yes/Recalc performed > 10% frequency | NA | 10% | |
| 13) Field QC Results | Not apparent | NA | None | None | None | None | |
| 14) Usability Usable/Supporting | Yes | Usable | Usable | Usable | Usable | Usable | |
| Qualifiers | Qualifiers mentioned but not defined. | Y/Minor J Quals due to detections below PQL. | Yes - Qualifiers due to CCV %D outliers, BS results, surrogate outliers, lab dups, SRM results and interferences | Yes - Minor J Flags due to low surrogate recovery or below PQL and above MDL. | Yes - Minor J Flags due to poor lab RPD | None | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1989 - 1990 GREEN BAY MASS BALANCE STUDY | 1995 - 1996 WDNR FISH TISSUE | 1996 USFWS/ HAGLER BAILLY DATA | 1995 WDNR BELOW DEPERE | | |
|---------|--------------|---|---------------------------------|-----------------------------------|------------------------|----------|----------|
| Types | Parameters: | PCBs | PCB | PCB | PCBs | TOC | Metals |
| | Requirements | Sediment | Fish Tissue | Fish Tissue | Sediment | Sediment | Sediment |
| SAP | | N/Study Plan | | N | Y | | |
| QAPP | | Y | | Y/Tech Memo | Y | | |
| Lab QAM | | Answer Pending/U of M SOPs? | Y | Y/Tech Memo | Y - Hazleton SOPs | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | LOWER LAKE MICHIGAN MASS BALANCE | 1998 FOX RIVER NRDA | | | | |
|--|---|--|--|---|---|---|---|
| Types | Parameters: | Asst. Conventionals, Pest/PCB, Hg, Atrazine,DEA, DIA Water (Open Lake,Tributary), Air, Sediment, Phytoplankton | PCB | PCB Congener | PCB Congener | Pesticide | Mercury |
| | Requirements | | Fish Tissue | Fish Tissue | Fish Tissue | Fish Tissue | Fish Tissue |
| SDG#s | | BALN, GPLN, GRAN, GRLN, IUAA, IUAP, LHTL, LHTM, LHTN, LHTP, MDLH, MIAH, MNPH, RUAP, RULA, RUTA, SSSP, USTN, WSAA, WWTH, WWTN | Enchem Multiple SDGs | Michigan State University | Quanterra | Enchem Multiple SDGs | Enchem Multiple SDGs |
| Data Review | 1) Third Party Validation Performed | No- data reviewed by QC Coordinators | Exponent | Exponent | Exponent | Exponent | Exponent |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | Yes |
| | 2) Hard copy | Unknown | Yes | Yes | Yes | Yes | Yes |
| Data Review Details | 1) Package Completeness | Not addressed | Y | Y | Y | Y | Y |
| | 2) Chain of Custody Procedures | Not addressed | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable |
| | 3) Holding Times | NO DV reports provided | Y | Some exceedences Samples J/UJ | Y | Some exceedences Samples J/UJ | Y |
| | 4) Initial Calibration | NO DV reports provided | Y | Y | Y | Y | Y |
| | | Curve - # of standards | NO DV reports provided | Y | Y | Y | Y |
| | 5) Calibration Verification | NO DV reports provided | 20% | 20% | 20% | 20% | 10% |
| | | Secondary Column | NO DV reports provided | Y | Y | Y | Y |
| | 6) Laboratory Blanks | NO DV reports provided | Y | Y- U based on blank contamination | Y | Y | Y |
| | 7) Surrogate Recoveries, # required | NO DV reports provided | Y | Y | Y | Y | Y |
| | 8) Matrix Spike, # required | NO DV reports provided | Y - no quals for %R outliers | Y - no quals for %R outliers | Y - no quals for %R outliers | Y | Y |
| | 9) Lab Duplicate | NO DV reports provided | Y - MS/MSD | Y - MS/MSD | Y - MS/MSD | Y - MS/MSD | Y |
| | | Lab Control Sample (SRM results?) | NO DV reports provided | Y | Y | Y | Y |
| | 10) Gel Permeation/Forisil Cleanup | NO DV reports provided | Not Mentioned | Not mentioned | Not mentioned | Not mentioned | NA |
| | 11) Detection Limit | NO DV reports provided | NA | NA | NA | NA | NA |
| 12) Calc and transposition verification. Qualitative verification? | No recalculations were provided unable to determine if transcription checks were done | No recalculations were provided unable to determine if transcription checks were done | No recalculations were provided unable to determine if transcription checks were done | No recalculations were provided unable to determine if transcription checks were done | No recalculations were provided unable to determine if transcription checks were done | No recalculations were provided unable to determine if transcription checks were done | No recalculations were provided unable to determine if transcription checks were done |
| 13) Field QC Results | Not addressed | None identified | None identified | None identified | None identified | None identified | |
| Usability Usable/Supporting | Supporting | Usable | Usable - Some results rejected for low surrogate %R | Usable | Usable | Usable | |
| 14) Qualifiers | Y - Specific LLMB 3 character Qual codes | Y/ holdtimes, surrogate %R, LCS %R | Y/ surr %R, blank contamination -U, coplanars- J/UJ diff between GC and HRGCMS, interference, coelutions | Y/ Coelutions, greater than calibration range | Y/ Holdtimes, MS/MSD %R, Surr %R, PCB interference -all + J | Y/ Duplicate RPD | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | LOWER LAKE MICHIGAN MASS BALANCE | 1998 FOX RIVER NRDA | | | | |
|---------|--------------|---|---------------------|--------------|--------------|-------------|-------------|
| Types | Parameters: | Asst. Conventionals, Pest/PCB, Hg, Atrazine,DEA, DIA | PCB | PCB Congener | PCB Congener | Pesticide | Mercury |
| | Requirements | Water (Open Lake,Tributary), Air, Sediment, Phytoplankton | Fish Tissue | Fish Tissue | Fish Tissue | Fish Tissue | Fish Tissue |
| SAP | | | | | | | |
| QAPP | | | | | | | |
| Lab QAM | | | | | | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1994 SAIC/GAS REMEDIAL INVESTIGATION/FEASIBILITY STUDY DATA SETS | | | | | | | |
|---------------------|--|--|--|--|--|---|--|--|--|
| Types | Parameters: | PCBs | PCBs | PCBs | PCBs | PCBs | PCBs | PCBs | PCBs |
| | Requirements | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment |
| SDG#s | | ARI M172 | ARI M174 | ARI M176 | ARI M177 | ARI M178/M179/M364 | ARI M365 | ARI M367/M368 | ARI M370 |
| Data Review | 1) Third Party Validation Performed | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | 2) Hard copy | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed |
| Data Review Details | 1) Package Completeness | Y | Y | Y | Y | Y | Y | Y | Y |
| | 2) Chain of Custody Procedures | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined |
| | 3) Holding Times | Y (Frozen) | Y/Some exceed | Y | Y | Y/some exceedances. one sample qualified J for gross exceedances (M178) | Yes exceedances. several sample qualified J for gross exceedances (M365) | Yes/Minor violations | Yes/Minor violations |
| | 4) Initial Calibration | Y | Y | Y | Y | Y | Y | Y | Y |
| | Curve - # of standards | 3-5pt | 3-5pt | 5-pt | 5-pt | 5-pt | 5-pt | 5-pt | 5-pt |
| | 5) Calibration Verification | 15%D but Ave was higher. Results flagged (J/UJ). | 15%D but Ave was higher. Results flagged (J/UJ). | 15%D but Ave was higher. Results flagged (J/UJ). | 15%D but Ave was higher. Results flagged (J/UJ). | 15%D but Ave was higher. Results flagged (J/UJ). | 15%D but Ave was higher. Results flagged (J/UJ). | 15%D but Ave was higher. Results flagged (J/UJ). | 15% |
| | Secondary Column | Not indicated | Not indicated | Not indicated | Not indicated | Not indicated | Not indicated | Not indicated | Not indicated |
| | 6) Laboratory Blanks | Y | Y | Y | Y | Y | Y | Y | Y |
| | 7) Surrogate Recoveries, # required | TCMX 55-115%/DCB 70-125% | TCMX 55-115%/DCB 70-125% | TCMX 55-115%/DCB 70-125% | TCMX 55-115%/DCB 70-125% | TCMX 55-115%/DCB 70-125% | TCMX 55-115%/DCB 70-125% | TCMX 55-115%/DCB 70-125% | TCMX 55-115%/DCB 70-125% |
| | 8) Matrix Spike, # required | 35% min - 130% max | 35% min - 130% max | 35% min - 130% max | 35% min - 130% max | 35% min - 130% max | 35% min - 130% max | 35% min - 130% max | 35% min - 130% max |
| | 9) Lab Duplicate | N | Not mentioned | Not mentioned | Not mentioned | Not mentioned | Not mentioned | Not mentioned | Not mentioned |
| | Lab Control Sample (SRM results?) | Y | Y | Y | Y | Y | Y | Y | Y |
| | 10) Gel Permeation/Forisil Cleanup | Y - If necess. | Y - If necess. | Not sure | Not sure | Not sure | Not sure | Not sure | Not sure |
| | 11) Detection Limit | 50 ppb wet wt | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| | 12) Calc and transposition verification. Qualitative verification? | Y / 10%? | N - No chros | ID and Quants Could not be verified. Raw data not provided | ID and Quants Could not be verified. Raw data not provided | ID and Quants Could not be verified. Raw data not provided | Data verified | N | Not verified |
| | 13) Field QC Results | None | None | None | Not identified | Not identified | Not identified | Not identified | Not identified |
| | 14) Usability Usable/Supporting | Usable | Usable | Usable | Usable | Usable | Usable | Usable | Usable |
| | Qualifiers | Yes - Minor quals assigned due to CCV (J/UJ) | Yes - Minor quals assigned due to CCV (J/UJ) | Yes - Minor quals assigned due to CCV, surrogate recoveries J/UJ | Yes - Minor quals assigned due to CCV, surrogate recoveries J/UJ | Yes - Minor quals assigned due to CCV, surrogate recoveries J/UJ | Yes - Minor quals assigned due to CCV, surrogate recoveries J/UJ | Yes - Minor quals assigned due to CCV, surrogate recoveries J/UJ | Yes - Minor quals assigned due to CCV, surrogate recoveries J/UJ |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1994 SAIC/GAS REMEDIAL INVESTIGATION/FEASIBILITY STUDY DATA SETS | | | | | | | |
|---------|--------------|--|----------|----------|----------|----------|----------|----------|----------|
| Types | Parameters: | PCBs | PCBs | PCBs | PCBs | PCBs | PCBs | PCBs | PCBs |
| | Requirements | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment |
| SAP | | Y | | | | | | | |
| QAPP | | Y | | | | | | | |
| Lab QAM | | | | | | | | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1994 SAIC/GAS REMEDIAL INVESTIGATION/FEASIBILITY STUDY DATA SETS (cont.) | | | | | | |
|--|--|---|---|---|---|---|---|-----------------------------|
| Types | Parameters: | Dioxins | CLP Pest/PCBs | CLP SVOCs | CLP Metals | TCLP Metals | Mercury | Mercury |
| | Requirements | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment |
| SDG#s | | Triangle Lab SDG #35589 | Swanson/SDG 948521 | Swanson/SDG 948521 | Swanson/SDGs 12718, 12724, 12745, 12806, 12816, 12941 | Swanson/SDGs 12718, 12724, 12730, 12827, 12718, 12802, 12833, 12844 | Swanson WL12941 | Swanson WL12745 |
| Data Review | 1) Third Party Validation Performed | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | 2) Hard copy | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed |
| Data Review Details | 1) Package Completeness | Y | Y | N/Forms 1 not supplied by lab | Y | Y | N/Forms 1 not supplied by lab | Y |
| | 2) Chain of Custody Procedures | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined | Not determined |
| | 3) Holding Times | Yes/Minor violations | N/Samples sent to TL 10 days after collection | N/All samples exceeded HT and are qualified as estimated (J, UJ). | Y/Hg results are flagged for exceeding HT by 27 to 42 days (J/UJ) | Y | N/All samples exceeded HT and are qualified as estimated (J, UJ). | Y |
| | 4) Initial Calibration | Y | Y/Not consistent with CLP protocol | Y/Not consistent with CLP protocol | Y (Validator recalc HG results) | Y | Y/exceedance | Y/exceedance |
| | 4) Curve - # of standards | 5-pt | 5-pt | 5-pt | Lin Reg | Lin Reg | 5pt | 5pt |
| | 5) Calibration Verification | 20%RSD | N/correct concentration not used. Certain analytes outside RT window | %15D/Some exceedances qualified samples as estimated J/UJ | 10 % D | 10 % D | Y/15% | Y/15% |
| | 5) Secondary Column | NA | Not indicated | Not indicated | NA | NA | NA | NA |
| | 6) Laboratory Blanks | Y | Y | Y | Y | Y | Y | Y |
| | 7) Surrogate Recoveries, # required | TCFD 25-150%/TCDD 25-150% | TCMX 55-115%/DCB 70-125% | 8 Required/ 18% min - 137% max | NA | NA | NA | NA |
| | 8) Matrix Spike, # required | TCDD/TCDF 54-162 | 18/9 Required 29 min - 152 max | 11 Required/11% min - 142% max | 75-125% | 75-125% | 75-125% | 75-125% |
| | 9) Lab Duplicate | Not mentioned | Not mentioned | Not mentioned | Y 20%/some exceedances qualified J/UJ | Y | Y | Y |
| | 9) Lab Control Sample (SRM results?) | Y | Y | Y/acenaphthene fell outside @53% | Y | Y | Y | Y |
| | 10) Gel Permeation/Forisil Cleanup | Not sure | Not sure | Not sure | N/A | N/A | N/A | N/A |
| | 11) Detection Limit | Elevated in some samples due to blank cont. and noise | Elevated in some samples due to blank cont. and noise | N/A | N/A | N/A | N/A | N/A |
| 12) Calc and transposition verification. Qualitative verification? | Y - Sample identifications. Sample Quant not reviewed. | Not Verifiable | Y | Y. Some calc errors. | Y | N | N | |
| 13) Field QC Results | Not identified | Not identified | Not identified | None | N | Y/Field Duplicate > | N | |
| 14) Usability Usable/Supporting | Usable | Third party validation considers it unusable. | Usable | Usable - 1 data point rejected for Zn | Usable | Usable | Usable | |
| 14) Qualifiers | Yes/Due to blank cont. and elevated matrix spike recovery sample results may be biased positive (J+) | Yes/Major issues about overall quality of data. Associated with RT drift, quality of work poor. | Yes/Minor qualifications due to HT exceedances and low surr and spike recoveries (J/UJ) | Yes/Minor and Major qualifications due poor spike recoveries (J/UJ) and (R) on Zinc | No Qualifications | Yes - Minor J Flags | Yes - Minor UJ/J Flags | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1994 SAIC/GAS REMEDIAL INVESTIGATION/FEASIBILITY STUDY DATA SETS (cont.) | | | | | | |
|---------|--------------|--|---------------|-----------|------------|-------------|----------|----------|
| Types | Parameters: | Dioxins | CLP Pest/PCBs | CLP SVOCs | CLP Metals | TCLP Metals | Mercury | Mercury |
| | Requirements | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment |
| SAP | | | | | | | | |
| QAPP | | | | | | | | |
| Lab QAM | | | | | | | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1994 SAIC/GAS REMEDIAL INVESTIGATION/FEASIBILITY STUDY DATA SETS (cont.) | | | | |
|--|-------------------------------------|--|--|---|---|------------------------------|
| Types | Parameters: | Mercury | Mercury | Mercury | Mercury | Mercury |
| | Requirements | Sediment | Sediment | Sediment | Sediment | Sediment |
| SDG#s | | Swanson WL12806 | Swanson WL12812/12724/12718 | Swanson WL12816/12882/12929/12922/12853/12852/12851 | Swanson WL12688/12725/12783/12777 | Swanson WL12693 |
| Data Review | 1) Third Party Validation Performed | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC | Y/SAIC |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes |
| | 2) Hard copy | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed | Yes but not easily accessed |
| Data Review Details | 1) Package Completeness | Y | Y | Y | Y | Y |
| | 2) Chain of Custody Procedures | Not determined | Not determined | Not determined | Not determined | Not determined |
| | 3) Holding Times | Y | Y | N/Quals J/UJ | Y | Y |
| | 4) Initial Calibration | Y/exceedance | Y (Validator recalc results) | Y (Validator recalc results) | Y (Validator recalc results) | Y (Validator recalc results) |
| | | Curve - # of standards | 5pt | 5pt | 5pt | 5pt |
| | 5) Calibration Verification | Y/15% | Y/15% | Y/15% | Y/15% | Y/15% |
| | | Secondary Column | NA | NA | NA | NA |
| | 6) Laboratory Blanks | Y | Y | Y | Y | Y |
| | 7) Surrogate Recoveries, # required | NA | NA | NA | NA | NA |
| | 8) Matrix Spike, # required | 75-125% | 75-125% | 75-125% | 75-125% | 75-125% |
| | 9) Lab Duplicate | Y | Used MS/MSD | Y/Occ. Used MS/MSD SDG 12922 >35% | Y/Used MS/MSD | Y |
| | | Lab Control Sample (SRM results?) | Y | Y (not always performed) CLs were 75-125% | Used MS/MSD (75-125%) | Used MS/MSD (80-120%) |
| | 10) Gel Permeation/Forisil Cleanup | N/A | N/A | N/A | N/A | N/A |
| | 11) Detection Limit | N/A | N/A | N/A | N/A | N/A |
| 12) Calc and transposition verification. Qualitative verification? | N | Y | Y/Recalc | Y/Recalc | Y/Recalc | |
| 13) Field QC Results | N | Y/Ok on rinsate/FD (12812) failed No Action | Y/Ok on rinsate/<35% on FD | Y/Ok on rinsate/<20% on FD | Y/Ok on rinsate/OK on FD | |
| 14) Usability Usable/Supporting | Usable | Usable | Usable | Usable | Usable | |
| Qualifiers | Yes - Minor U/J/J Flags | Yes/Minor qualifications due to incorrect ICB calc. | Yes/Minor J/UJ Flags due to HT exceedances/SDG 12853 also qualified on poor FD values. | No Qualifications | Not apparent if no or some minor qualifications | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1994 SAIC/GAS REMEDIAL INVESTIGATION/FEASIBILITY STUDY DATA SETS (cont.) | | | | |
|---------|--------------|--|----------|----------|----------|----------|
| Types | Parameters: | Mercury | Mercury | Mercury | Mercury | Mercury |
| | Requirements | Sediment | Sediment | Sediment | Sediment | Sediment |
| SAP | | | | | | |
| QAPP | | | | | | |
| Lab QAM | | | | | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1998 FOX RIVER GROUP | | | | | | | | |
|--|---|--|--|--|--|--|--|--|--|---|
| Types | Parameters: | PCB | Conventionals | PCB | PCB Congeners | Pesticides | SVOC | Metals | TOC/Ammonia | PCB |
| | Requirements | Surface Water | Surface Water | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment | Fish Tissue |
| SDG#s | | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs | Quanterra Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem Multiple SDGs |
| Data Review | 1) Third Party Validation Performed | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee | Blasland Bouck & Lee |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | 2) Hard copy | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| Data Review Details | 1) Package Completeness | Y | Y | Y | Y | Y | Y | Y | Y | Y |
| | 2) Chain of Custody Procedures | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable |
| | 3) Holding Times | Y | Y/TSS samples J flagged | Y/ Dilutions done out of hold, diluted Aroclors J | Y | Y | Y/ 1 missed hold time sample J/UJ | Y | Y/ Some TOC and ammonia samples J | Y |
| | 4) Initial Calibration | Y | Y | Y | Y | Y | Y | Y | Y | Y |
| | 4) Curve - # of standards | | | | NA | NA | NA | NA | NA | NA |
| | 5) Calibration Verification | 20% | 10% | 20% | 30% Target analytes 40% Internal stds | 20% | 20% | 10% | 10% | 20% |
| | 5) Secondary Column | 20% qualitative only | NA | 20% qualitative only | NA | 20% qualitative only | NA | NA | NA | 20% qualitative only |
| | 6) Laboratory Blanks | Y | Y | Y | Y | Y | Y | Y | Y | Y |
| | 7) Surrogate Recoveries, # required | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided |
| | 8) Matrix Spike, # required | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided | Y/ Control limits not provided |
| | 9) Lab Duplicate | Y - MS/MSD/ Control limits not provided | Y / Control limits not provided | Y - MS/MSD/ Control limits not provided | Y - MS/MSD/ Control limits not provided | Y - MS/MSD/ Control limits not provided | Y - MS/MSD/ Control limits not provided | Y/ Control limits not provided | Y / Control limits not provided | Y - MS/MSD/ Control limits not provided |
| | 9) Lab Control Sample (SRM results?) | Y | Y | Y - not addressed | Y | Y | Y | Y | Y | Y |
| | 10) Gel Permeation/Forisil Cleanup | Not mentioned | NA | Not mentioned | Not mentioned | Not mentioned | Not mentioned | NA | NA | Not mentioned |
| | 11) Detection Limit | NA | NA | NA | NA | NA | NA | NA | NA | NA |
| 12) Calc and transposition verification, Qualitative verification? | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | |
| 13) Field QC Results | Field Duplicates -OK Rinsates had contamination | Field Duplicates -OK Rinsates had contamination | Field Duplicates -OK | None identified | Field Duplicates -OK | Field Duplicates -OK | Field Duplicates -OK | Field Duplicates -OK | None identified | |
| 14) Usability Usable/Supporting | Usable | Usable - except some TOC/DOC rejected | Usable | Usable | Usable | Usable | Usable - except hexachlorocyclopentadiene rejected | Usable | Usable | |
| 14) Qualifiers | Y/ Aroclor 1242 ND based on rinsate cont./ UJ extraction errors/ J/UJ low surrogate % R | Y/TOC/DOC R DOC > TOC, All parameters U rinsate, TSS J hold time | Y/ Aroclor 1242 & 1254 J spectral overlap/ J dilutions out of hold time/ minor CCAL %L | Y/1 compound J/UJ CCAL D, MS/MSD/LCS low %R, poor peak resolution | Y/ HCCP R 0% MS/MSD, minor CCAL %d, low surr %R, and missed hold time | Y/ Blank contamination, low MS %R, RPD | Y/ holdtimes | Y/ Aroclor 1242 & 1254 J spectral overlap, J /UJ due to extraction error | | |

TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA

| | | 1998 FOX RIVER GROUP | | | | | | | | |
|---------|--------------|----------------------|---------------|----------|---------------|------------|----------|----------|-------------|-------------|
| Types | Parameters: | PCB | Conventionals | PCB | PCB Congeners | Pesticides | SVOC | Metals | TOC/Ammonia | PCB |
| | Requirements | Surface Water | Surface Water | Sediment | Sediment | Sediment | Sediment | Sediment | Sediment | Fish Tissue |
| SAP | | | | | | | | | | |
| QAPP | | | | | | | | | | |
| Lab QAM | | | | | | | | | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1992/1993 DEPOSIT A SEDIMENT DATA | | | | | DEPOSIT N DEMONSTRATION PROJECT 1998 | | | | |
|--|-------------------------------------|---|---|--|---|--|--|---|---|-------------------------------------|----------------------|
| Types | Parameters: | VOA | SVOC | PCB | Pesticides | Metals/CN | PCB | PCB Congener | TOC/DOC/TSS | PCB | |
| | Requirements | Soil | Soil | Soil | Soil | Soil | Slurry, Soil, Liquid | Slurry, Soil, Liquid | Slurry, Soil, Liquid | Sludge | |
| SDG#s | | Hazleton 104116 203257 | Hazleton 104116 203242 | Hazleton SDG-1, SDG-2, SDG-3, SDG-4, SDG-5 | Hazleton 104135 203256 | Hazleton BASD34 SD01 BASD08 | Severn Trent VT. Fox9, Fox10, Fox11, Fox12, Fox13, Fox14, Fox16 | Severn Trent VT. Fox9, Fox10, Fox11, Fox12, Fox13, Fox14, Fox16 | WSLH | Severn Trent VT. Fox17 and Fox18 | |
| Data Review | 1) Third Party Validation Performed | EcoChem | EcoChem | EcoChem | EcoChem | EcoChem | MA Kuehl Co | MA Kuehl Co | MA Kuehl Co | MA Kuehl Co | |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | |
| | 2) Hard copy | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | |
| Data Review Details | 1) Package Completeness | Y | Y | Y | Y | Y | Yes | Yes | Yes | Yes | |
| | 2) Chain of Custody Procedures | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | |
| | 3) Holding Times | Y | Y | Y | Y | Y | Y - some exceedences | Y- some results J/UJ, Some results Rejected (greater than 14 days) | Y - some exceedences | Yes | |
| | 4) Initial Calibration | | Y | Y | Y | Y | Y | Y | Y | Y | Y |
| | | Curve - # of standards | Y - As required by method | Y - As required by method | Y - As required by method | Y - As required by method | Y - As required by method | NA | NA | NA | NA |
| | 5) Calibration Verification | | 20% | 20% | 20% | 20% | 10% | 15% | Y | Y | Y |
| | | Secondary Column | NA | NA | Yes | Yes | NA | Y - some %D exceedences | Y | NA | Y - %D outliers |
| | 6) Laboratory Blanks | Y - Tics rejected due to contamination | Y - Tics rejected due to contamination | Y | Y | Y | Y | Y - some results U based on MB cont. | Y | Y | |
| | 7) Surrogate Recoveries, # required | Y | Y | Y | Y | Y | Y | Y | Y | Y | |
| | 8) Matrix Spike, # required | Y - No MS/MSD for SDG 203257 J/UJ | Y - No MS/MSD for SDG 203242 J/UJ | Y | Y | Y | Y | Y | Y | Y | |
| | 9) Lab Duplicate | | Y - No MS/MSD for SDG 203257 J/UJ | Y - No MS/MSD for SDG 203242 J/UJ | Y | Y | Y | Y | Y | Y | Y |
| | | Lab Control Sample (SRM results?) | Y - No LCS for SDG 203257 J/UJ | Y - No LCS for SDG 203242 J/UJ | Y | Y | Y | Y - some %R outliers | Y - some %R outliers | Y | Y - some %R outliers |
| | 10) Gel Permeation/Forisil Cleanup | NA | NA | NA | NA | NA | NA | Not addressed | Not Addressed | NA | Not Addressed |
| | 11) Detection Limit | NA | NA | NA | NA | NA | NA | NA | NA | NA | NA |
| 12) Calc and transposition verification. Qualitative verification? | Yes | Yes | Yes | Yes | Yes | Yes | Y | Y | Yes | Yes | |
| 13) Field QC Results | None identified | None identified | Yes | Yes | None identified | Y | Y - some outliers, no quals assigned | Y - DOC RPD outlier | Y | | |
| 14) Usability Usable/Supporting | | Usable - Tics rejected due to contamination | Usable - Tics rejected due to contamination | Usable | Usable | Usable | Usable - some results rejected due to possible cross contamination | Usable - some results rejected due to exceeded holding times | Usable | Usable | |
| | Qualifiers | Y/ blank contamination U, lcal RSD, CCAL %D, no LCS MS/MSD TICs rejected due to blank contamination | Y/ blank contamination, CCAL %D, Internal std %R, NO LCS MS/MSD, TICs rejected due to blank contamination | Y/ surrogate %R, LCS %R, Field Dup RPD 1242 | Y/ RPD between main and confirmation columns NJ | Y/ Blank contamination, ICV %R CN, MS %R, GFAA post spike %R | Y - cooler temps, CCAL %D, holding time, LCS %R, Dual Column %D | Y - hold times, cooler temps, CCAI %D, method blank contamination, LCS %R, over cal | Y - holding times, cooler temps, Field Dup RPD, DOC>TOC | Y - Dual column %D outliers | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 1992/1993 DEPOSIT A SEDIMENT DATA | | | | | DEPOSIT N DEMONSTRATION PROJECT 1998 | | | |
|---------|--------------|-----------------------------------|------|------|------------|-----------|--------------------------------------|----------------------|----------------------|--------|
| Types | Parameters: | VOA | SVOC | PCB | Pesticides | Metals/CN | PCB | PCB Congener | TOC/DOC/TSS | PCB |
| | Requirements | Soil | Soil | Soil | Soil | Soil | Slurry, Soil, Liquid | Slurry, Soil, Liquid | Slurry, Soil, Liquid | Sludge |
| SAP | | | | | | | | | | |
| QAPP | | | | | | | | | | |
| Lab QAM | | | | | | | | | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | | |
|--|--|-------------------------------------|----|
| | | | |
| Types | Parameters: | PCB Congener | |
| | Requirements | Sludge | |
| SDG#s | | Severn Trent VT. Fox17 and Fox18 | |
| Data Review | 1) Third Party Validation Performed | MA Kuehl Co | |
| Deliverables | 1) Electronic Deliverables | Yes | |
| | 2) Hard copy | Yes | |
| Data Review Details | 1) Package Completeness | Yes | |
| | 2) Chain of Custody Procedures | Acceptable | |
| | 3) Holding Times | Yes | |
| | 4) Initial Calibration | | Y |
| | | Curve - # of standards | NA |
| | 5) Calibration Verification | | Y |
| | | Secondary Column | Y |
| | 6) Laboratory Blanks | Y | |
| | 7) Surrogate Recoveries, # required | Y | |
| | 8) Matrix Spike, # required | Y - some %R and RPD outliers | |
| | 9) Lab Duplicate | | Y |
| | | Lab Control Sample (SRM results?) | Y |
| | 10) Gel Permeation/Forisil Cleanup | Not addressed | |
| | 11) Detection Limit | NA | |
| 12) Calc and transposition verification. Qualitative verification? | Yes | | |
| 13) Field QC Results | Y - some outliers, no quals assigned | | |
| Usability Usable/Supporting | Usable | | |
| 14) Qualifiers | Y - CCAL %D outliers, MS/MSD %R and RPD outliers, LCS %R, over cal | | |

TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA

| Types | Parameters: | PCB Congener |
|---------|--------------|--------------|
| | Requirements | Sludge |
| SAP | | |
| QAPP | | |
| Lab QAM | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | DEPOSIT N DEMONSTRATION PROJECT 1998 (cont.) | | | | 2000/2001 FOX RIVER GROUP-LITTLE LAKE BUTTE DES MORTS | | | | | |
|---------------------------------|--|--|--|--|---------------------------|---|--|--|--|--|--|
| Types | Parameters: | TOC | PCB Congener | PCB | PCB Congener | VOC | Cyanide | PCB Aroclors | Metals | Semivolatiles | |
| | Requirements | Sludge | Surface Water | Fish | Minnow | Wood Chips | Sediment | Sediment | Sediment | Sediment | |
| SDG#s | | Severn Trent VT. Fox17 and Fox18 | WSLH | Severn Trent VT. Fox7 | WSLH | Enchem 913915 | Enchem 913915 | Enchem Multiple SDGs | Enchem 913426/913915 | Enchem 913426/913904 | |
| Data Review | 1) Third Party Validation Performed | MA Keuhl Co | MA Keuhl Co | MA Keuhl Co | MA Keuhl Co | CH2M Hill | CH2M Hill | CH2M Hill | CH2M Hill | CH2M Hill | |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | |
| | 2) Hard copy | Yes | Yes | Yes | Yes | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | |
| Data Review Details | 1) Package Completeness | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | |
| | 2) Chain of Custody Procedures | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | |
| | 3) Holding Times | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | |
| | 4) Initial Calibration | Curve - # of standards | Y | Y | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | | Calibration Verification | Y | Y | Yes | Yes | unknown | Yes | Yes | Yes | Yes |
| | 5) Secondary Column | | NA | Y | Yes | Yes | NA | NA | qualitative only | NA | NA |
| | | 6) Laboratory Blanks | Y | Y - some results U because of MB cont. | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | 7) Surrogate Recoveries, # required | Y | Y | Yes | Yes | Y/ Low recoveries | NA | Yes | NA | Y/ 2 samples J/UJ for low %R. | |
| | 8) Matrix Spike, # required | Y | N- not enough sample | No | Y | No | Y/ Lab limits | Yes/MS/MSD | Yes | Yes-MS/MSD - 1 sample J for high %R | |
| | 9) Lab Duplicate | Lab Control Sample (SRM results?) | Y - some RPD outliers | Y | Yes | Yes | No | Yes-criteria met | No | Yes | No |
| | | | Y - one outlier | Y | Yes | Yes | Yes-some low recoveries | Yes-criteria met | Yes-acceptable | Yes-acceptable | Yes-acceptable |
| | 10) Gel Permeation/Forisil Cleanup | | NA | Not addressed | Not Addressed | Not Addressed | NA | NA | Not mentioned | NA | Not mentioned |
| | 11) Detection Limit | | NA | NA | NA | NA | ppb-varies by sample and compound | ppm-varies by sample | ppb-varies by sample | ppm-varies by sample and analyte | ppb-varies by sample and compound |
| | 12) Calc and transposition verification. Qualitative verification? | | Yes | Yes | Yes | Yes | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done |
| 13) Field QC Results | | Y - some RPD outliers | Y - some outliers, no quals assigned | Yes | Yes | Field Dups & Trip Blanks -OK | Field Duplicates -OK | Field Duplicates -some high RPD with no qualifiers | Field Dup for Hg only | Field Duplicates -OK | |
| 14) Usability Usable/Supporting | | Usable | Usable | Usable | Usable | Usable | Usable | Usable | Usable | Usable | |
| Qualifiers | | Y- LCS %R, Dup RPD, Field Dup RPD | Y- blank contamination, results < LOQ, | No | Y- reported results < LOQ | Yes-All results U/UJ for low surrogate %R | No | Yes/ Many Aroclor 1254 & some 1260 qualified J due to spectral overlap | No | Yes/due to surrogate and MS %R outliers | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | DEPOSIT N DEMONSTRATION PROJECT 1998 (cont.) | | | | 2000/2001 FOX RIVER GROUP-LITTLE LAKE BUTTE DES MORTS | | | | |
|---------|--------------|--|---------------|------|--------------|---|--------------|--------------|--------------|---------------|
| Types | Parameters: | TOC | PCB Congener | PCB | PCB Congener | VOC | Cyanide | PCB Aroclors | Metals | Semivolatiles |
| | Requirements | Sludge | Surface Water | Fish | Minnow | Wood Chips | Sediment | Sediment | Sediment | Sediment |
| SAP | | | | | | Not provided | Not provided | Not provided | Not provided | Not provided |
| QAPP | | | | | | Not provided | Not provided | Not provided | Not provided | Not provided |
| Lab QAM | | | | | | Not provided | Not provided | Not provided | Not provided | Not provided |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 2000 FOX RIVER GROUP-SUPPLEMENTAL MONITORING PROGRAM-SURFACE WATER | | | | 2000/2001 FOX RIVER GROUP-SUPPLEMENTAL MONITORING PROGRAM-SEDIMENTS | | |
|---------------------|--|--|--|--|--|--|--|--|
| Types | Parameters: | Fuels (GRO/DRO) | Conventionals | PCB Aroclors | PCB Congeners | Conventionals | PCB Aroclors | PCB Congeners |
| | Requirements | Sediment | Water & XAD Resins | Water & XAD Resins | Water & XAD Resins | Sediment | Sediment | Sediment |
| SDG#s | | Enchem 913426/913904 | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem & STL Multiple SDGs | Enchem & CQM Multiple SDGs | Enchem Multiple SDGs | STL GOL020161 |
| Data Review | 1) Third Party Validation Performed | CH2M Hill | BBL | BBL | BBL | BBL | BBL | BBL |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | 2) Hard copy | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem |
| Data Review Details | 1) Package Completeness | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | 2) Chain of Custody Procedures | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable |
| | 3) Holding Times | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | 4) Initial Calibration | Yes | Yes | Yes | Yes | Yes | Yes | Yes |
| | Curve - # of standards | Lin Reg | Per method | Lin Reg | 5 pt | Per method | Lin Reg | 5 pt |
| | 5) Calibration Verification | Yes | Yes | Yes | Yes-all samples in 3 SDG qualified 1+ congeners J/UJ | Per method | Yes | Yes |
| | Secondary Column | NA | NA | qualitative only | NA | NA | qualitative only | NA |
| | 6) Laboratory Blanks | Yes | Yes | Yes | Yes-several congeners in several samples qualified U | Yes-TOC only | Yes | Yes |
| | 7) Surrogate Recoveries, # required | Yes | NA | Yes | Yes | NA | Yes | Yes |
| | 8) Matrix Spike, # required | No | Yes- TOC only | Yes-MS/MSD | No | Yes-TOC only; 20 samples J for high %R | Yes-MS/MSD | No |
| | 9) Lab Duplicate | No | Yes-criteria met | No | No | No duplicates for grain size & %moisture | No | No |
| | Lab Control Sample (SRM results?) | Yes-acceptable | Yes-criteria met | Yes-acceptable | Yes-acceptable | Yes-TOC only | Yes-acceptable | No |
| | 10) Gel Permeation/Forisil Cleanup | Not mentioned | NA | Not mentioned | NA | NA | Not mentioned | NA |
| | 11) Detection Limit | ppm-varies by sample | ppm-varies by sample | ppb-varies by sample | ppb-varies by sample & congener | TOC-ppm-varies by sample | ppb-varies by sample | ppt-varies by sample & congener |
| | 12) Calc and transposition verification. Qualitative verification? | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done |
| | 13) Field QC Results | Field Duplicates -all DRO results J due to high RPD | Field Duplicates -OK | Field Duplicates -some high RPD with no qualifiers | Field Dup for Hg only | Field Duplicates TOC only | Field Duplicates -acceptable | No |
| | Usability Usable/Supporting | Usable | Usable | Usable | Usable | Usable | Usable | Usable |
| | 14) Qualifiers | Yes/all DRO results J due to high RPD | No | No | Yes-due to blank cont., ccal, IS %R, & linear range exceed. | Yes-TOC 20 samples J for high % R | No | No |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 2000 FOX RIVER GROUP-SUPPLEMENTAL MONITORING PROGRAM-SURFACE WATER | | | | 2000/2001 FOX RIVER GROUP-SUPPLEMENTAL MONITORING PROGRAM-SEDIMENTS | | |
|---------|--------------|--|--------------------|--------------------|--------------------|---|--------------|---------------|
| Types | Parameters: | Fuels (GRO/DRO) | Conventionals | PCB Aroclors | PCB Congeners | Conventionals | PCB Aroclors | PCB Congeners |
| | Requirements | Sediment | Water & XAD Resins | Water & XAD Resins | Water & XAD Resins | Sediment | Sediment | Sediment |
| SAP | | Not provided | Not provided | Not provided | Not provided | Not provided | Not provided | Not provided |
| QAPP | | Not provided | Not provided | Not provided | Not provided | Not provided | Not provided | Not provided |
| Lab QAM | | Not provided | Not provided | Not provided | Not provided | Not provided | Not provided | Not provided |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 2001 FOX RIVER GROUP-GREEN BAY SEDIMENT SAMPLING | | 2001 FOX RIVER GROUP-WATER COLUMN - HIGH FLOW STUDY | | | |
|-----------------------------|--|--|--|--|---|--|--|
| Types | Parameters: | Conventionals | PCB Aroclors | Conventionals | PCB Aroclors | PCB Congeners | |
| | Requirements | Sediment | Sediment | Water & XAD Resins | Water & XAD Resins | Water & XAD Resins | |
| SDG#s | | Enchem & CQM 914351, 914390 | Enchem 914351, 914390 | Enchem Multiple SDGs | Enchem Multiple SDGs | Enchem & STL Multiple SDGs | |
| Data Review | 1) Third Party Validation Performed | EcoChem & BBL | EcoChem & BBL | BBL | BBL | BBL | |
| Deliverables | 1) Electronic Deliverables | Yes | Yes | Yes | Yes | Yes | |
| | 2) Hard copy | Yes | Yes | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | Yes-but only Form 1s reviewed by EcoChem | |
| Data Review Details | 1) Package Completeness | Yes | Yes | Yes | Yes | Yes | |
| | 2) Chain of Custody Procedures | Acceptable | Acceptable | Acceptable | Acceptable | Acceptable | |
| | 3) Holding Times | Yes | Yes | Yes-several TVS samples J/UJ | Yes | Yes | |
| | 4) Initial Calibration | Curve - # of standards | Per method | Lin Reg | Per method | Lin Reg | 5 pt |
| | | Calibration Verification | Per method | Yes | Per method | Yes | Yes-all samples in 1 SDG qualified 1+ congeners J/UJ |
| | 5) Secondary Column | | NA | qualitative only | NA | qualitative only | NA |
| | | 6) Laboratory Blanks | Yes-TOC only | Yes | Yes-TOC only | Yes | Yes-10 SDG had mult. congeners qualified U |
| | 7) Surrogate Recoveries, # required | NA | Yes-1 sample J due to high % R | NA | Yes-1 sample J/UJ & 1 sample J/R due to low %R | Yes-several results R due to low %R; several SDG J/UJ due to low % R | |
| | 8) Matrix Spike, # required | Yes-TOC only MS/MSD | Yes-MS/MSD | Yes-TOC only; 20 samples J for high %R | Yes-MS/MSD | No | |
| | 9) Lab Duplicate | | No duplicates for grain size & %moisture | No | No duplicates for grain size & %moisture | No | No |
| | | Lab Control Sample (SRM results?) | Yes-TOC only | Yes-acceptable | Yes-TOC only | Yes-acceptable | Yes-results in 16 samples J/UJ due to low %R |
| | 10) Gel Permeation/Forisil Cleanup | NA | Not mentioned | NA | Not mentioned | NA | |
| | 11) Detection Limit | TOC-ppm-varies by sample | ppb-varies by sample | TOC-ppm-varies by sample | ppb-varies by sample | ppt-varies by sample & congener | |
| | 12) Calc and transposition verification. Qualitative verification? | EcoChem performed recalcs and transcription checks | EcoChem performed recalcs and transcription checks | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | No recalculations were provided; unable to determine if transcription checks were done | |
| 13) Field QC Results | No | No | Field Duplicates-acceptable; Rinse blank (TOC only)-contamination | Field Duplicates -acceptable | Yes-high RPD, no action taken | | |
| Usability Usable/Supporting | Usable | Usable | Usable | Usable | Rejected (R) data not usable; all other data usable | | |
| 14) Qualifiers | Yes-TOC data estimated due to high RSD between injections | No | Yes-Several TOC samples U due to rinse blank contamination. Several TVS samples J/UJ due to HT exceedance. | Yes-1 sample J/UJ & 1 sample J/R due to low %R | Yes-several results R due to low %R. Results J/UJ due to surrogate, LCS, CCAL, co-elution & ion ratio outliers. Results U due to blank contamination. | | |

**TABLE 3-2
QC Elements for Data Sets
Supporting the Fox River RI/FS and RA**

| | | 2001 FOX RIVER GROUP-GREEN BAY SEDIMENT SAMPLING | | 2001 FOX RIVER GROUP-WATER COLUMN - HIGH FLOW STUDY | | |
|---------|--------------|--|--------------|---|--------------------|--------------------|
| Types | Parameters: | Conventionals | PCB Aroclors | Conventionals | PCB Aroclors | PCB Congeners |
| | Requirements | Sediment | Sediment | Water & XAD Resins | Water & XAD Resins | Water & XAD Resins |
| SAP | | Not provided | Not provided | Not provided | Not provided | Not provided |
| QAPP | | Not provided | Not provided | Not provided | Not provided | Not provided |
| Lab QAM | | Not provided | Not provided | Not provided | Not provided | Not provided |

APPENDIX B

TIME TRENDS ANALYSIS (MOUNTAIN-WHISPER-LIGHT, 2002)

Time Trends in PCB Concentrations in Sediment and Fish

Lower Fox River and Green Bay, Wisconsin

Prepared by:

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RETEC Project No.: WISCN-14414-215

Prepared for:

**Wisconsin Department of Natural Resources
101 S. Webster Street
Madison, Wisconsin 53707**

December 2002

Time Trends in PCB Concentrations in Sediment and Fish

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December 2002

Executive Summary

Introduction

PCBs were introduced into the Fox River, Wisconsin, from the manufacture and recycling of carbonless, multi-copy paper. PCBs were deposited in river sediments and were also passed along the food chain to fish and other wildlife. The fate of the PCBs is an important issue. This report presents rates of change of PCB concentrations in sediment and fish over time.

Methods

Sediment

Sediment samples were grouped into 23 newly designated geographic deposits that were spatially relatively compact within each river reach (see Figure 5 through Figure 8). Depth strata within each deposit were defined consistent with earlier studies: 0 to 10, 10 to 30, 30 to 50, 50 to 100, and 100+ cm. A total of 1,618 observations in 46 combinations of deposit and depth were included in the sediment time trends analysis. PCBs were analyzed as the logarithm of PCB concentration (in ppb) due to the approximately lognormal distribution of these values.

Samples were determined to be spatially correlated, and a method was used which spatially clusters observations into groups that are then approximately independent, and the statistical significance of time trends can be appropriately calculated (Lumley and Heagerty, 1999; Heagerty and Lumley, 2000).

Regression models for log PCB concentration in sediment versus time, depth, and spatial coordinates were fitted using the method of maximum likelihood, which readily incorporates the observations below detection limit. A meta-analysis was performed to yield an average time trend of PCB concentrations in surface deposits (0 to 10 cm) in each reach.

Fish

There were 19 combinations of reach, species, and sample type (whole body or fillet with skin) that had a sufficient sample size and a sufficient time spread for analysis of time trends. These 19 combinations included 867 samples. Carp and walleye provided the largest number of observations of any species.

Regression models for log PCB concentrations versus time were fitted using the logarithm of percent lipid content and time as independent variables. A linear spline function was included in some time trends analyses to accommodate a “breakpoint” and different rates of change in PCB concentrations during earlier versus later periods. The maximum likelihood method was used to accommodate observations below detection limit.

The differences in fish PCB concentrations between De Pere to Green Bay Reach and Green Bay Zone 2 were analyzed using cross-sectional data (1989–1991, five analyses) and data over time (1989–1998, four time trends analyses).

Results and Conclusions

Concentrations of PCBs in fish tissue and surface sediments have generally declined following the elimination of PCB point source discharges. However, there are statistically significant breakpoints in the decline for most of the fish species examined, suggesting that the decline has slowed down or, in some cases, that tissue concentrations of PCBs have increased. The time trends appear to be quite changeable and confidence intervals for rates are quite wide so that it is not possible to project PCB concentrations into the future for fish or sediment with much confidence.

Data on PCBs in surface sediment samples suggest that PCB concentrations have generally declined over time. Trends in concentrations of PCBs in subsurface sediments are mixed—some deposits show declining trends, while others show trends either close to zero or not significantly different from zero and yet others show increasing trends.

Specific conclusions of the time trends analysis include:

- **Fish tissue concentrations have generally declined over the period of time for which there are data in the Lower Fox River and Green Bay Zone 2.** Fish tissue PCB concentrations generally showed a slow rate of decline throughout the Lower Fox River and Green Bay Zone 2. Most time trend slopes were negative, and all statistically significant slopes for the most recent period were negative except one.
- **Significant “breakpoints” in the decline were identified for some of the fish species examined, suggesting that rates of decline in PCB tissue concentrations are changeable and may be slowing and, in some cases, tissue concentrations may be increasing.** Fish tissue concentrations have not declined at a constant rate since the 1970s. Among fish time trends analyzed, seven out of 19 combinations of reach, species, and sample type showed a statistically significant change in trend between earlier and later periods. In Little Lake Butte des Morts, De Pere Reach, and in Green Bay Zone 2, there were steep declines in fish tissue PCB concentrations from the 1970s, but with significant breakpoints in declines for some species beginning around 1980. A meta-analysis of the most recent time trends was carried out for three reaches, yielding 5 to 7 percent rates of decline per year averaged across species. Six species showed an increasing rate in their final slope, but only two of these rates were statistically significant. The existence of breakpoints and an additional analysis

showing non-constant rates suggests that rates of change are not stable and could be different in the future.

- **PCBs in surface sediment samples have generally declined over the period of time for which there are data for the Lower Fox River.** Surface sediment PCB concentrations combined within each reach by meta-analysis showed statistically significant decreasing trends in all reaches (10 to 15 percent decline per year) except Appleton to Little Rapids (1 percent increase per year). Surface sediments of individual deposits within the reaches included a mixture of positive and negative slopes. Among the 16 negative slopes, six were statistically significant; and neither of the two positive slopes was statistically significant. There were wide confidence intervals for rates of change, both for individual deposits and combined deposits, indicating that rate estimates are not precise. This imprecision and other uncertainties associated with the data prevent truly accurate future projections of sediment PCB concentrations.
- **Time trends in PCB concentrations in sediments below the surface sediment are quite varied—some indicate a decline, others indicate no change, others indicate an increase.** There is a strong trend toward fewer and weaker negative slopes at increasing depths. For Little Lake Butte des Morts, subsurface trends are mixed. The only statistically significant subsurface trend shows an increase and the other trends are a mixture of positive and negative trends. In the Appleton and De Pere reaches, there are both positive and negative trends that, taken together, are not clearly distinguishable from an overall zero trend. For Little Rapids to De Pere, there are consistently negative trends in the 10- to 30-cm strata, but in the lower strata, the data are consistent with either a zero trend (30 to 50 cm), or an increasing trend (50 to 100 cm and 100+ cm).
- **Projection of PCB concentrations into the future for fish or sediments are speculative because of imprecision and other uncertainties identified in the analysis.** The analyses carried out cannot assure a continued decline in PCB concentrations in fish and sediments over time. Even though there are a number of negative time trends that suggest PCB declines, future projection is speculative. Increases in PCB concentrations in some deeper sediments and breakpoints and other indications of changing rates in fish PCB time trends suggest that the river, its sediment, and its fish species could experience an arrest or reversal of such a decline at some time in the future.
- **PCB concentrations may increase or decrease in the future.** Some, perhaps all, of the changes in slope from before to after a breakpoint in the fish time trends may be genuine, due to

unpredictable events, such as floods accompanied by scouring and deposition. As discussed in the RI, sediment bed elevations have been altered historically and may also undergo changes in the future due to scouring and redistribution of sediments. The occurrence of these breakpoints in the past suggests that the river may change again in the future. The presence of non-constant rates of change in the post-breakpoint period also suggests unpredictability. These findings support the notion of a dynamic process, liable to change, rather than a steady state with future constant rates of change. Thus, the data do not provide assurance of a continuing future decline in PCB concentrations.

- **PCB concentrations in fish in the De Pere Reach differ from concentrations from the same species in Green Bay Zone 2.** Comparison of samples from the De Pere to Green Bay Reach (Green Bay Zone 1) and Green Bay Zone 2 showed statistically significant differences between alewife, carp, gizzard shad, and walleye in the two reaches in seven out of eight analyses. A given species and sample type differed between the reaches in one or more ways: 1) average PCB concentration differed, 2) time trend in PCB concentration differed, or 3) the relationship of PCB concentration to lipid content differed.

Discussion

Some of the considerable variability observed in the data may be accounted for by changes in river profile, burial, scour by flood or ice, and propeller wash in the lower reaches of the river during the period of data collection. Changes in sediment bed elevations have been documented and are discussed in Technical Memorandum 2g (WDNR, 1999a) and in the Remedial Investigation (ThermoRetec, 2001a). These potential mechanisms could not be introduced into the statistical analysis and could not be controlled. The time trends analysis is dependent upon the existing hydraulic conditions in the Lower Fox River. Any changes in those conditions might result in exposure of underlying PCB-laden sediments or burying of such sediments, and lead to new trends that may not be similar to the trends from the present analysis.

The conclusions of a general decrease in PCB burdens in sediments and fish of the Lower Fox River and in Zone 2 of Green Bay are consistent with findings from other research on PCBs in the Great Lakes (Offenberg and Baker, 2000; DeVault *et al.*, 1996; Lamon *et al.*, 1998; Gobas *et al.*, 1995; Smith, 2000). Some of these reports have also noted slowing of trends. Based on the present and previous studies, there may continue to be slow, gradual declines, or a steady state in PCB concentrations in fish and sediment in the future. The possibility of some increases cannot be ruled out.

Controlling for lipid content of fish samples distinctly helped in calculating more accurate time trends. The lipid content is best used as an independent variable in

regression analysis rather than as the denominator of a ratio (*PCB concentration ÷ percent lipid content*) used in traditional “lipid normalization.”

Some strengths of the study include the methods used to handle data below a detection limit, methods used to detect and handle spatial correlation of sediment samples, approaches to quantifying and testing for non-constant rates of change in fish time trends, data-driven modeling of lipid content as a factor in PCB concentrations, and meta-analysis of rates to increase precision and power. The inherent very great variability of the PCB concentrations has been thoroughly described quantitatively and graphically, and clear statements about confidence in and statistical significance of the various quantitative trends have been provided to guide the reader in the use of the trends.

Sources of Uncertainty

The data used for both sediment and fish time trends analyses are inherently quite variable. Of the 46 sediment deposit group analyses and four surface sediment meta-analyses, only 16 of the analyses can offer us a reasonably firm conclusion that PCB concentrations are changing. Two of the 16 analyses indicate increasing trends and 14 indicate decreasing trends. The remaining 34 analyses show trends with wide confidence intervals. Among the 19 analyses of individual fish species and three meta-analyses, 17 clearly demonstrate a non-zero trend. The other five analyses or meta-analyses do not support a solid “no change,” zero-slope conclusion, but yield an uncertain rate, consistent with a fairly wide range of plausible increasing or decreasing trends.

Relative depth was used rather than absolute depth. Depth of sediment is closely related to PCB concentration. We used depth defined as the distance of a sample to the sediment-water interface. Some of the time trends noted here may possibly be due to a change in the depth due to deposition or scouring over time, so that different parcels of sediment are identified with the same depth label at different times. Some changes that have occurred in sediment or fish tissue concentrations may be due to flooding, ice scouring, propeller wash, or other mechanisms that would have caused changes to the hydraulic conditions in the river or may have changed the relative depth of a deposit or a sample.

Age of fish may be related to their PCB concentrations, due to different feeding habits and locations during the lifecycle. Incorporating age proxy variables (either length or mass, unavailable in this study) might reduce unexplained variance and increase power to detect trends.

A “laboratory effect,” whereby different laboratories would produce a different mean PCB concentration on split samples, is possible. In addition, analytical techniques may have changed over the 1989-through-1998 period of sediment sample collection and both the laboratory and analytical variation may have introduced spurious positive or negative trends, or masked real trends.

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1 Introduction

PCBs are toxic chemicals that may pass through the food chain via fish, birds, and other wildlife to ultimately reach humans. PCBs were introduced into the Fox River, Wisconsin, from the manufacture and recycling of carbonless, multi-copy paper between 1954 and 1971. A number of studies have been carried out on the burden of PCBs in sediment, wildlife, water, and other media. The issue of time trends in PCB concentrations motivates our analysis. Carbonless paper manufacturing no longer introduces PCBs into the river, and other sources negligibly add to the PCB burden. Therefore, one can determine the rate at which the original store of PCBs is changing over time in fish and sediments.

In this report, therefore, we analyze the trends of PCB concentrations in sediment and fish over time. We provide quantitative estimates of rates of change of PCBs concentrations in sediments for:

- River reaches,
- Deposits within those reaches,
- Depth strata within the deposits, and
- Surface sediments combined within each reach.

We also provide quantitative estimates of time trends of PCBs in fish tissue for:

- Individual species, by reach, and
- Estimates combined across species, by reach.

In addition, we compare time trends in PCB concentrations in fish between the De Pere to Green Bay Reach and Green Bay Zone 2.

The analysis proves challenging due to the following features of the data:

- Concentrations below detection limits (both in sediment and fish),
- Spatial correlation of observations in sediment (due to the proximity of many of the samples in space),
- Potentially confounding spatial trends in sediment concentrations,
- A decline in fish PCB concentrations that, for several cases, is neither linear on the original scale of concentration per unit mass nor on a logarithmic scale,
- A limited number of sampling episodes for sediment and fish, typically leading to just a few distinct points in time for each analysis,

- Limited sample sizes for some deposits and some fish species, and
- Generally wide confidence intervals for estimates of rates of change.

Our methodology attempts to address each of the issues noted above. Despite the somewhat daunting methodology (a discussion of our methodology occupies more space in this report than our findings), the key results boil down to some fairly simple values: slope coefficients that represent the rate of change of the logarithm of PCB concentrations in sediment or fish over time. From the slope coefficients we calculated the following items of interest:

- The annual percent rate of decrease (or increase) of PCB concentrations in fish and sediments, and
- The statistical significance of the rate of change over time compared to a zero rate of change over time.

The last item refers to a “hypothesis test.” Specifically, we test the null hypothesis that a given rate of change (of sediment or fish) is zero (no change over time) versus the alternative hypothesis that the PCB concentration is either decreasing or increasing over time.

2 Methods for Sediment Analysis

2.1 Sediment Data

Sediment data were obtained from EcoChem, the contractor responsible for maintaining the Fox River database. An initial selection from the Fox River Database (FRDB) yielded 2,776 observations for the following restrictions: analyte = total PCBs; matrix = sediment; and location = Little Lake Butte des Morts, Appleton, Little Rapids, or De Pere reaches.

2.1.1 Variables of Interest

Each sediment sample was described by a number of variables, of which the following variables were used in this study:

- Sample ID (used to identify records in case of unusual values or problems),
- Location (reach designation),
- Deposit (traditional deposit designations supplied with each record within the FRDB and used in other reports on the Fox River),
- “Depth from” and “depth to” (minimum and maximum depth of a sample),
- Sample date (date sample was obtained),
- Analyte (we used only total PCB concentration),
- Qualifier (indicates whether PCBs were detected or were below detection limits, and, also, data quality),
- Northing and easting (geographic location in meters), and
- “Result,” which, in this case, gives the PCB concentration or the detection limit in $\mu\text{g}/\text{kg}$ (or parts per billion, ppb).

2.1.2 Preliminary Data Handling

We excluded the following types of data:

- Ninety-four (94) samples with northing and easting coordinates outside the river boundaries, or with no northing or easting coordinates. These were typically side samples from creeks and

tributaries, unusual samples such as bottled samples collected by divers with no exact location specified, or samples with sediment type indicated as coal composite, coarse-screened material, sand, stockpile, or non-TSCA pile;

- Thirty-four (34) samples from Appleton Deposit N, collected after January 1, 1999 (after dredging operations, which would have disturbed the natural action of the river); and
- Thirty (30) duplicated records, samples the data from which were present in more than one record in the database.

After these initial exclusions, a total of 2,618 observations were available. Any samples with a quality qualifier of R (rejected value—do not use) were ineligible for inclusion, but no samples were excluded on this basis alone.

Some data were missing the month and day, or just the day of the sample acquisition. Samples missing the day, but including month and year, were assigned to the midpoint of the month (i.e., day set to 15). Samples missing both day and month were set to the midpoint of the year (July 1). Because the time trends span data covering several years, these date imputations have a minor impact on the trend analysis.

To handle the fairly dramatic differences in concentrations and potential trends by depth, we incorporated the framework for stratifying observations by depth used in many other Fox River studies. The depth strata were right-endpoint inclusive (e.g., the interval 10 to 30 cm includes all samples with a depth greater than 10 cm and less than or equal to 30 cm): 0 to 10 cm, 10 to 30 cm, 30 to 50 cm, 50 to 100 cm, and 100+ cm). Samples were placed into a stratum based on their average depth (the mean of the minimum and maximum depth of the sample).

2.1.3 Logarithmic Transformation

We analyzed sediment and fish concentrations of PCBs after a logarithmic transformation. We implemented the \log_{10} transform for two main reasons. First, plotting the logarithm of the concentrations generated a far more normally distributed (bell-shaped) curve than plotting values on the original scale.

Second, an analysis on the log scale corresponds to modeling percent change. Expressing the rate of change as percent change per year rather than absolute change in concentration is generally more meaningful. Percentages are a common way to express rates of change (e.g., “3 percent per year”). A fixed percentage rate of change per year (analogous to compound interest) corresponds to an exponentially increasing or decreasing curve. Such a curve on the natural scale transforms to an easily modeled straight line on the logarithmic scale.

Stated another way, fixed multiplicative increments on the natural scale (as in compound interest) become fixed additive increments on the log scale.

We note, also, that the logarithmic transform is consistent with the analysis of halving and doubling times for a PCB concentration. Like the percentage rate of change, the halving (or doubling) time can readily be calculated from a model for the logarithm of PCB concentration versus time. However, throughout this report, we favor the use of the percentage rate of change over halving and doubling times. The reported percentage estimates the actual rate of change during the period when the data were collected. The halving and doubling times, however, refer to a halving or doubling of concentration that would occur only if the rate of change of log concentration remains constant over the stated halving or doubling period. For example, suppose the coefficient of time (in years) for a model of \log_{10} PCB concentration versus time is -0.01 per year during the period 1989 through 1998. The average rate of change of the PCB concentration during that period is, then, -2.3 percent per year $[=100\%(10^{-0.01} - 1)]$ and the calculated halving time is 30.1 years $[=(\log_{10} 0.5) \div (-0.01)]$. On the one hand, the -2.3 percent per year is a confident statement about a real period of time, 1989 through 1998. On the other hand, the 30.1 years for halving assumes a steady state that may not occur in a changeable river during a speculative 30.1 years. There is a one-to-one correspondence between the percentage rate of change, P , and the halving time, T ($-T$ for doubling): $P = 100(0.5^{1/T} - 1)$. Both bear the same information. We avoid, however, the connotation of possible long-term stability implied by the “doubling” and “halving” terms.

Figure 1 provides an example of a distribution of PCB concentrations plotted on the original scale (ppb, left plot), which can be compared to a plot on a logarithmic scale (\log_{10} ppb, right plot). The X-axis is an arbitrary scale for each plot, expressed as positive or negative deviations from the mean. The Y-axis shows the number of cases in each bin. A bell-shaped curve has been superimposed on each plot. The logarithmic plot shows a more symmetrical distribution and no outliers, compared to the plot on the natural scale. Generally speaking, for the hypothesis tests used in this study, such as those used to detect non-zero time trends, a more normal or “bell-shaped” distribution is less likely to lead to biased results. An exact or approximate normal distribution is desirable because the hypothesis tests used in our study assume a normal distribution. Moderate departures from this assumption are acceptable.

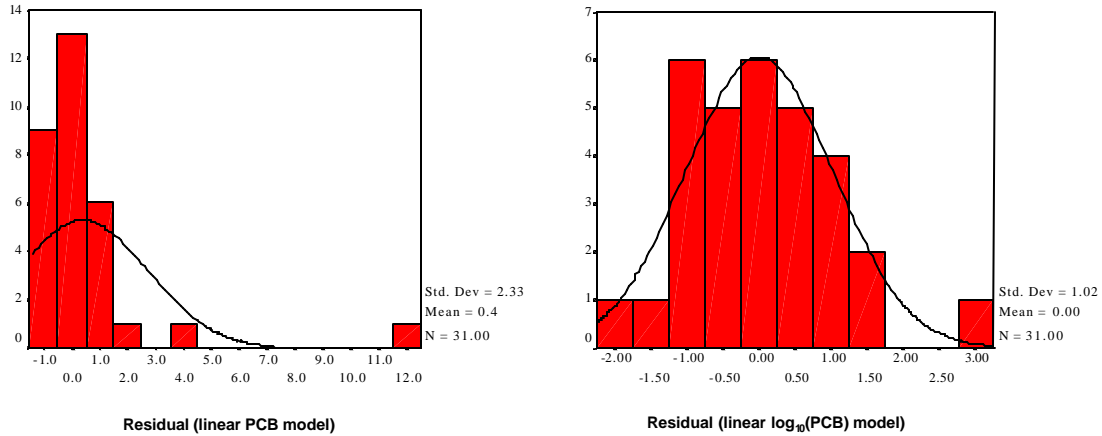


Figure 1 Example of PCB Concentration Distribution on Natural and Logarithmic Scales

Time trend estimates based on less skewed, more normal distributions are less likely to be influenced by extreme observations. A measure of skewness is the classical skewness coefficient, which is zero for symmetrical distributions and increasingly positive or negative for distributions that are increasingly stretched toward large values or small values, respectively. The normal distribution has a skewness coefficient of zero. The Appendix contains the skewness coefficients for the PCB concentrations and \log_{10} (PCB concentration). Almost all distributions of sediment PCB concentrations had smaller skewness coefficients (closer to zero) on the logarithmic scale than on the natural scale. In addition, use of the logarithmic transformation passed an important visual test for the bell-shaped normality, based on “residuals.” A residual here is defined as an observed value of \log PCB concentration minus the corresponding predicted value from the fitted regression model. If the residuals have a bell-shaped distribution, then estimates from the fitted model are more likely to be correct. To check the bell shape, we commonly use a visual display called the QQ, or “cum-cum” plot. One plots the cumulative distribution of residuals against the corresponding cumulative normal distribution. If the residuals are normally distributed, the points will all huddle along the 45 degree line. If the residuals are not normally distributed, the points will stray therefrom.

Figure 2 shows an example of a cum-cum plot. The \log PCB data (right plot) lie closer to the straight line representing the normal distribution than the PCB data on the original scale (left plot).

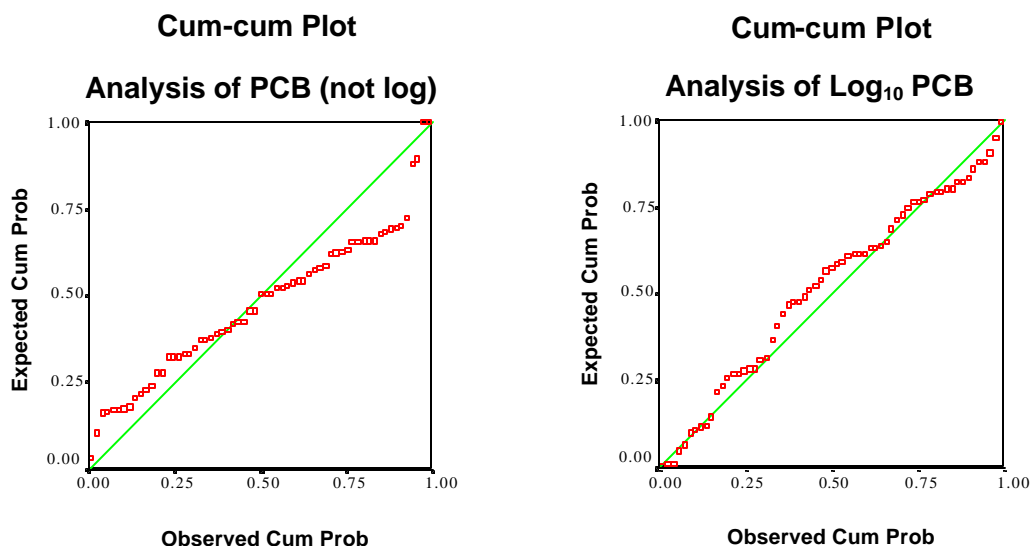


Figure 2 Comparison of Cum-cum Plots Based on Untransformed and Log-transformed Data

We have not carried out a formal hypothesis test that the distributions of \log_{10} PCB concentrations are normal. With the sample sizes used in this study (both for sediment and fish), the visual checks noted here are adequate and consistent with good statistical practice. Formal tests for normality, such as that of Kolmogorov and Smirnov, have low power for these sample sizes. In addition, regression and other procedures used in this study are robust, even if distributions are only approximately normal.

Given the good fit of the lognormal distribution to most of the observed distributions of PCB concentrations, we analyzed PCBs as the logarithm to base 10 of the PCB concentration in parts per billion. Throughout the report, our reference to PCB concentrations denotes this logarithm, unless stated otherwise. In plots and tables, the log carries the usual, easily interpreted quantification: a log value equal to 0 means an untransformed value of 1, a value of 1 represents 10, 2 represents 100, and so on.

Later, we develop models for log PCB concentration over time, i.e., “time trends” models. Given a correct model for time trend in a particular deposit, the predicted value of log PCB concentration at a specific time from the model is an unbiased estimate of the corresponding true mean log concentration at that time. The anti-logarithm of this predicted mean is an unbiased estimate of the geometric mean (GM_{est}) of PCB concentrations on the natural scale at the specified time. Because of the skewness toward large values on the natural scale, however, the geometric mean underestimates the arithmetic mean at the specified time. The arithmetic mean PCB concentration is a value of particular interest. Equation 1 provides an estimate of the arithmetic mean (AM_{est}) that can be calculated from the geometric mean.

Equation 1

$$AM_{est} = GM_{est} \exp(s^2 \cdot 5.302 / 2).$$

where GM is the geometric mean and s^2 is the estimated variance on the \log_{10} scale, calculated from a regression model. The quantity 5.302 comes from use of a \log_{10} scale rather than a \log_e scale. If a \log_e scale is used, the 5.302 can be dropped.

2.1.4 Core Averaging

We refer to the combination of some samples from the same vertical core sample as “core averaging.” As described below, proximate samples were correlated (showed similar PCB values). Thus, we replaced the log PCB concentrations of multiple samples from the same core in a given depth range with their mean (on the log scale), yielding one core-averaged sample per core per depth stratum. Twenty-five (25) percent of the sediment observations included in the analysis resulted from core averaging. A mean of 2.4 single observations contributed to a core-averaged observation. After core averaging, there were 1,980 observations.

Core averaging offers several advantages. Samples taken from exactly the same location constitute a distinct spatial sampling pattern with, possibly, different correlations than may be found among samples taken at distinct locations. Spatial correlation typically varies inversely with distance, so that samples taken close together possess stronger correlations than samples taken far apart. A distance of zero, and its infinite inverse, arising from samples taken at exactly the same location may not fit into the spatial correlation pattern present among samples collected from dispersed locations. Specifically, if $r(d)$ is the correlation between samples separated by distance d , the value of $r(0)$ may not equal the limit of $r(d)$ as d approaches zero; i.e., $r(0)$ may be an isolated discrete value. Taking the average of multiple samples from a single location will likely yield a concentration that fits better with the spatial correlation pattern from other, spatially dispersed samples. Also, multiple samples from a single location would weight that location more heavily in subsequent analyses than locations represented by a single sample. Core averaging equally weights each location.

Other than addressing an unusual correlation scenario and a statistical weighting issue, core averaging probably has little influence on the calculated time trends. A scatter plot of \log_{10} PCB concentration (Y -axis) versus time (X -axis) would spread the multiple PCB concentrations from the single location vertically around the core-averaged value at the same value for time, $X = t_0$. If the individual sample concentrations are given the same total statistical weight as the single core-averaged value, then a least-squares regression analysis of log PCB concentration versus time would yield identical slopes for either representation of the samples—core-averaged or individual. This simplified example ignores the spatial variables that we used in our regression analysis. However, the point is that core averaging is unlikely to influence the slope of a time trend.

Core averaging probably does not affect statistical significance because of two offsetting factors:

1. Heightened precision of a core-averaged log PCB concentration (compared to the less precise individual concentrations) would tend to add power to detect a non-zero slope and designate it as statistically significant.
2. Reduced sample size from core averaging would tend to subtract power to detect non-zero slopes, and would then be less likely to designate a real non-zero slope as statistically significant.

These two factors may balance out.

Core averaging imputes the mean log PCB concentration to the mean depth of the samples (all within the same stratum). Thus, core averaging reduces the information available to determine and control spatial trends. This is probably a small effect, because 75 percent of \log_{10} PCB concentrations used in the time trends analysis did not result from core averaging.

In summary, core averaging protects against a mixture of two possibly distinct spatial correlation patterns, offers equal statistical weight to each location sampled, and likely will have little influence on both estimated time trend slopes and statistical significance. It may result in slightly less precise estimates of spatial trends.

In subsequent calculations, a core-averaged value counted as one observation, on par with other single observations that had not been core averaged.

2.1.5 Observations Below Detection Limit

A number of observations dropped below detection limits. We used the maximum likelihood method (see next section) to handle these observations. In statistical parlance, observations below detection limits are designated as “censored,” which simply refers to truncated observations. Note that “censored” does not mean that observations have been excluded from the analysis. Observations both above and below detection limits contribute to the analysis. By using the maximum likelihood methodology, an observation below the detection limit brings all the information that it contains to the analysis—namely, a concentration observed as not exceeding a certain limit—and obviates the need to impute a replacement value, such as half the detection limit.

2.2 Maximum Likelihood Method

Maximum likelihood (ML) is a method very commonly used in statistics to estimate parameters such as coefficients in a regression model or in other types of models (Lawless, 1982). The precision of an estimated parameter depends on

the size of the dataset, the complexity of the model, and other factors. One expresses the precision as the standard error. In many situations, adding and subtracting twice the standard error to the estimated parameter value, as obtained from the sample, provide a 95 percent confidence interval for the true population value. That is, we are 95 percent confident that the interval includes the true population parameter. Like other estimation methods, including normal-based least-squares, ML yields: 1) an estimate of the parameter; 2) a standard error of the parameter, which indicates the precision of the estimate; and 3) a statement of statistical significance (p -value), which tells us the strength of evidence that the true parameter is not zero. One can conduct tests for statistical significance using either: 1) the parameter compared to its standard error (the ratio would be approximately normally distributed with an expected mean of zero if the true value of the parameter, such as a slope, is zero), or 2) a likelihood ratio test (LRT).

Specifying some distribution for the data is integral to the ML method. This assumption of a particular distribution is part of our model for the observed data. The models used in the current study, both for sediment and fish PCB concentrations, assume that the PCB concentration depends on some known variables. For PCB in sediments, the variables are spatial dimensions and time. For PCB concentrations in fish tissue, the variables are time, position within the annual seasonal cycle, and lipid content of the tissue. For specified values of these other variables (e.g., specified time, sediment depth, and northing and easting coordinates), the observations are assumed to occur randomly above or below an expected value. This random variation constitutes “noise.” As part of the maximum likelihood approach, one must specify the distribution of this “noise.” In our analysis, we have assumed a normal distribution for log PCB concentrations and, equivalently, a lognormal distribution for the original data. As noted earlier (Section 2.1.3), this assumption fit the distribution of log-transformed sediment concentrations exceptionally well. The normal distribution then was assumed when using the ML method with log PCB concentrations.

Data analysis customarily assumes a model, such as that noted here, for generating observations: random variation generates observations scattered around the “truth.” In this study, “the truth” of sediment time trends has been modeled as a straight line (logarithm of PCB concentration versus time) corresponding to an exponential decay of the actual PCB concentration, with appropriate adjustment for spatial coordinates. The “noise” has been modeled as the normal distribution—a bell-shaped curve.

The idea behind maximum likelihood estimation for the coefficients in a model can be illustrated by a simple example. We can visualize a scatter plot of a dependent variable (y) versus time (t) with some apparent linear trend to the scatter of points. When attempting to fit a straight line to the data, we can imagine taking the line and shifting it around the plot until we see a “best fit.” We can get residuals from this line of predicted values to the observed data

points. For a given point, the residual is the observed value minus the predicted value. Generally large residuals imply a poorer fit than generally small residuals. Given the assumption of a normal distribution of points around the line (a bell-shaped curve) at each time t and an estimate of the “width” or variance of the normal distribution around the straight line, we then can calculate the probability of getting a particular collection of residuals around the line. (The reader should note that this simplified example of PCB concentration versus time does not include spatial coordinates. The actual models developed later do include spatial coordinates.)

A straight line that does not pass through most of the data would produce a very unlikely collection of residuals. As such, the probability of such a line being a good fit would be low. Similarly, a straight line driving right through the data would produce a far more likely collection of residuals. The “best fit” line is the one with the most probable collection of residuals.

The maximum likelihood method lets us actually calculate the probability, given a particular straight line, that we would get a certain set of residuals scattered around the straight line. Each residual would contribute to that probability. For a concentration below the detection limit, we can calculate the probability, given the line, that an observation would occur at or below the specified detection limit. By multiplying together the probabilities for all residuals, we would calculate one overall probability that the given configuration of residuals would occur around this line.

We can think of the maximum likelihood method as calculating probabilities for infinitely many lines, with infinitely many values of noise around the line. The method allows one to identify the line and the value of the noise around the line with the maximum probability for the data. (The maximizing and probability concepts lead to the name “maximum likelihood.”)

One can then find the statistical significance of the slope of the line—the probability that the non-zero slope could have arisen randomly when the “truth” is a zero, or horizontal, slope. The statistical significance (or of lack thereof) of the departure of a fitted line from zero slope involves comparing a model with that slope set to zero (in this simple example a horizontal, straight line) to the model with a sloping straight line. A small change in the likelihood from the horizontal, straight line to the sloping line suggests a non-statistically significant difference, and, similarly, a non-statistically significant non-zero slope. That is, random variation could easily generate a line with this magnitude of slope.

Conversely, if we have to tilt the line quite a bit in order to get a better representation of the data, and the likelihood of that fit increases dramatically compared to the horizontal, zero-slope line, then we would probably declare the slope “statistically significant.” Such an impressively sloping line probably could not have arisen by chance if “the truth” had a zero slope. So, we would reject the hypothesis of zero slope.

The typical output from the maximum likelihood method includes:

- The estimate for each parameter,
- The standard error of the parameter estimate, and
- The statistical significance (p -value) for the null hypothesis that the true parameter is zero.

One can extend the ML method to more complex models including spatial coordinates with relative ease. Either simple or complex models will have residuals. As in the simple linear case, the more complex models also involve multiplying probabilities together and adjusting parameters in the model to get the largest overall probability of producing the observed set of residuals.

Throughout the report, significance levels of $p < 0.05$, from regression analysis or from any other analyses, have been designated as “statistically significant.” “ $p < 0.05$ ” means that there is less than 5 percent probability that an observed non-zero slope could arise randomly and differ from zero to the extent observed, if the true slope were zero.

2.3 Spatial Dependence

Analysis of sediment PCB concentrations for the Fox River data revealed a close-range spatial dependence. As will be shown later, measured total PCB concentration from samples obtained within a few centimeters or meters of one another tended to have similar values. Samples located hundreds of meters apart were more dissimilar. Thus, PCB concentrations appear to be spatially correlated.

Standard statistical methods typically assume independent observations. When data show spatial correlation, standard statistical methods may provide an unbiased parameter estimate, but they will also underestimate the standard error of the estimate, generate anticonservative p -values and confidence intervals, and overstate claims of statistical significance. This occurs because two observations that show spatial correlation do not produce as much information as two independent observations. Hence, standard statistical methods overestimate effective sample size.

Consider the following illustration of dependence, polling voters on their choice of a presidential candidate: asking five people in each of two households to choose the next president will yield 10 answers, but the true sample size will be closer to two, not 10, as people within households tend to vote more similarly than people in separate households. Asking the same question of 10 individuals from separate households in different neighborhoods across the country will yield much more information than asking five individuals within two households. As

an extreme example, we cannot obtain a precise percentage estimate of the popular vote by asking one person repeatedly 10,000 times how they expect to vote.

We investigated spatial correlation using semivariogram analysis (Cressie, 1993), a method developed in the field of mining geostatistics for assessing close-range correlation of mineral concentrations in soil samples. In our context, the semivariogram vertical axis shows the average squared difference in \log_{10} (PCB concentration) between pairs of observations, and the horizontal axis shows the distance between the observation pairs. If the observed difference in PCB concentrations is smaller for pairs close together, this curve will rise from zero up to a “sill” level, where the curve flattens out, as in Figure 3. Beyond the sill level, the approximately constant difference in concentration indicates independence between data pairs at that level of separation. The semivariogram in Figure 3 also sports a smooth curve, added to aid in assessing the sill level; these smoothed curves do not always accurately show the initial rise to the sill (as in Figure 4), due to the particular algorithm used for smoothing. The leftmost data values help to visually assess the “rise to the sill.” The leftmost point(s) are lower on the Y-axis than other points, indicating that points close together have more similar PCB concentrations than the concentrations of points farther apart. Around any trend, however, one finds considerable scatter.

The log of core-averaged concentrations was used in calculating and plotting the semivariograms. Because most observations (75 percent) did not arise from core averaging, semivariogram plots based on the original concentrations (not core averaged) would be expected to differ little from plots based on core averaging. Without core averaging, points on the plot would tend to shift upward (toward larger variances).

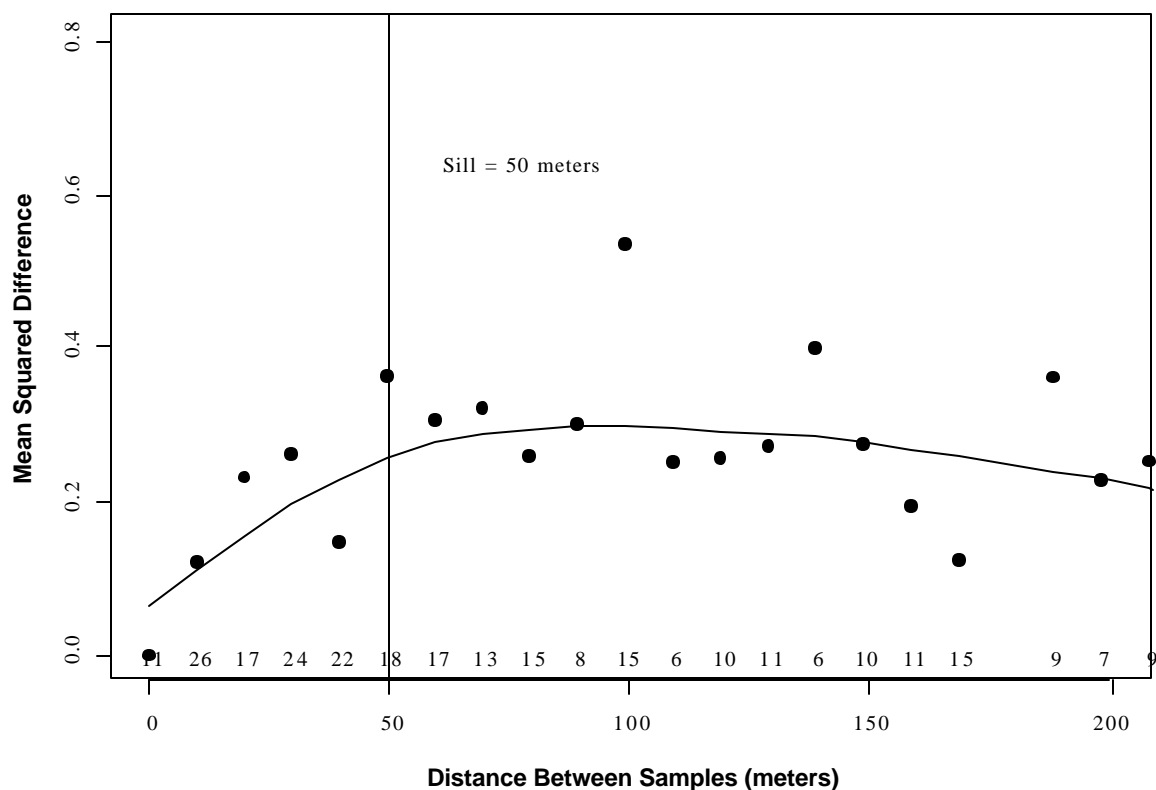


Figure 3 Semivariogram Plot of Appleton Deposit Group N Pre-dredge, 10+–30 cm Depth

The semivariogram considers all possible pairs of n samples. That is, sample #1 and sample #2 are a pair, sample #1 and sample #3 are a pair, and so on, up to the last pair, sample # n and # $(n - 1)$. There are $n(n - 1) \div 2$ total pairs. The vertical axis shows the mean squared difference in \log_{10} (PCB concentration) between a pair of samples, and the horizontal axis shows the distance between the pair. The distance between pairs of samples binned (i.e., all pairs of samples closer than about 10 meters are pooled into one bin). For each sample pair in this bin, the squared difference of their \log_{10} PCB concentration is calculated and the mean of the squared values is plotted above the bin location on the X-axis. The next bin represents pairs of samples separated by about 10 to 20 meters. Again, the mean squared difference is calculated and plotted. A similar process of calculation and plotting is carried out for all possible pairs of samples. Note that a given sample will appear in $(n - 1)$ pairs (once with each other sample). Moreover, it may occur in multiple bins as a member of some pairs that are close together and other pairs that are far apart. A smooth curve has been added to represent the trend of increasing mean squared difference with increasing distance between pairs of samples. The number of sample pairs in each bin shows just above the horizontal axis, directly beneath the estimated mean squared difference point for that bin. Samples obtained very close together show small differences, as their measured PCB values tend to be quite similar; i.e., samples obtained close together are not statistically independent. The average squared difference rises from zero as distance between points increases, up to the “sill” value (marked as 50 meters in the plot), where the average squared difference levels off and reflects the distance beyond which points are effectively independent. Semivariogram plots were used to detect spatial dependence, but no quantitative results from the semivariograms entered calculation of time trends.

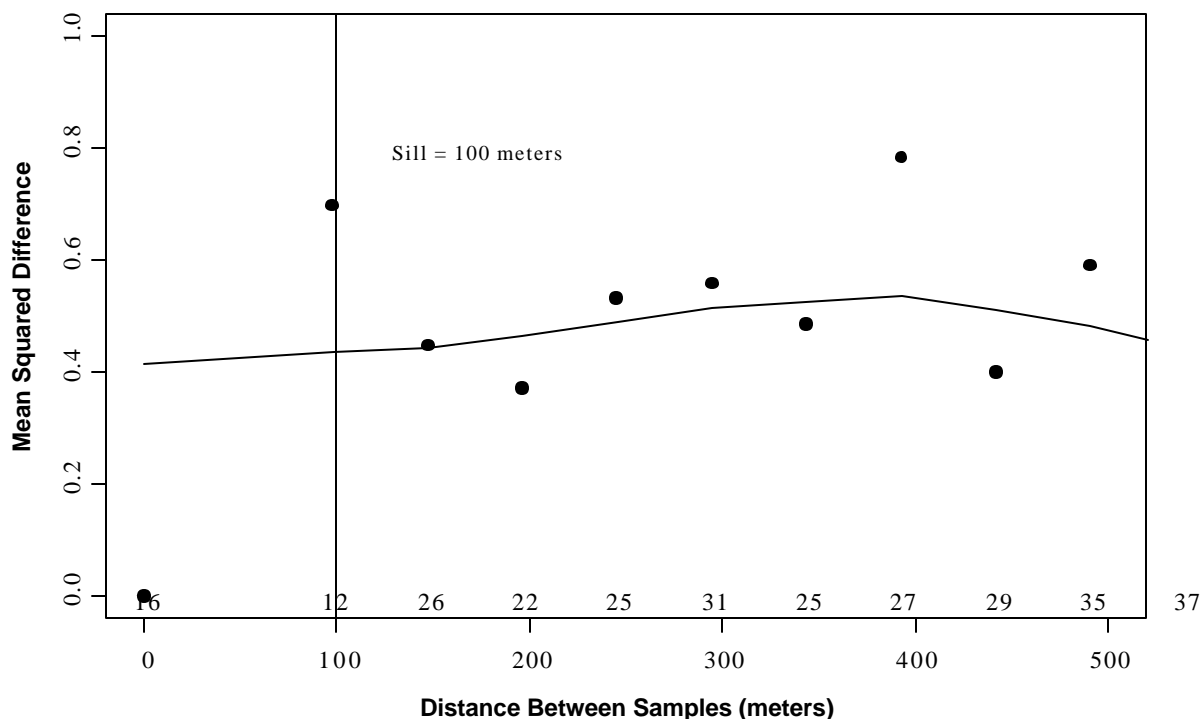


Figure 4 Semivariogram Plot of De Pere to Green Bay Deposit Group 2025, 10–30 cm Depth

(See explanation in legend of Figure 3.) Semivariogram plot portraying a case where smoother curve shows the “sill” level adequately, but does not curve up from zero due to lack of sample pairs close together. The leftmost data point indicates a very low mean squared difference for sample pairs located closer than about 25 meters. Beyond 100 meters, the average squared difference is fairly constant, indicating that samples separated by at least 100 meters are effectively independent.

Semivariograms were plotted for each of the many combinations of deposit and depth that were ultimately analyzed. The plots showed that short-range spatial dependence was pervasive in these data. Semivariogram analysis was used only to visually display spatial dependence. No quantitative results from the semivariogram analysis were used in subsequent time trends calculations. Spatial dependence was handled through the WSEV estimate, discussed below.

2.4 Addressing Spatial Dependence Using the WSEV Method

Lumley and Heagerty (1999) and Heagerty and Lumley (2000) have developed a method for more accurately assessing variability in the presence of spatial correlation using Window Subsampling Empirical Variance (WSEV) estimation. The problem being addressed is that the effective sample size is smaller than the total sample size because correlated observations do not contain as much total information as totally independent, uncorrelated observations. The WSEV method tends to lump correlated observations together into groups that are then

approximately uncorrelated. In the WSEV method, one divides up the geographic region over which the data values are obtained into a collection of windows, or subregions. We can think of the subregions being defined by a rectangular grid (with rectangular grid cells) placed over the map of sample locations. With a grid of the right spacing, the observations in different subregions of the grid will tend to be independent. The mean of the observations in a subregion can represent that subregion. The WSEV method works with means of regions, though one actually uses a more complex function than the mean. The WSEV method is analogous to using a sample size that is more closely related to the number of independent regions, rather than the number of samples available. This smaller effective sample size yields a more accurate estimate of the standard error of a parameter, more accurate confidence intervals, and a more accurate statement of statistical significance.

The ML method discussed earlier provides estimates of regression coefficients, such as a time trend slope, that do not need any adjustment. Only the standard error of these regression coefficients is adjusted by the WSEV method. In turn, the standard error is used to calculate statistical significance (a p -value).

Implementing Lumley's WSEV method involves dividing the spatial region using a coarse mesh grid, then averaging particular functions of the data within grid cells and using the averages to obtain standard error estimates for the regression model parameters. One repeats the procedure with decreasing grid mesh sizes (i.e., decreasing size of subregions), typically investigating five to ten mesh sizes. As the mesh size decreases, parameter standard errors initially increase and then decrease.

Inordinately large grid sizes result in too much averaging and subregions exhibit too little variation among themselves. As the grid size initially decreases, the estimated standard error will increase. As the grid size continues to decrease, at some point the estimated standard error will now stop increasing and begin to decrease. This occurs because neighboring cells will show too little variation due to their correlation with one another. The WSEV method uses the standard error of the regression model intercept as an aid in determining the proper grid size. We fit all of our regression models with an intercept (constant term). The WSEV standard error of the intercept will show the increasing-then-decreasing magnitude with increasing grid size as just described. In the WSEV method, the grid size that yields the largest standard error for the intercept term of the regression model is selected. From this grid size, we then calculate the WSEV standard error for the coefficient of time (the time trend slope). This standard error fully accounts for spatial dependence and is selected in an objective way.

In each analysis, we used ML estimation S-PLUS functions "SurvReg" and "CensorReg" to fit regression models and calculate the time trend slope coefficients. (The two S-PLUS functions SurvReg and CensorReg provide the same estimates of slope, but each generates different quantities used in the WSEV analyses.) Using the WSEV method, we then calculated the standard error

of the time trend slope coefficient. We wrote our own software routines (in S-PLUS) to calculate the WSEV estimates of standard error, based on output from SurvReg and CensorReg (S-PLUS 2000, Release 2, MathSoft, Seattle).

We calculated the statistical significance (p -value) of each time trend slope using the t -distribution; i.e., a “ t -test.” The t -statistic was calculated as the ratio of: 1) the time trend slope coefficient (the coefficient of time, t , in Equation 1); and 2) the WSEV standard error. The degrees of freedom for the t -statistic was the number of grid cells, at the chosen grid mesh size, which contained at least one sample. This is analogous to the number of independent groups of observations. The Appendix includes this number of non-empty grid cells.

2.5 Geographic Grouping of Data

Our need for geographically grouping samples for statistical analysis led to the creation of new “deposit groups.” The sample deposit designations in the FRDB were unsuited to defining spatially cohesive subsets, as many samples fell outside the original deposits (and had no deposit designation). Furthermore, some deposit designations spanned stretches of a river reach too long to allow adequate control of spatial variation in PCB concentration. We examined the spatial layout of all samples in each river reach. Based on this plotting and mapping exercise, we defined new “Deposit Groups,” forming data subsets with spatial variation far more amenable to statistical analysis. We named the deposit groups to reflect, to some extent, the original deposit designations already in place, with the added benefit that these groups designated non-overlapping spatial sets that included all samples. The geographic size of deposit groups is a compromise between a desire for large sample sizes in each group and a desire for tiny areas with homogeneity (i.e., relatively similar PCB concentrations within each depth stratum).

There was an isolated sample, labeled as “POG,” located by Wrightstown in the Appleton to Little Rapids Reach. The sample was located at least 2 miles from upstream samples and at least 3 miles from downstream samples. The sample was excluded.

Table 1 through Table 4 show how the original sample designations (identified in table rows) correspond to our “deposit group” designations (positioned in table columns). For example, the new “Little Lake Butte des Morts Deposit **Group E**” primarily contains samples from the original Little Lake Butte des Morts Deposit E (40 samples), but also includes four samples from the original Little Lake Butte des Morts Deposit D and nine from Deposit POG. Samples with no deposit designation in the FRDB constitute from 5 to 70 percent of samples within each of the four reaches (Table 1 through Table 4). Little Lake Butte des Morts had 5 percent of samples with no deposit designation (presumably samples located spatially outside the original deposit designations). The corresponding percentages of samples without designations in other reaches were 7 percent for Appleton Reach, 12 percent for Little Rapids Reach, and 72 percent for the De

Pere Reach. The large percentage for De Pere Reach arises because the original deposit designations were noted only for SMU Deposits 50–67. Our new “deposit group” designation includes all samples and thus increases sample sizes available for trend estimates and hypothesis tests. In any case, having an original deposit designation became irrelevant with the formation of our new deposit groups. Furthermore, the lack of an original deposit designation had no role in disqualifying a sample from inclusion in our time trends analysis. Finally, not having an original deposit designation does not suggest poor data quality.

Table 1 Little Lake Butte des Morts Deposit Groups Defined for Time Trends Analysis

| Original Deposit Designation | Time Trends Analysis: Deposit Group Designation | | | | | | | Total |
|------------------------------|---|-----------------------|-------------------------|-----------------------|-----------------------|-----------------------|------------------------|------------|
| | LLBdM Deposit Group AB | LLBdM Deposit Group C | LLBdM Deposit Group POG | LLBdM Deposit Group D | LLBdM Deposit Group E | LLBdM Deposit Group F | LLBdM Deposit Group GH | |
| Deposit A | 281 | 0 | 0 | 0 | 0 | 0 | 0 | 281 |
| Deposit B | 5 | 0 | 0 | 0 | 0 | 0 | 0 | 5 |
| Deposit C | 0 | 52 | 0 | 0 | 0 | 0 | 0 | 52 |
| Deposit D | 0 | 0 | 1 | 49 | 4 | 8 | 0 | 62 |
| Deposit E | 0 | 0 | 2 | 1 | 40 | 68 | 32 | 143 |
| Deposit F | 0 | 0 | 0 | 0 | 0 | 12 | 0 | 12 |
| Deposit G | 0 | 0 | 0 | 0 | 0 | 0 | 3 | 3 |
| Deposit H | 0 | 0 | 0 | 0 | 0 | 0 | 3 | 3 |
| Deposit POG | 0 | 0 | 27 | 0 | 9 | 0 | 0 | 36 |
| No Designation | 13 | 2 | 4 | 5 | 0 | 10 | 0 | 34 |
| Total: | 299 | 54 | 34 | 55 | 53 | 98 | 38 | 631 |

Note:

Column entries show number of samples from original deposits included in each time trends deposit group.

Table 2 Appleton Deposit Groups Defined for Time Trends Analysis

| Original Deposit Designation | Time Trends Analysis: Deposit Group Designation | | | | | Total |
|------------------------------|---|--------------------------|----------------------------|---------------------------|---------------------------|------------|
| | Appleton Deposit Group IMOR | Appleton Deposit Group N | Appleton Deposit Group VCC | Appleton Deposit Group SU | Appleton Deposit Group DD | |
| Deposit AA | 0 | 0 | 1 | 0 | 0 | 1 |
| Deposit BB | 0 | 0 | 3 | 0 | 0 | 3 |
| Deposit CC | 0 | 0 | 9 | 0 | 0 | 9 |
| Deposit DD | 0 | 0 | 0 | 0 | 20 | 20 |
| Deposit I | 4 | 0 | 0 | 0 | 0 | 4 |
| Deposit J | 2 | 0 | 0 | 0 | 0 | 2 |
| Deposit K | 3 | 0 | 0 | 0 | 0 | 3 |
| Deposit L | 3 | 0 | 0 | 0 | 0 | 3 |
| Deposit M | 2 | 0 | 0 | 0 | 0 | 2 |
| Deposit N | 0 | 136 | 0 | 0 | 0 | 136 |
| Deposit O | 7 | 0 | 0 | 0 | 0 | 7 |
| Deposit P | 12 | 0 | 0 | 0 | 0 | 12 |
| Deposit Q | 12 | 0 | 0 | 0 | 0 | 12 |
| Deposit R | 2 | 0 | 0 | 0 | 0 | 2 |
| Deposit S | 0 | 0 | 0 | 7 | 0 | 7 |
| Deposit T | 0 | 0 | 0 | 15 | 0 | 15 |
| Deposit U | 0 | 0 | 0 | 3 | 0 | 3 |
| Deposit V | 0 | 0 | 7 | 0 | 0 | 7 |
| Deposit W | 0 | 0 | 39 | 0 | 0 | 39 |
| Deposit X | 0 | 0 | 46 | 0 | 0 | 46 |
| Deposit Y | 0 | 0 | 3 | 0 | 0 | 3 |
| Deposit Z | 0 | 0 | 2 | 0 | 0 | 2 |
| No Designation | 9 | 0 | 15 | 0 | 0 | 24 |
| Total: | 56 | 136 | 125 | 25 | 20 | 362 |

Note:

Column entries show number of samples from original deposits included in each time trend deposit group.

Table 3 Little Rapids Deposit Groups Defined for Time Trends Analysis

| Original Deposit Designation | Time Trends Analysis: Deposit Group Designation | | | | Total |
|------------------------------|---|--------------------------------------|--------------------------------|----------------------------------|------------|
| | Little Rapids Deposit Group Upper EE | Little Rapids Deposit Group Lower EE | Little Rapids Deposit Group FF | Little Rapids Deposit Group GGHH | |
| Deposit EE | 100 | 96 | 94 | 145 | 435 |
| Deposit FF | 0 | 0 | 3 | 5 | 8 |
| Deposit GG | 0 | 0 | 0 | 75 | 75 |
| Deposit HH | 0 | 0 | 0 | 49 | 49 |
| No Designation | 4 | 22 | 0 | 52 | 78 |
| Total: | 104 | 118 | 97 | 326 | 645 |

Note:

Column entries show number of samples from original deposits included in each time trend deposit group.

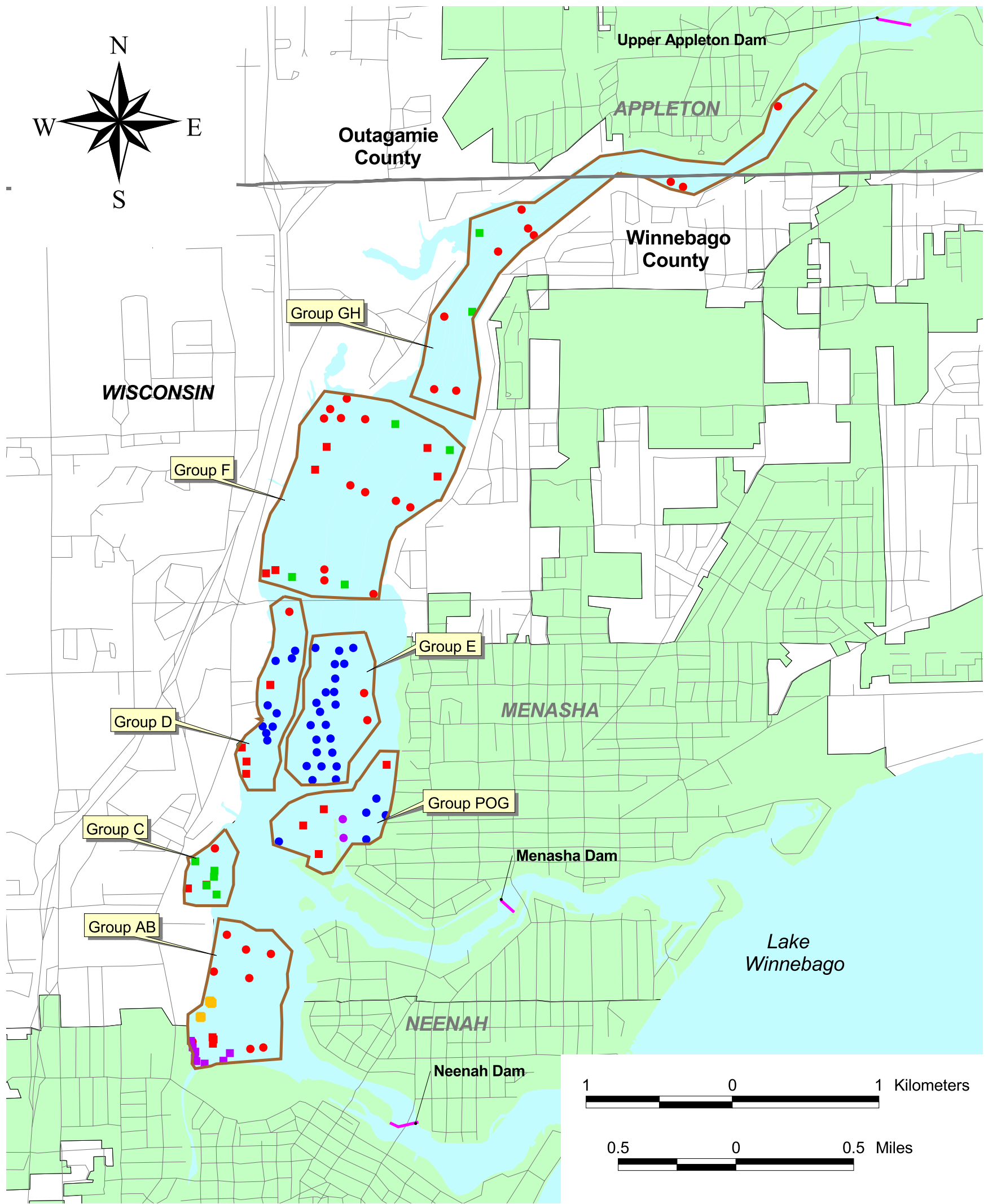
Table 4 De Pere SMU Groups Defined for Time Trends Analysis

| Original Deposit Designation | Time Trends Analysis: Deposit Group Designation | | | | | Total |
|------------------------------|---|------------------------|------------------------|------------------------|-------------------------|--------------|
| | De Pere SMU Group 2025 | De Pere SMU Group 2649 | De Pere SMU Group 5067 | De Pere SMU Group 6891 | De Pere SMU Group 92115 | |
| SMU56/57 | 0 | 0 | 282 | 0 | 0 | 282 |
| No Designation | 201 | 284 | 97 | 88 | 61 | 731 |
| Total: | 201 | 284 | 379 | 88 | 61 | 1,013 |

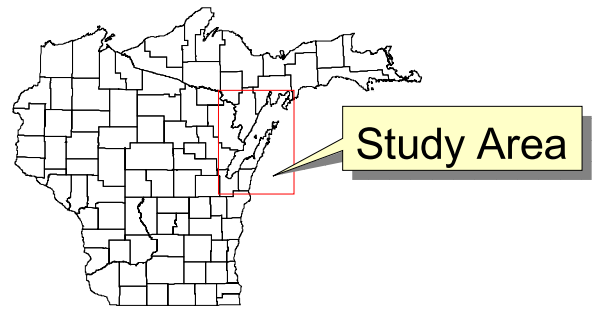
Note:

Column entries show number of samples from original deposits included in the time trends SMU group.

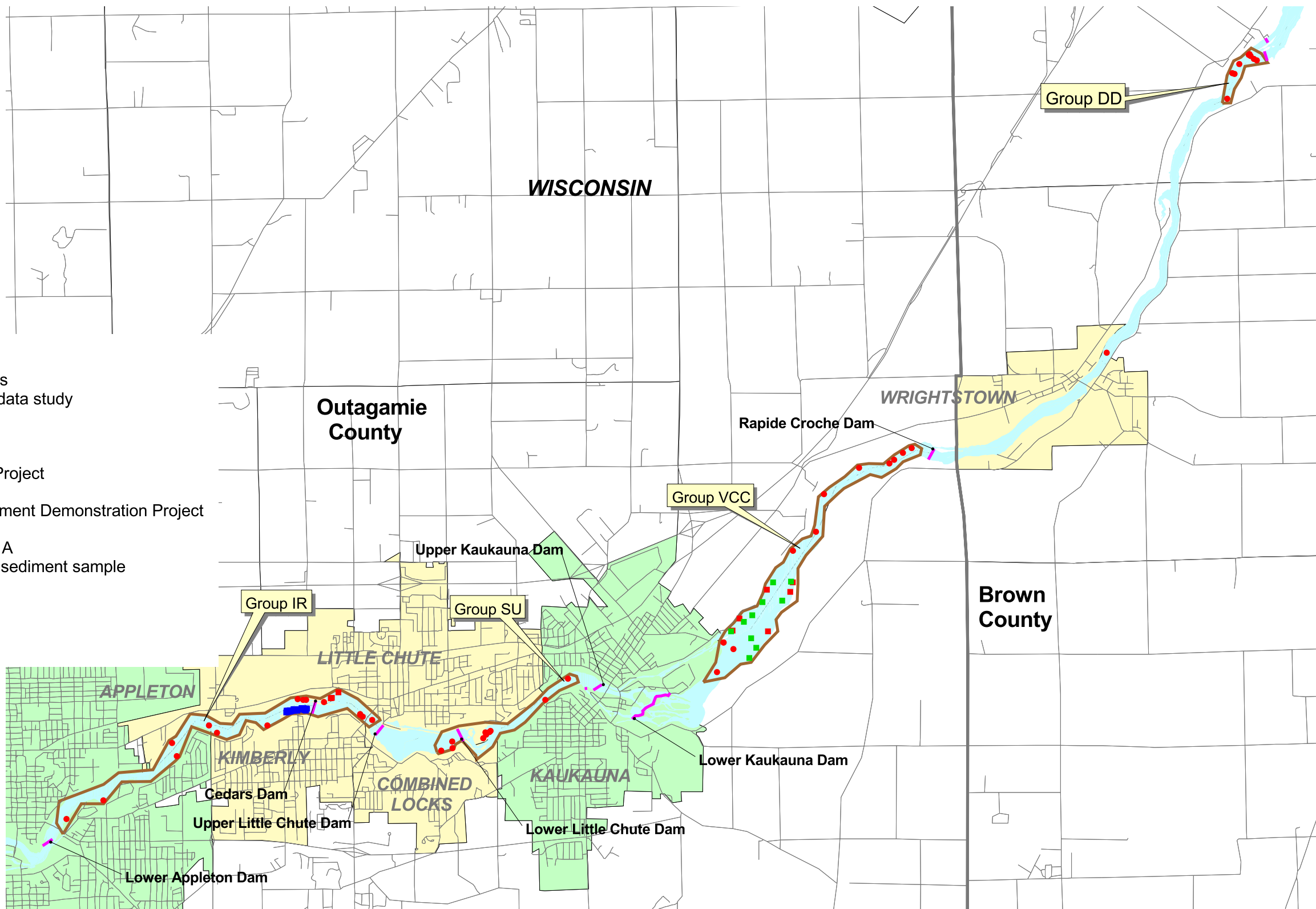
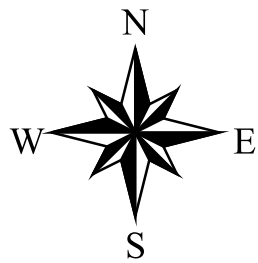
Figure 5 through Figure 8 map the location of samples and our deposit groups in the four river reaches. The boundaries separating the deposits were approximations drawn by eye, as formal definitions were unnecessary. Figure 8 breaks our SMU groups into smaller units than actually used, showing some of the original SMU designations. Our SMU Group 2025 aggregated (approximately) the original SMU designations 20–25; our SMU Group 2649 aggregated the original SMU designations 26–49; and so on for our SMU groups 5067 (aggregating 50–67), 6891 (aggregating 68–91), and 92115 (aggregating 92–115).



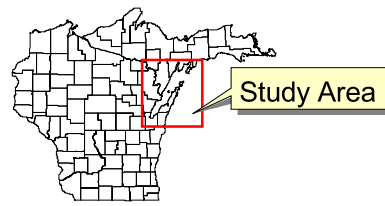
- Revised Deposit Outlines
- Sediment Sample Data Collection Points**
- 1989/90 Mass Balance sediment data study
- 1994 SAIC and GAS study
- 1995 WDNR
- 1996 BBL data study
- 1997 SMU 56/57 Demonstration Project
- 1998 BBL data study
- 1998 Deposit N Post-Dredge sediment Demonstration Project
- 1998 RI/FS Supplemental data
- 1992/1993 LLBDM RI/FS Deposit A
- 1994 Woodward Clyde Deposit A sediment sample
- ~ Dam Locations
- Roads
- County Boundary
- Water



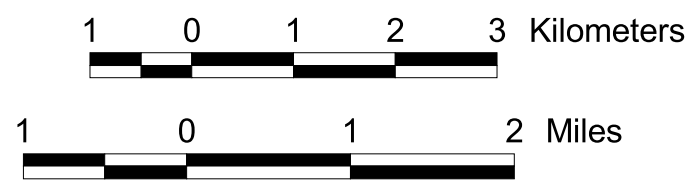
NOTES:
 1. Basemap generated in ArcView GIS, version 3.2, 1998, and TIGER census data, 1995.
 2. Sediment sample point data obtained from Wisconsin Dept. of Natural Resources, 1999, and are included in the Fox River database.
 3. Revised deposit outlines created by Nayak Polissar, Mountain-Whisper-Light Statistical Consulting, and Stephen Jesse, ThermoRetec Engineering Consultants, 2000.




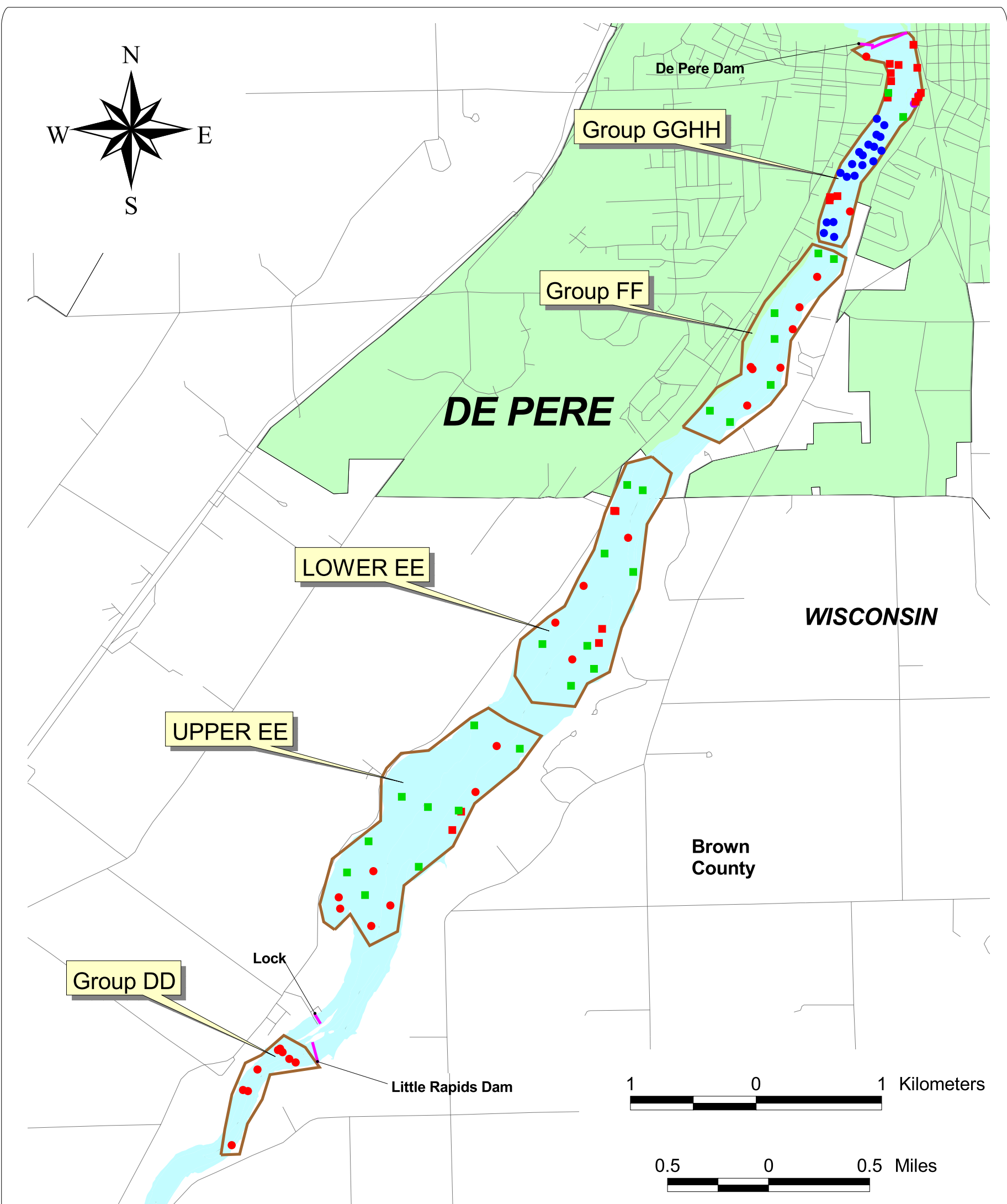
- Revised Deposit Outlines
- Sediment Sample Data Collection Points**
- 1989/90 Mass Balance sediment data study
- 1994 SAIC and GAS study
- 1995 WDNr
- 1996 BBL data study
- 1997 SMU 56/57 Demonstration Project
- 1998 BBL data study
- 1998 Deposit N Post-Dredge sediment Demonstration Project
- 1998 RI/FS Supplemental data
- 1992/1993 LLBDM RI/FS Deposit A
- 1994 Woodward Clyde Deposit A sediment sample
- Dam Locations
- Roads
- County Boundary
- Water



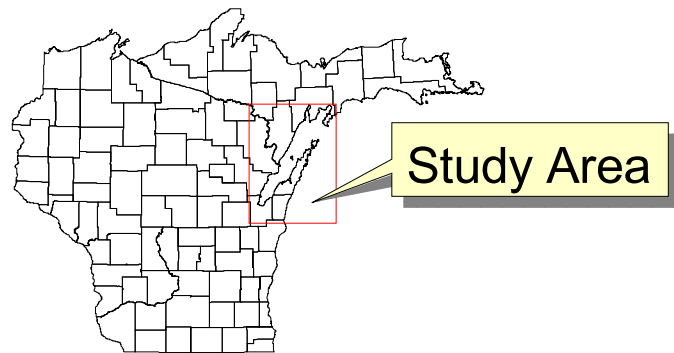
NOTES:
 1. Basemap generated in ArcView GIS, version 3.2, 1998, and TIGER census data, 1995.
 2. Sediment sample point data obtained from Wisconsin Dept. of Natural Resources, 1999, and are included in the Fox River database.
 3. Revised deposit outlines created by Nayak Polissar, Mountain-Whisper-Light Statistical Consulting, and Stephen Jesse, ThermoRetec Engineering Consultants, 2000.



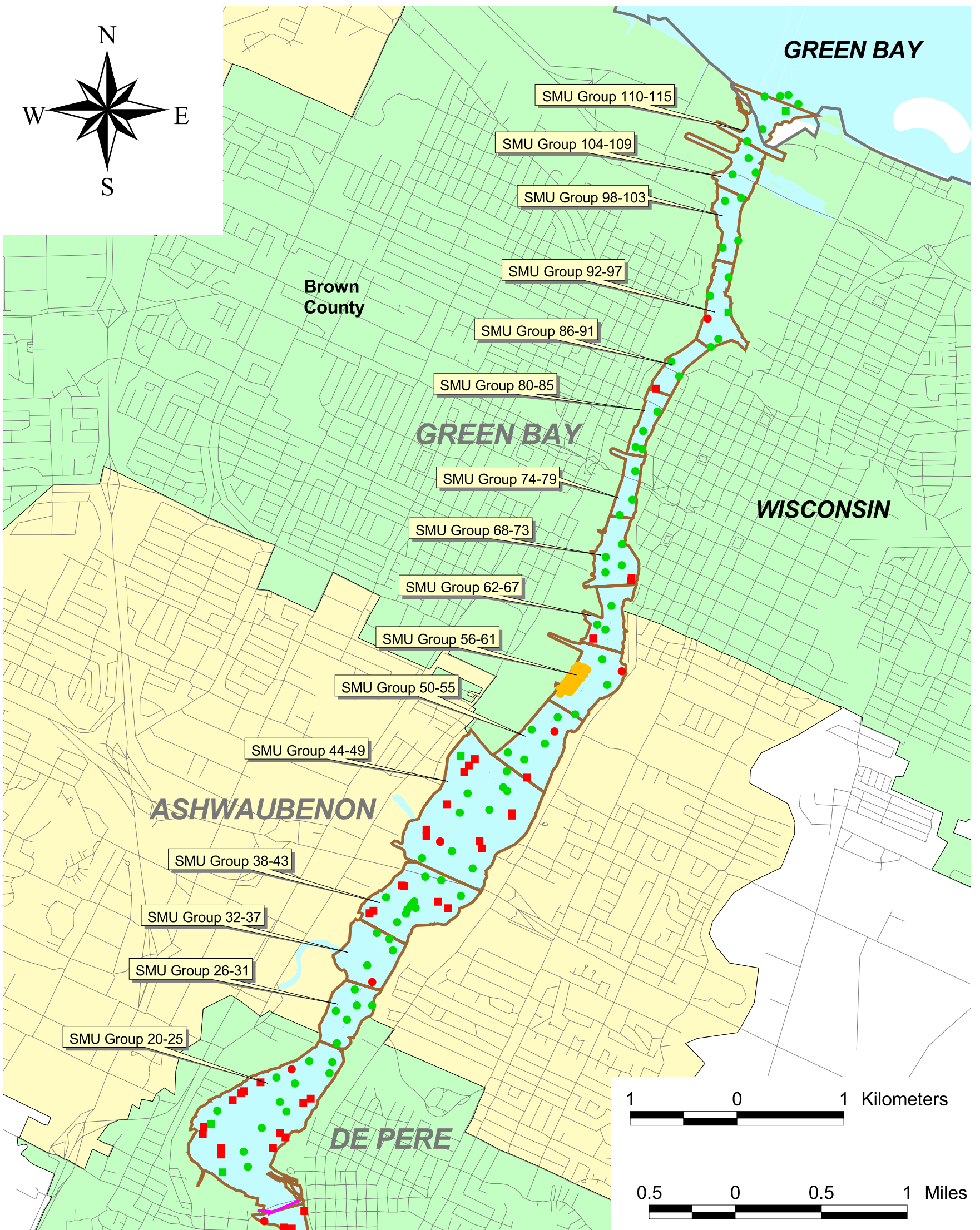
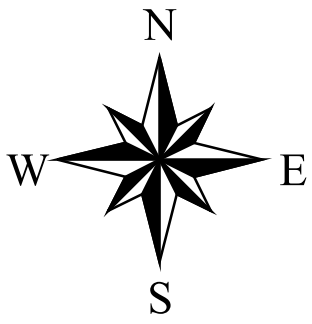
| | | | | |
|---|------------------------------------|------------------------|---|--|
|  <small>Smart Solutions. Proven Outcomes.</small> | Natural Resource Technology | Risk Assessment | Sample Point Groups for Sediment Time Trend Analysis: Appleton to Little Rapids FIGURE 6 | FIGURE NO: RA-14414-425-2 PRINT DATE: 1/23/01 CREATED BY: SCJ APPROVED: AGF |
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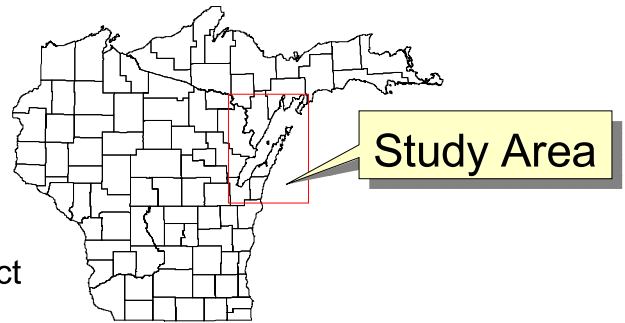
- Revised Deposit Outlines
- Sediment Sample Data Collection Points**
- 1989/90 Mass Balance sediment data study
- 1994 SAIC and GAS study
- 1995 WDNR
- 1996 BBL data study
- 1997 SMU 56/57 Demonstration Project
- 1998 BBL data study
- 1998 Deposit N Post-Dredge sediment Demonstration Project
- 1998 RI/FS Supplemental data
- 1992/1993 LLBDM RI/FS Deposit A
- 1994 Woodward Clyde Deposit A sediment sample
- Dam Locations
- Roads
- County Boundary
- Water



NOTES:
 1. Basemap generated in ArcView GIS, version 3.2, 1998, and TIGER census data, 1995.
 2. Sediment sample point data obtained from Wisconsin Dept. of Natural Resources, 1999, and are included in the Fox River database.
 3. Revised deposit outlines created by Nayak Polissar, Mountain-Whisper-Light Statistical Consulting, and Stephen Jesse, ThermoRetec Engineering Consultants, 2000.



- SMU Deposits
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Figure 9 through Figure 12, show the location of each sample in a rectangular coordinate system devoid of map features. The “northing” and “easting” rectangular coordinates locate each sample along a north-south and east-west axis, respectively, based on a standard geographic coordinate system for Wisconsin State. Northing and easting are expressed in meters relative to an origin not shown on the plot.

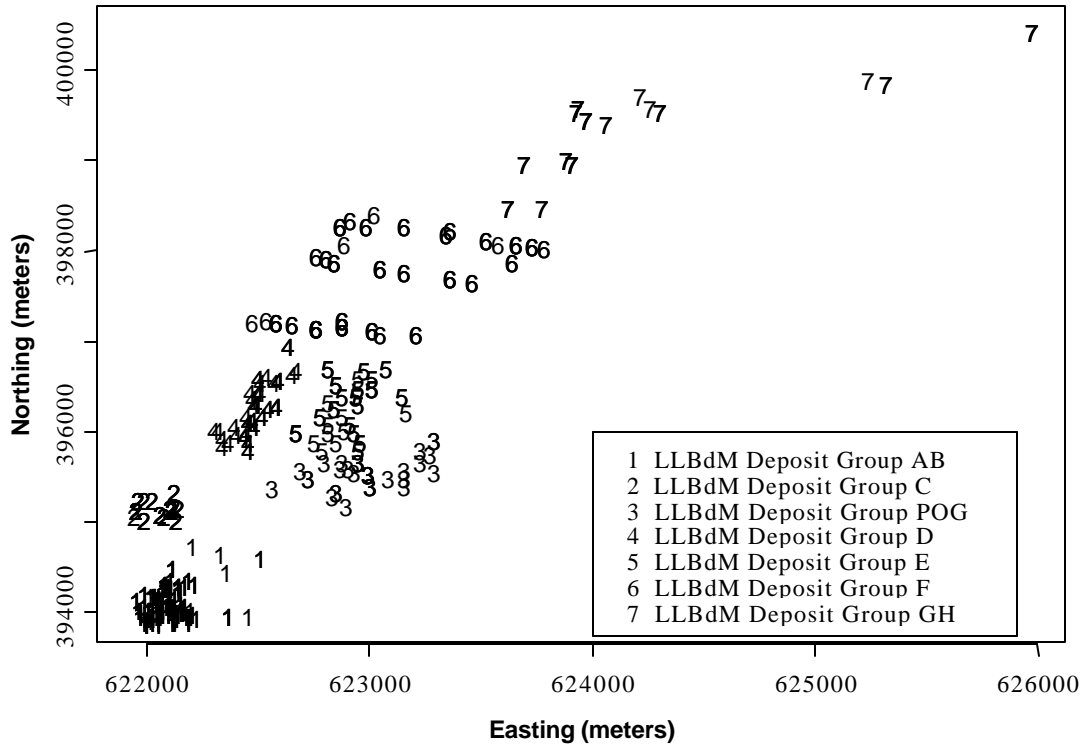


Figure 9 Locations of Deposit Groups in Little Lake Butte des Morts Reach

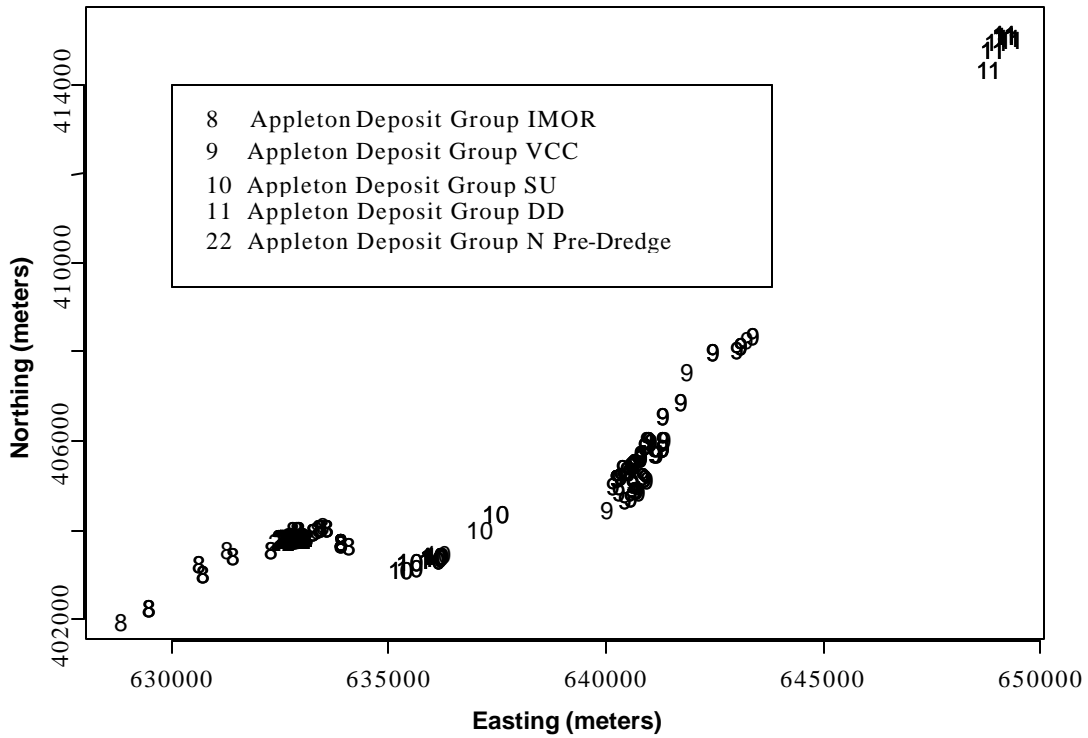


Figure 10 Locations of Deposit Groups in Appleton Reach

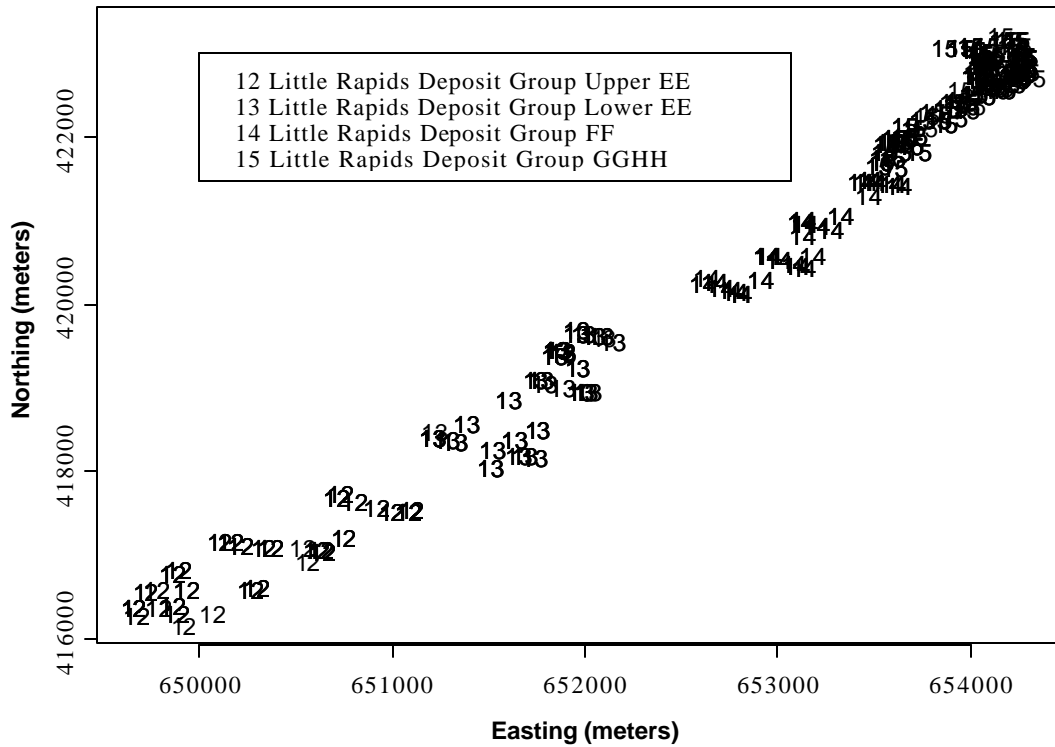


Figure 11 Locations of Deposit Groups in Little Rapids Reach

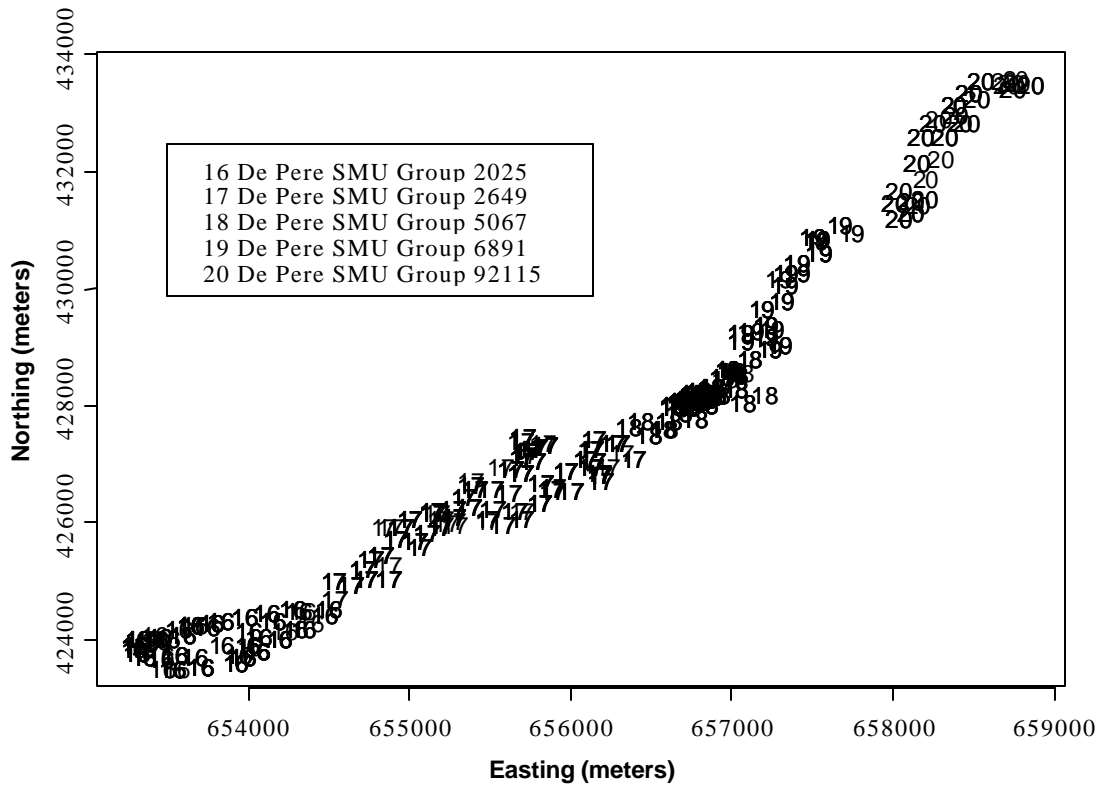


Figure 12 Locations of Deposit Groups in De Pere Reach

2.6 Models for Variation in PCB Concentration in Space and Time

Because PCB concentrations vary spatially as well as over time, we included spatial dimensions in our regression models. To characterize the spatial component in the models, we used linear and quadratic variables for “easting” and “northing” coordinates (east and north distances in meters) and a linear term for depth. For every stratum, depth was measured from a value of zero at the top of the 0- to 10-cm layer. Depth, thus, means simply distance from the surface of the river sediment at the time the sample was taken. We centered the northing and easting coordinates for each depth stratum in each deposit group. “Centering” involved finding the spatial centroid of the samples used in the regression analysis for the specific deposit group and depth stratum. Given a set of northing and easting coordinates, the sample centroid sits at the mean of the northing and easting coordinates. We produced the centered northing (*N*) and easting (*E*) coordinates by subtracting the centroid from the northing and easting coordinates of each sample.

Under this new coordinate system, the centroid of each deposit group at each depth stratum is the origin of a coordinate system with coordinates (0, 0). By centering, one avoids round-off problems when using the fitted regression models. Without centering, calculating a fitted concentration would involve subtracting a very large number from a second large number. The difference of interest (a log PCB concentration) is usually relatively close to zero. Thus, the later digits for the two large numbers must be tabulated accurately. A simple hypothetical example illustrates this point. Let us ignore time and consider only easting, where an equation

$$\log_{10} PCB = 2.24 + 0.016E_c$$

indicates that \log_{10} PCB concentration increases by 0.016 for each meter to the east of the centroid of a deposit group. At the centroid, the \log_{10} PCB concentration is 2.24 (the value of the intercept). E is the centered easting coordinate. If E^* is the original (uncentered) easting coordinate and the deposit group centroid E^* mean = 622,347 meters (a realistic value for this study), then the equation for \log_{10} PCB concentration with the original easting coordinate would be

$$\log_{10} PCB = -9955.312 + 0.016E^*.$$

If this cumbersome second equation is used with $E^* = 622,347$ (the centroid), $\log_{10} PCB = 2.24$ is calculated accurately for the centroid location. However, if -9955.312 is casually rounded to 9955, an estimate of 2.552 is obtained (instead of the correct 2.24), off by +0.312 units, which, on the natural scale (not log), corresponds to approximately a doubling of the concentration. Thus, centering helps computation and presentation. For the same reason, time was measured from January 1, 1989, taken as $time = 0$.

The specific regression model fitted to the PCB concentrations was:

Equation 2

$$\log_{10} PCB = b_0 + b_t \cdot t + b_D \cdot D + b_E \cdot E + b_N \cdot N + b_{E^2} \cdot E^2 + b_{N^2} \cdot N^2$$

where

- $\log_{10} PCB$ = the logarithm (base 10) of the PCB concentration in $\mu\text{g}/\text{kg}$ (ppb) by weight,
- t = time in years since January 1, 1989,
- D = depth in centimeters from the sediment-water interface,
- E = the centered easting coordinate for the particular deposit group and depth stratum (meters), and
- N = the centered northing coordinate (meters).

The intercept is b_0 and b_t , b_D , etc., are regression coefficients. E^2 and N^2 are the quadratic terms for centered easting and northing.

Based on scatter plots of PCB concentrations versus easting coordinate or northing coordinate, we included the quadratic terms (E^2 and N^2) for easting and northing in the regression models whenever we analyzed at least 20 samples. For sample sizes smaller than 20, we included the quadratic terms whenever we suspected a potential curvilinear trend of \log_{10} PCB concentration versus northing or easting.

We note that we included up to five variables to describe spatial variation: D , E , N , E^2 , and N^2 . These five variables are sometimes needed to describe five unique kinds of spatial variation in concentrations of PCBs: linear trends in depth, easting and northing, and curvilinear trends in easting and northing. When there is a deposit group and stratum with little variation in one of these variables (e.g., little curvilinear trend in the easting direction), then the coefficient of that variable will be zero or close to zero, and it is virtually harmless to include it in a model. Because of widely varying sample sizes, we did not wish to tailor the spatial model to each deposit group and stratum; in some cases, the small sample sizes yield insufficient power to formally accept or reject a given type of spatial variation, such as curvilinearity. Due to low power to detect the need for variables for the spatial dimensions, one errs on the side of safety by including all, rather than erroneously excluding some. With fewer than 20 observations, however, we were concerned about over-fitting models to the data. (See discussion of over-fitting in the context of fish analysis, Section 5.2.1, subsection on Green Bay Zone 2.) Thus, we included the curvilinear terms (E^2 and N^2) only in the face of a visually apparent curvilinear trend in diagnostic plots (see below). We note that, regardless of their number, including appropriate spatial variables in a regression model increases the power to detect time trends notwithstanding a slight possibility that inappropriately including extra spatial terms could decrease power if there are correlations between space and time variables.

In addition to the spatial variables in the regression models, we introduced time as a simple linear term in all analyses. In each analysis, there was an insufficient number of distinct times of sampling to implement a curvilinear model for time. For this brief discussion, we considered a “distinct” time of sampling as a period of several months, or even a year, with at least two samples taken (see Figures A-44 through A-89, upper left panel). Of the 46 analyses ultimately carried out (specific combinations of deposit group and depth), 23 had observations at only two distinct points in time (e.g., 1989 and 1998), 20 had observations at three points in time, and 3 had observations at four points in time.

The dependent variable in all analyses was the \log_{10} PCB concentration with a companion variable indicating whether the observation was below the detection limit or was a detected concentration. We examined residual plots for all regression analyses to detect outliers and assess the assumption of normality. Table 5 notes the removal of only one exceptional value from the formal sediment regression analysis. This sample is considered in the context of time trends in the results section.

Table 5 Sample Removed from Time Trends Analysis

| Database ID | Reach | Original Deposit | Time Trends Deposit Group | Depth | Total PCBs (ppb) |
|-------------|---------|------------------|---------------------------|---------|------------------|
| A3_0-4 | De Pere | SMU56/57 | SMU Group 5067 | 0-10 cm | 99,000 |

Note:

Other PCB values range from 400 to 7,800 in this depth stratum and SMU group.

Figure 13 through Figure 17 show examples of plots we used to determine choice of linear or quadratic terms for northing and easting in the regression models. The plots also show log PCB concentration versus time and log PCB concentration versus depth. We added a “smoother” line to the plots to depict the general trend for these variables taken one at a time. As can be noted in some of the plots, a common structure of the deposit groups shows PCB concentrations rising from minima at one or both sides of the deposit group to a maximum in the middle (e.g., see Figure 17). The quadratic terms (E^2 and N^2) for northing and easting in the regression models capture this curvilinear trend. Separate plots evaluate each variable (time, depth, easting, northing), though a single regression model uses them all.

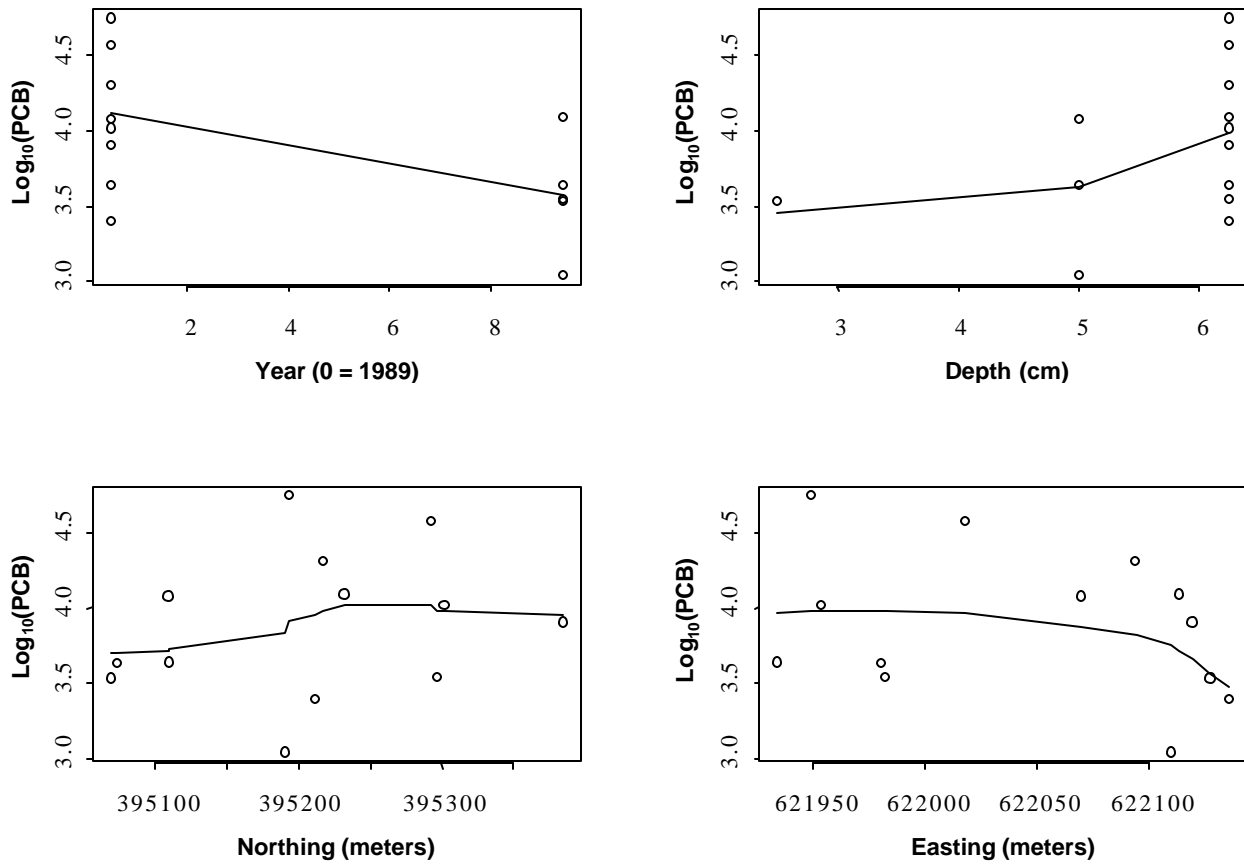


Figure 13 Log_{10} PCB Concentration versus Time, Depth, Northing and Easting for Little Lake Butte des Morts Deposit Group C (0 to 10 cm) Including Fitted Smoothed Line

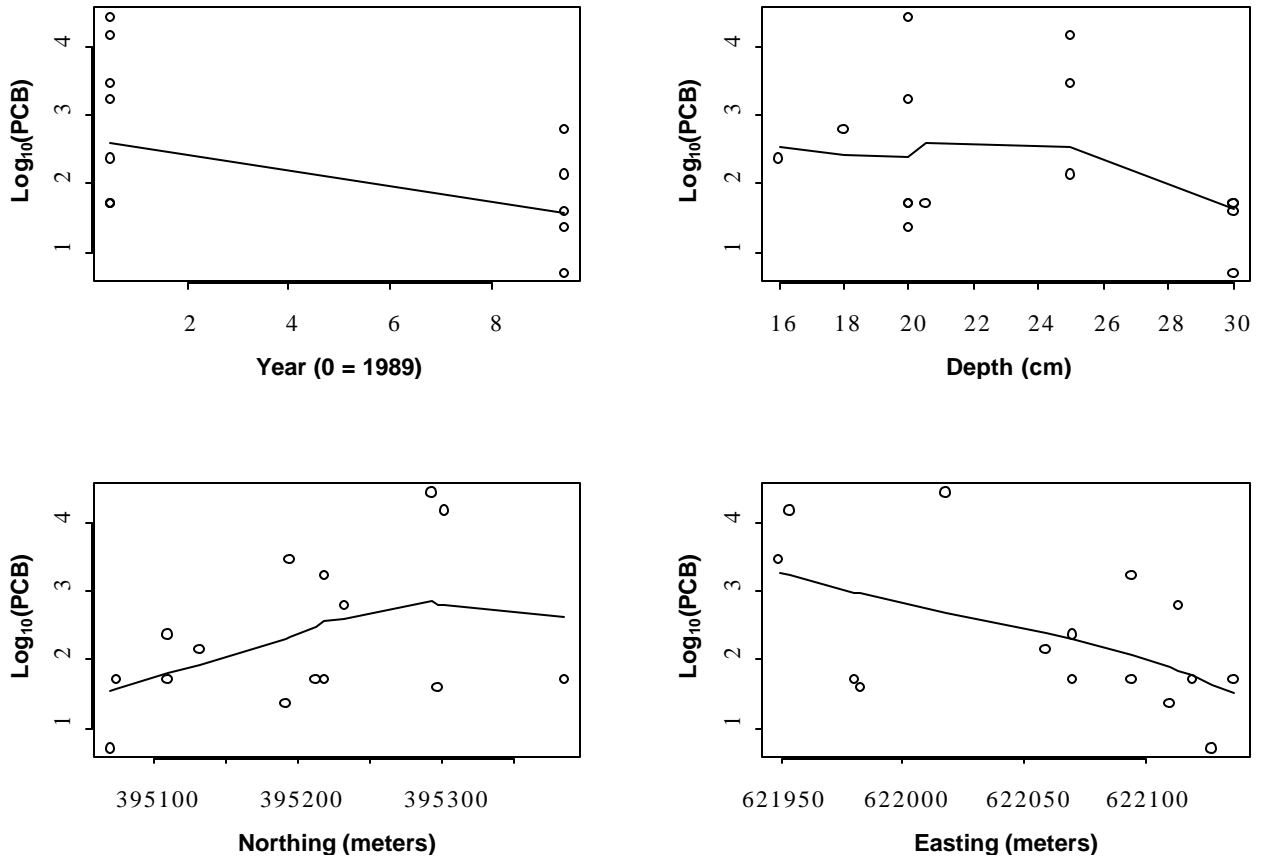


Figure 14 Log_{10} PCB Concentration versus Time, Depth, Northing and Easting for Little Lake Butte des Morts Deposit Group C (10 to 30 cm) Including Fitted Smoothed Line

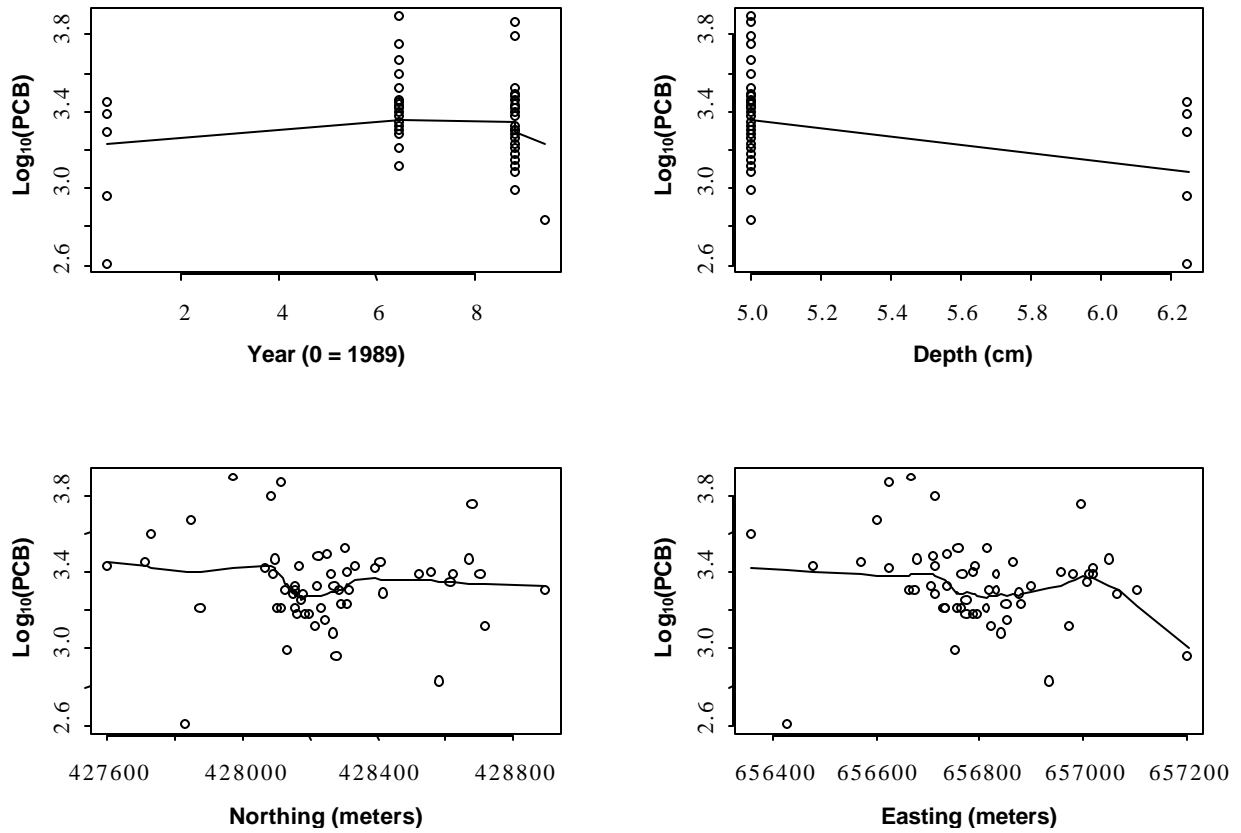


Figure 15 Log_{10} PCB Concentration versus Time, Depth, Northing and Easting for De Pere SMU Group 5067 (0 to 10 cm) Including Fitted Smoothed Line

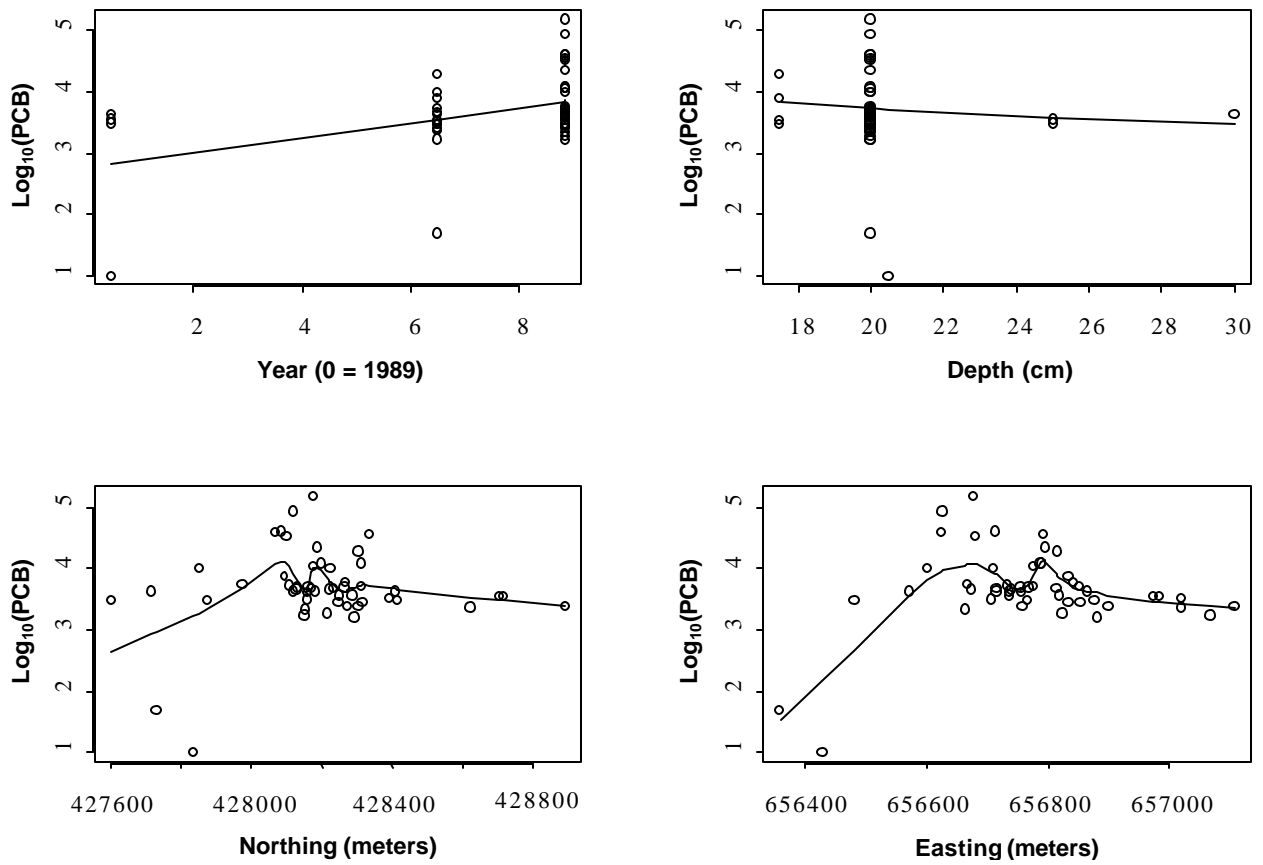


Figure 16 Log_{10} PCB Concentration versus Time, Depth, Northing and Easting for De Pere SMU Group 5067 (10 to 30 cm) Including Fitted Smoothed Line

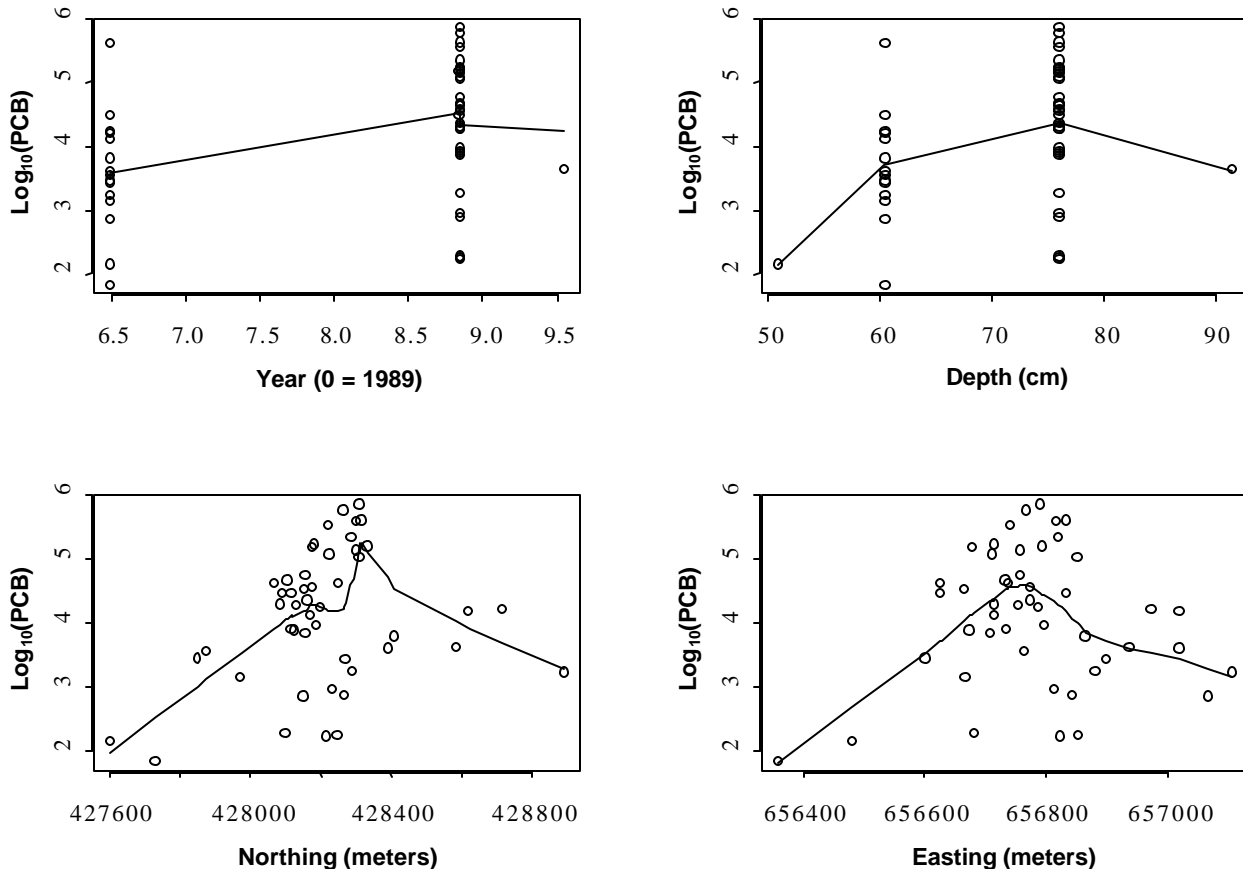


Figure 17 Log₁₀ PCB Concentration versus Time, Depth, Northing and Easting for De Pere SMU Group 5067 (50 to 100 cm) Including Fitted Smoothed Line

PCB concentration shows strong spatial variation, as shown on Figure 13 through Figure 17 and in the Appendix (i.e., space and PCBs are correlated). Controlling for spatial variation in the analysis allows for proper estimation of time trends in PCB concentrations. Similarly, the date and location of sampling may be correlated. These correlations can induce a spurious correlation between PCB concentration and time. This might happen, for example, if early samples were taken in the “hotter” location of a deposit (higher PCB concentrations) and later samples were drawn from a “cooler” location.

In order to determine the extent of the time-location correlation (which might create false time trends), we calculated the Pearson correlation coefficient between time and spatial variables. This correlation coefficient is +1.0 for perfect positive correlation, -1.0 for perfect negative correlation, and 0.0 (zero) if no correlation exists. We encountered a number of statistically significant correlations between the time that samples were drawn and either their depth

within the stratum, their easting (centered) or easting-squared coordinates, or northing or northing-squared. Among the 46 combinations of deposit group and depth we analyzed, 22 had statistically significant correlation coefficients between time and depth, eight between time and easting or easting-squared, and nine between time and northing or northing-squared. Among all the correlations between time and spatial coordinates, one-quarter were of magnitude 0.3 or larger, and 10 percent of the correlations were of magnitude 0.5 or larger (corresponding to a moderate correlation or stronger), with a maximum observed correlation of 0.97. These numerous non-zero correlations between time and the spatial variables show the importance of controlling for spatial variables, lest spatial trends in the time of sampling combine with spatial trends in PCB concentrations to induce false time trends in PCB concentrations.

The values of \log_{10} PCB also correlate with spatial coordinates. Again, among 46 analyzed combinations of deposit group and depth, six had statistically significant Pearson correlations between \log_{10} PCB and depth within the stratum, 18 between \log_{10} PCB and easting or easting-squared, and 10 between \log_{10} PCB and northing or nothing-squared. The 75th and 90th percentile and maximum of all of the correlations of \log_{10} PCB with spatial coordinates were of magnitude 0.3, 0.5 and 0.7, respectively. Peppered throughout these data are significant spatial trends either in time of sample acquisition or in PCB concentration. Thus, it behooves the analyst to include spatial variables in regression models for time trends of PCB concentrations in order to minimize the opportunity for a spatial trend in PCB concentration to masquerade as a time trend. (For purposes of exploring these correlations, concentrations below detection limits entered the analysis with the value of the detection limit. These limits and actual PCB concentrations were all log-transformed and used in the calculation of correlations.)

We also carried out an inspection of visual displays to detect glaring shifts over time in location of samples within a deposit group.

Figure 19 displays an example of these plots, showing northing and easting location of each sample, for each depth stratum, and for two time periods for Little Lake Butte des Morts Deposit Group AB. The key to interpretation of symbol size is included as Figure 18. Circles and squares indicate measured concentrations and concentrations below detection limits, respectively, and the size of the symbol indicates the magnitude of the PCB concentration. The upper row of the figure shows northing and easting location of each sample taken during 1989 through 1993 and the lower row corresponds to a later period, 1994 through 1999.

Working through the 0- to 10-cm plots (Figure 19, upper and lower left panels) as an example will help to clarify the role of space and its interaction with PCB concentrations and time. This is intended as a descriptive exploration. Note that in the 0- to 10-cm stratum, a larger fraction of early samples (upper panel, 1989–1993) occurs in the north of the deposit group than samples taken in the later

period (lower panel, 1994–1999). Correlation coefficients can help to summarize such trends. The Pearson correlation coefficient ranges from $r = -1$ (perfect negative association) to $r = +1$ (perfect positive association). In a scatter plot, when $r = +1$, all points would fall on an upward sloping straight line. A correlation of $r = 0$ means no association between two variables. The correlation of the time of sampling and the northing coordinate is $r = -0.3$ ($p = 0.02$, statistically significant), indicating that sampling locations have a southward trend across the deposit over time. The correlation coefficient is negative because later (“larger”) sampling times tend to occur with smaller northing coordinates. Smaller northing coordinates are farther south than larger ones. Also, earlier samples (upper plot) spread out more in the east and west directions than the samples from the later period (lower plot). The statistically significant correlation of -0.3 ($p = 0.03$) between time of sampling and the centered easting-squared term provides evidence for this. Over time, therefore, the sampling effort became more concentrated toward the south and west-center of this deposit group. This shift readily appears by comparing the upper and lower panels of Figure 19. In statistical parlance, time and spatial coordinates are confounded (and correlated). It is important to control for one when examining the role of the other.

We also found strong and highly significant spatial trends in \log_{10} PCB concentrations. The correlation between \log_{10} PCB concentration and easting is $r = -0.6$ ($p < 0.0001$). The negative correlation indicates that PCB concentration generally decreases from west to east. The correlation is $r = -0.5$ ($p < 0.0001$) for easting-squared, meaning that PCB concentrations decrease from the middle of the deposit to the east and west. The correlations of $r = -0.5$ ($p < 0.0001$) for northing, and $r = -0.6$ ($p < 0.0001$) for northing-squared, have similar interpretations to those just offered. The strong correlation of PCB concentration with linear and curvilinear (quadratic) spatial dimensions suggests a deposit group with a peak concentration near one edge of the area sampled. Concentrations taper off on all sides, but particularly to the east and north. In the upper plot for the 0- to 10-cm stratum (still Figure 19), the smaller circles toward the upper right corroborate this trend. Given that the PCB concentrations in the 0- to 10-cm stratum of Little Lake Butte des Morts Deposit Group AB have a distinct spatial structure, we have incorporated that structure in our model for a time trend in this deposit group. We also note that Figure 19 presents two time periods although time in the continuous form has been used in the analysis of time trends.

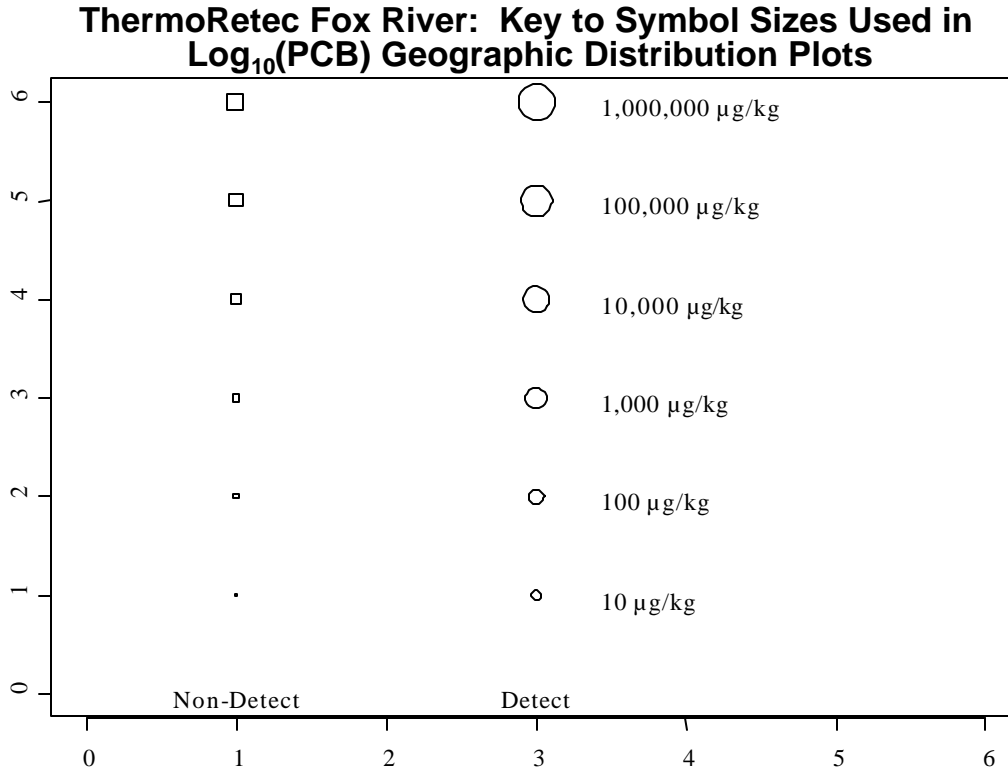


Figure 18 Northing/Easting Plot Key

Scale plot showing the size of circles (for samples with detected PCBs) and squares (for samples with PCBs below detection limit, the square conveys the level at which the PCBs would have been detected as reported by the various testing agencies) used to convey total PCB concentration in the northing/easting plots of sample locations.

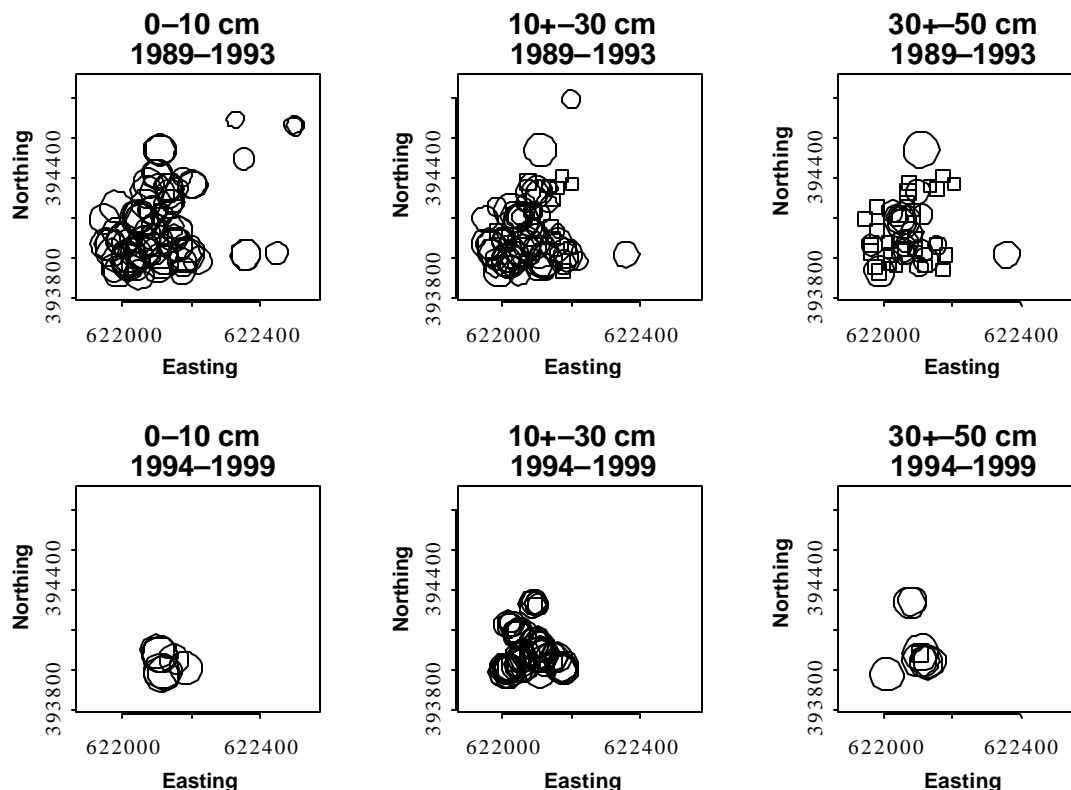


Figure 19 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group AB

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

3 Methods for Fish Analysis

For the reasons discussed earlier (“Logarithmic Transformation,” Section 2.1.3), we used the log of PCB concentration as the outcome variable in all the regression models fitted. There are good reasons for using the log transformation. Expressing rate of change as percent change per year has more meaning than absolute change in concentration, which can lead to absurd negative concentration predictions. An analysis on the \log_{10} scale corresponds to modeling percent change. The data have an approximately normal distribution on the log scale, but a strongly skewed distribution on the original scale.

We included two potential confounding factors in all regression models for \log_{10} PCB concentration versus time: percent lipid in the sample by weight and seasonality. As described below in the results section, both of these factors added significantly to prediction of PCB concentrations in most analyses. The following paragraphs describe how we incorporated these two factors into the models. We could not introduce any procedures to handle spatial dependence of fish data due to the lack of easting and northing coordinates for the fish samples in each reach. Not being able to model or investigate spatial dependence of fish samples does not imply the absence of such dependence. We simply have no means to study or address it. Because fish move more than sediments do, we expect that fish samples are closer to independent than sediment samples.

3.1 Lipid Normalization

Analyses of PCB concentration in fish often utilize “lipid normalization” in order to account for the relationship between PCB concentration and percent lipid in fish tissue. PCBs tend to concentrate in fat tissue so that, in general, fatter fish have higher concentrations of PCBs per total weight than leaner fish. The direct lipid normalization commonly used consists of dividing the PCB concentration by the percent lipid content (by weight) of the sample. This results in a variable showing PCB concentration per unit weight of lipid. We have chosen a somewhat different approach, similar to that of Larsson *et al.* (1993) and Herbert *et al.* (1995). We regard the lipid variable as an independent variable rather than a direct divisor of the PCB concentration. This approach allows the data itself to specify the relationship of PCBs to lipids. The model we use is:

Equation 3

$$\log(\text{PCB}) = b_0 + b_1 \log_{10}(\text{lipid}) + \dots + e$$

where

- b_0 = the intercept term,
- b_1 = the regression coefficient on log of percent lipid,
- e = random error, and

additional variables such as time are included in the model as well (the time variable is considered below).

This model yields a predicted value for PCB concentration per unit tissue weight. Since the public consumes fish tissues (rather than just the lipid in the tissue), this offers a more useful prediction in many applications than the other normalization based on PCB per unit of lipid content.

An interesting fact should be noted about this model for PCB concentration (Equation 3). The model can be directly compared to the traditional “lipid normalization.” Subtracting $\log(\text{lipid})$ from both sides of Equation 3 gives the equivalent model:

Equation 4

$$\log_{10}\left(\frac{PCB}{lipid}\right) = b_0 + (b_1 - 1)\log_{10}(lipid) + \dots + e$$

Comparing the two (Equation 3 and Equation 4), one clearly sees that as long as we treat \log of percent lipid as a predictor, then it does not really matter whether we lipid-normalize the PCB concentration on the left-hand side of the equation or use PCB concentration without lipid normalization. Except for the coefficient for \log of percent lipid differing by 1, all other coefficient and standard error estimates will remain unchanged. An analysis that has \log of lipid-normalized PCB concentration on the left-hand side (such as in Equation 4), but does not include \log of percent lipid on the right-hand side, amounts to forcing b_1 to be 1, so that $b_1 - 1$ will be zero. If direct lipid normalization represents the best model for the observed data, then we will estimate b_1 close to 1 in the regression approach. It is an advantage of the regression approach, model 3, that it will reduce to the direct lipid normalization if that is the correct model for the data considered in a given analysis. If PCB concentration and percent lipid do not have a directly proportional relationship, then we will estimate b_1 as something different than 1 (usually less than 1, as seen in the results below).

3.2 Seasonality

To account for the possibility that PCB concentration may vary by time of year, we incorporated into the model a sine curve as a function of the time of year.

Equation 5

$$\log_{10}(PCB) = b_0 + b_1 \log_{10}(lipid) + b_2 \sin(2\pi t^*) + b_3 \cos(2\pi t^*) + \dots + e$$

where

t^* = time of year expressed as a fraction between 0 and 1.

Trigonometry shows that the weighted sum of the sine and cosine function in this equation gives a sine curve with a maximum at the time $\arctangent(-b_2/b_3)$ and

an amplitude equal to $(b_2^2 + b_3^2)^{0.5}$. We present these more meaningful quantities, time of maximum and amplitude, in our results tables rather than the more abstract b_1 and b_2 . The time of maximum is coded to range from 1.0 (beginning of January) to 12.999... (end of December).

We note that the true seasonal cycle in PCB concentration may not be sinusoidal. Albeit likely, the presence of some average annual pattern of rise and fall of PCB concentration may not have the shape or smoothness of a sine curve. Nevertheless, the sine curve can serve as an approximation to seasonal variation. The statistical significance of the fitted sine curve (described later) strongly suggests that this simple function helps to capture and control seasonal variation in PCB concentration in fish.

Prior to model fitting, we centered the log of percent lipid variables. This step is analogous to the centering of northing and easting coordinates described earlier in the methodology section for sediment analysis. For each combination of reach, species, and sample type, we subtracted the mean \log_{10} lipid percent within that reach/species/type from the \log_{10} lipid value for each sample. Table 22 of the results displays these mean values. We also centered the sine and cosine terms by subtracting off the value of the sine and cosine variables at midyear (i.e., July 1). The advantage of the centering is that for forward projection we need only use the intercept and slope coefficient from the fitted models for PCB concentrations. Then, when using the intercept term and the coefficient on final slope to predict values of PCB at future time points, we are estimating the PCB concentration for a fish with average lipid content sampled on July 1. For numerical stability in estimating the slope coefficient for time, we centered time at the beginning of 1989 by subtracting January 1, 1989 from each sample date.

3.3 Time Trend Models

The simplest model for time trend in PCB concentration is a linear relationship between log of PCB concentration and time. A negative slope corresponds to an exponential decay in PCB concentration at a constant rate (for example, 5 percent per year). The first step in our analysis involved testing whether this simple model fit the data well, for each unique combination of reach, species and sample type (whole body, or fillet with skin). In statistical terms, this means testing the null hypothesis of a constant exponential rate of decay over all years versus the alternative of decay rate that is not constant over time. To perform such a hypothesis test, one must specify an alternative model, which we consider to be a competing model for the change in PCB concentration over time.

The simple linear model has the following equation:

Equation 6

$$\log_{10}(PCB) = b_0 + b_1 \log_{10}(lipid) + b_2 \sin(2\pi t^*) + b_3 \cos(2\pi t^*) + b_5 time + e$$

We modeled the alternative nonlinear model as a two-slope model in the form of a linear spline, which appears as two straight lines joined at a kink, or breakpoint (Cressie, 1993). This is modeled in a linear regression equation as:

Equation 7

$$\log_{10}(PCB) = b_0 + b_1 \log_{10}(lipid) + b_2 \sin(2\pi t^*) + b_3 \cos(2\pi t^*) + b_4 early + b_5 time + e$$

The variable *early* equals (*time* – *breakpoint*) if time predates the breakpoint and 0 for time after the breakpoint. The coefficient of *time* (b_5) represents the slope of log PCB concentration versus time after the breakpoint, and the coefficient of *early* (b_4) measures how much the early slope differs from the late slope. That is, the early slope equals $b_4 + b_5$ and the late slope equals b_5 .

This model offers simplicity and intuitive clarity: it means that PCBs were changing at two different constant rates of change—one before and one after the breakpoint. This model has been applied to PCB and DDT concentrations in fish in the Great Lakes (De Vault *et al.*, 1996). A visual inspection of scatter plots of log PCB versus time shows that, for many reach/species/type combinations, this model gives a good representation of the pattern apparent in the data. Since the model incorporates a constant rate of change after the breakpoint (coefficient b_5), it facilitates straightforward projections of concentrations into the future.

One could fit more complex models to the data. Given the fairly small number of distinct time points at which data were collected for each reach/species/type combination, however, one can not reliably fit models containing many parameters used to describe the time effect. The linear spline model, which includes a seasonal time effect, already uses five parameters explicitly modeling change with time: two seasonal terms (sine and cosine), early and late slope, and the location of the breakpoint.

3.4 Model Fitting and Hypothesis Testing

Fitting models and testing hypotheses involved several analyses. The first key steps were: 1) finding the best-fitting linear spline model, 2) determining if the spline model (Equation 7) offered a significant improvement over a simple linear model (Equation 6), and 3) choosing a spline or simple linear model accordingly.

If the breakpoint is specified, Equation 7 is a linear regression model that can be fitted using standard statistical software that accommodates concentrations below the detection limit. We used the SPLUS procedure *CensorReg* for this analysis. As described earlier for sediment samples, *CensorReg* uses the maximum likelihood method to estimate parameters in the model while correctly accounting for the values below the detection limit. In order to find the optimal location of the breakpoint, we fit models using different possible breakpoint locations. To reduce the computation time required to a manageable level, we considered only one breakpoint per year, on January 1 for each year across the range of data. For all analyses, the 1-year span of uncertainty in the breakpoint is

small compared to the total range of the observations over time. We considered only breakpoint locations that provided data extending at least 2 years on both sides of the breakpoint. This 2-year rule would provide at least a minimum of data needed to calculate slopes before and after the breakpoint. The best linear spline model, including the optimum breakpoint location was determined using the maximum likelihood method.

The best linear spline model (Equation 7) and the simple model (Equation 6) were compared and a choice between them was made, as follows:

In comparing the two models using the maximum likelihood method, a quantity called the “deviance” is calculated. The change in deviance relates to the change in probability (i.e., improvement in fit) when extra parameters are added to a model. For a given model, the deviance is $-2 * \log(L)$, where L is the likelihood of the model, given the data, as described in the sediment methods section.

The linear spline model (Equation 7) has two additional parameters compared to the simple linear model (Equation 6)—the location of the breakpoint and the early slope difference. Under the null hypothesis, the spline model would not be a true improvement over the simple linear model. The difference in deviance between the linear model and the best linear spline model should have a chi-square distribution with two degrees of freedom if the null hypothesis is true. If the chi-square test statistic is too large, we reject the null hypothesis and accept the spline model. The spline model, if selected, includes the parameter estimates in Equation 7 and their standard errors and p -values based on the likelihood method. A small chi-square value prompts selection of Equation 6.

If we know the true location of the breakpoint, the method behind the S-PLUS procedure *CensorReg* produces correct standard errors and p -values for slopes and other parameters in the spline model, which are reported in the tables. As the breakpoint is not known with absolute certainty, the data are used to estimate it. Thus, the reported standard errors and p -values for the intercept, time trend slopes, and other coefficients in a model based on Equation 7 do not account for the additional variance due to the estimated breakpoint location. Without compensating for the uncertainty in the breakpoint, the p -values and standard errors for other parameters are too small. Through bootstrapping, we could compute more accurate standard errors. We did not use the quite computer-intensive bootstrap given the resources available to the project. Instead, we used a more informal sensitivity analysis to determine the role of the breakpoint in slope estimates. This analysis tells us how sensitive the conclusions concerning time trend slopes are to shifts in the breakpoint.

As part of the breakpoint sensitivity analysis, we initially created a plausible range of breakpoints for those combinations of species, reach, and sample type where a spline model (Equation 7) fit significantly better than the simple linear model (Equation 6). We considered as plausible all breakpoints having a value of the likelihood that was close to the value of the likelihood at the best breakpoint,

in that they fit the data almost as well as the best breakpoint. Formally, we settled on the plausible range of breakpoints as starting from the earliest and ending at the latest breakpoint year with a deviance within 3.84 of the best model. The value 3.84 corresponds to a p -value of 0.05 for a chi-square test with one degree of freedom and is analogous to testing whether the alternative breakpoint (and its associated early and late slopes and other parameters) fits the data significantly worse than the best breakpoint.

3.5 Testing for a Constant versus a Changing Final Slope

The fitted models assume that PCB decreases at a constant rate on the log scale (i.e., linear on the log scale) after the breakpoint, or for the entire range if there is no breakpoint. We tested the appropriateness of this assumption by fitting a model that includes a quadratic term in time for the interval after the breakpoint. This analysis simply adds a term to Equation 6 that is $b_6 \cdot (time^2)$ for time after breakpoint or $b_6 \cdot (0)$ for time before the breakpoint. This model allows for a curved rather than a linear relationship of log PCB concentration with time. A significant p -value for this quadratic term indicates that the curved model fits better than the model that assumes linearity after the breakpoint. Testing the quadratic model addresses the simple question: are the data consistent with a constant rate of change after the breakpoint (or entire range if there is no fitted breakpoint) or do the data imply a changing rate?

3.6 Meta Analyses—Combining Data on All Species Within a Reach

After completing all of the model fitting and hypothesis testing for each of the reach/species/type combinations, we performed analyses that combined results from all the species/type combinations within each reach. Three groups of hypothesis tests of interest emerged. The first group involved testing the null hypothesis that a simple linear model, without a breakpoint, for every species/type fits just as well as a spline model for all species/types within a reach. Formally, we accomplished this by summing up the chi-square statistics from the linear versus spline tests for each of the species/type combinations within the reach, and then comparing this sum to a chi-square distribution with degrees of freedom equal to twice the number of species/types combinations in the reach.

The second group of hypothesis tests is actually a single test. We tested the null hypothesis that the final slope is zero for all species/types in the reach versus the alternative that one or more species/types have a negative or positive slope. We accomplished this by first computing the directional or one-tailed p -value for each species/type. That is, p is close to 0 for large negative slopes and close to 1 for large positive slopes. Then, for each species/type within the reach we computed the statistic $X^2 = -2 \log(p - value)$, where \log is the natural log. Under the null hypothesis, X^2 has a chi-square distribution with two degrees of freedom.

Thus, summing up the X^2 values within a reach gives a quantity that should, under the null hypothesis that all final slopes are zero, have a chi-square distribution with degrees of freedom equal to twice the number of species/type combinations within the reach. We converted the statistic to a two-tailed p -value by counting either very large values or values very close to zero as rejecting the null hypothesis. These correspond to evidence for an overall negative or overall positive slope, respectively.

An average final slope estimate for the reach was defined as a weighted average of the final slope estimates for each species/type combination, where the weight was the inverse of the square of the standard error of the slope coefficient estimate. Thus, slope estimates with great precision (low standard error) have more weight than imprecise ones (high standard error). This weighting minimizes the variance of the resulting combined estimate and proves optimal if all of the true final slopes are in fact identical.

The third group of hypothesis tests examined the null hypothesis that the final slope is constant over time versus the curved alternative that the slope changes over time. We followed a similar procedure to that just described for testing for a zero final slope, since the null hypothesis corresponds to the coefficient on the quadratic term being zero. A positive coefficient on the quadratic term means the slope either curves upward or plateaus over time (on the log scale), while a negative coefficient means the slope curves downward or steepens over time.

3.7 Projecting into the Future

Predictions of concentration of PCBs in future years assumed that PCB concentration continues to decrease (or increase) at a constant rate, which is the final slope or the slope after the breakpoint. Based on this assumption, we can compute the estimate of the mean of log (PCB concentration) from the coefficients in Equation 6 or Equation 7a:

Equation 8

$$E[\log(\text{PCB at time})] = b_0 + b_3 \text{ time}$$

where E indicates the expected value and time is years since 1989, the year at which time was centered prior to fitting the model. The formula predicts the mean of log (PCB) for a fish with average percent lipid content sampled on July 1 of the year, as long as the year follows the breakpoint. We obtain this formula from Equation 6 or Equation 7 by setting all other covariates in the model equal to zero. Since we centered log (*lipid*) at its mean, a zero value for the centered lipid variable is the same as setting log (*lipid*) equal to its mean. The seasonal variables and sine and cosine of time were centered at zero on July 1. The variable *early* in Equation 7 equals zero for all times after the breakpoint.

One computes the confidence interval for this predicted mean by first calculating the standard error:

Equation 9

$$SE(\text{predicted mean at year } t) = \sqrt{[SE(b_0)]^2 + t^2 [SE(b_5)]^2 + 2t \text{cov}(b_0, b_5)}$$

where $\text{cov}(b_0, b_5)$ denotes the covariance between these two parameter estimates, b_0 and b_5 from Equation 6 or Equation 7. The predicted mean plus or minus twice the standard error gives the 95 percent confidence interval on the log scale.

One can convert the predicted mean on the log scale to an estimate of the mean on the original scale (i.e., ppb) by the formula:

Equation 10

$$E(\text{PCB at time}) = 10^{(E(\log(\text{PCB at time})) + (MSE \div 2))}$$

where MSE is the mean squared error from the regression model on the natural log scale and is an estimate of the residual variance around the fitted regression model. This is just the formula for the mean of a lognormal distribution, based on the mean and variance on the log scale. We applied this formula to the predicted mean on the log scale and the lower and upper bounds of the confidence interval on the log scale in order to get the mean and confidence interval on the original (ppb) scale. This confidence interval does not consider the variance due to estimating the location of the breakpoint. A confidence interval that corrects for breakpoint estimation could be wider.

We also computed predicted time until mean PCB concentration reaches a specified concentration, G . The formula is:

Equation 11

$$\text{time to specified concentration } (G) = \frac{(\log_e(G) - b_0 - (MSE \div 2))}{b_5}$$

where

- G = the specified level of PCB concentration in ppb,
- $time$ = time until that level is reached, in years since 1989,
- MSE = mean squared error from a regression model fit to \log_e of PCB concentration,
- b_0 = intercept from Equation 6 or Equation 7, and
- b_5 = coefficient of $time$ from Equation 6 or Equation 7.

Computing confidence intervals for the predicted time to reach a specified level would seriously complicate our analysis, so we did not attempt to do so. A confidence interval based on the estimated standard errors would be wide and one that correctly accounted for the uncertainty due to estimating the breakpoint would be exceptionally wide. Therefore, we regard these “time to specified level” estimates as very uncertain.

In addition to the need to account for variance due to estimating the location of the breakpoint, the predictions are uncertain for yet another reason. Predictions of concentration of PCBs in future years assume that PCB concentration continues to decrease (or increase) at a constant rate. One cannot test this assumption except to continue collecting data in future years. Moreover, the assumption of a constant rate of change may not be very reasonable. A positive final slope, for example, implies that the PCB concentration continues to increase “forever” to higher and higher levels, an absurd conclusion. A negative final slope means that PCB concentration continues to decline to values near zero. But a scouring event that uncovered buried sediment more contaminated than surface sediment would likely lead to an increase in PCB concentration at the surface. Also, even a decreasing rate may level off well above a PCB concentration of zero. These future projections depend for their validity on an unverifiable future steady state.

4 Sediment Results

4.1 Number of Observations

A total of 1,980 observations (core-averaged) were initially available for analysis. Table 6 shows the distribution of these observations by our deposit group designation and depth. Due to the requirement of a sufficient number of observations and a sufficient time spread for an appropriate time trend analysis, only 1,618 samples qualified for the time trend analysis (Table 7). The reasons for dropping particular depth strata in specific deposit groups are explained in Table 8. Over one-third of the 1,618 usable observations occurred in the upper 10 cm of sediment, approximately one-third in the 10- to 30-cm stratum, about one-eighth in the 30- to 50-cm stratum, and the balance at greater depths. The greatest fraction of unusable data (due to lack of sufficient number of observations or lack of sufficient time spread) occurred at depths of 30 cm or lower, where approximately one-third of the core-averaged observations were unusable.

The fraction of observations below detection limit (BDL) varied widely by reach, deposit group, and depth, from a minimum of 0 percent (no BDL observations) to a maximum of 82 percent BDL observations. A majority of analyses included 20 percent or fewer BDL observations. The fraction of BDL observations, however, sufficiently requires the use of the maximum likelihood (ML) methods noted earlier. The number and percent of BDL observations by deposit group and depth is included in an appendix table. As noted in Section 2, all observations available for a given deposit group and depth stratum were included in the calculation of time trends. Due to the use of ML methods, BDL observations were neither modified nor excluded.

Table 6 Sample Size by Deposit Group and Depth after Core Averaging

| TMWL Deposit Group | Sample Average Depth (cm) | | | | | Total |
|------------------------------------|---------------------------|------------|------------|------------|------------|--------------|
| | 0-10 | 10+-30 | 30+-50 | 50+-100 | 100+ | |
| <i>Little Lake Butte des Morts</i> | | | | | | |
| Deposit Group AB | 67 | 105 | 54 | 12 | 2 | 240 |
| Deposit Group C | 13 | 15 | 8 | 2 | 0 | 38 |
| Deposit Group POG | 13 | 10 | 4 | 3 | 2 | 32 |
| Deposit Group D | 18 | 15 | 9 | 6 | 0 | 48 |
| Deposit Group E | 6 | 7 | 21 | 14 | 2 | 50 |
| Deposit Group F | 29 | 28 | 10 | 2 | 2 | 71 |
| Deposit Group GH | 15 | 12 | 3 | 0 | 0 | 30 |
| <i>Appleton</i> | | | | | | |
| Deposit Group IMOR | 18 | 15 | 9 | 3 | 1 | 46 |
| Deposit Group N Pre-dredge | 51 | 40 | 18 | 4 | 0 | 113 |
| Deposit Group VCC | 41 | 34 | 17 | 9 | 3 | 104 |
| <i>Little Rapids</i> | | | | | | |
| Deposit Group Upper EE | 31 | 25 | 13 | 3 | 1 | 73 |
| Deposit Group Lower EE | 30 | 33 | 13 | 5 | 3 | 84 |
| Deposit Group FF | 32 | 31 | 8 | 0 | 0 | 71 |
| Deposit Group GGHH | 49 | 45 | 75 | 54 | 36 | 259 |
| <i>De Pere</i> | | | | | | |
| SMU Group 2025 | 43 | 31 | 13 | 30 | 25 | 142 |
| SMU Group 2649 | 66 | 48 | 10 | 46 | 45 | 215 |
| SMU Group 5067 | 57* | 51 | 34 | 48 | 50 | 240 |
| SMU Group 6891 | 20 | 18 | 2 | 16 | 15 | 71 |
| SMU Group 92115 | 27 | 15 | 3 | 7 | 1 | 53 |
| Total: | 626 | 578 | 324 | 264 | 188 | 1,980 |

Note:

* One additional sample, A3_0-4, not included in these sample sizes, had an exceptionally large PCB concentration and was considered separately.

Table 7 Sample Size by Deposit Group and Depth Included in Time Trends Analysis, after Core Averaging

| TMWL Deposit Group | Sample Average Depth (cm) | | | | | Total |
|------------------------------------|---------------------------|------------|------------|------------|------------|--------------|
| | 0-10 | 10+30 | 30+50 | 50+100 | 100+ | |
| <i>Little Lake Butte des Morts</i> | | | | | | |
| Deposit Group AB | 67 | 105 | 54 | — | — | 226 |
| Deposit Group C | 13 | 15 | — | — | — | 28 |
| Deposit Group POG | 13 | — | — | — | — | 13 |
| Deposit Group D | 18 | 15 | — | — | — | 33 |
| Deposit Group F | 29 | 28 | — | — | — | 57 |
| Deposit Group GH | 15 | — | — | — | — | 15 |
| <i>Appleton</i> | | | | | | |
| Deposit Group IMOR | 18 | — | — | — | — | 18 |
| Deposit Group N Pre-dredge | 32 | 27 | 17 | — | — | 76 |
| Deposit Group VCC | 41 | 34 | 17 | — | — | 92 |
| <i>Little Rapids</i> | | | | | | |
| Deposit Group Upper EE | 31 | 25 | 13 | — | — | 69 |
| Deposit Group Lower EE | 30 | 33 | 13 | — | — | 76 |
| Deposit Group FF | 32 | 31 | — | — | — | 63 |
| Deposit Group GGHH | 49 | 45 | 75 | 54 | 36 | 259 |
| <i>De Pere</i> | | | | | | |
| SMU Group 2025 | 43 | 31 | 13 | 30 | — | 117 |
| SMU Group 2649 | 66 | 48 | — | 46 | 45 | 205 |
| SMU Group 5067 | 57* | 51 | — | 48 | 50 | 206 |
| SMU Group 6891 | 20 | 18 | — | — | — | 38 |
| SMU Group 92115 | 27 | — | — | — | — | 27 |
| Total: | 601 | 506 | 202 | 178 | 131 | 1,618 |

Note:

* One additional sample, A3_0-4, not included in these sample sizes, had an exceptionally large PCB concentration and was considered separately.

A dash, “—,” indicates that the particular cell could not be analyzed for time trends. An explanation is provided in Table 8.

Table 8 Deposit Groups Analyzed, Or Reasons for No Analysis

| TMWL Deposit Group | Sample Average Depth (cm) | | | | | Total Yes |
|------------------------------------|---------------------------|-----------|----------|----------|----------|-----------|
| | 0-10 | 10+-30 | 30+-50 | 50+-100 | 100+ | |
| <i>Little Lake Butte des Morts</i> | | | | | | |
| Deposit Group AB | Yes | Yes | Yes | I, T | I, T | 3 |
| Deposit Group C | Yes | Yes | I | I, T | N | 2 |
| Deposit Group POG | Yes | I, T | I, T | I, T | I, T | 1 |
| Deposit Group D | Yes | Yes | I, T | I, T | N | 2 |
| Deposit Group E | I, T | I | T | I, T | I, T | 0 |
| Deposit Group F | Yes | Yes | I, T | I, T | I, T | 2 |
| Deposit Group GH | Yes | I, T | I, T | N | N | 1 |
| <i>Appleton</i> | | | | | | |
| Deposit Group IMOR | Yes | T | I, T | I, T | I, T | 1 |
| Deposit Group N Pre-dredge | Yes | Yes | Yes | I, T | N | 3 |
| Deposit Group SU | T | I, T | I, T | N | N | 0 |
| Deposit Group VCC | Yes | Yes | Yes | I, T | I, T | 3 |
| Deposit Sample POG | I, T | I, T | I, T | I, T | I, T | 0 |
| Deposit Group DD | I, T | I, T | I, T | N | N | 0 |
| <i>Little Rapids</i> | | | | | | |
| Deposit Group Upper EE | Yes | Yes | Yes | I, T | I, T | 3 |
| Deposit Group Lower EE | Yes | Yes | Yes | I, T | I, T | 3 |
| Deposit Group FF | Yes | Yes | I | N | N | 2 |
| Deposit Group GGHH | Yes | Yes | Yes | Yes | Yes | 5 |
| <i>De Pere</i> | | | | | | |
| SMU Group 2025 | Yes | Yes | Yes | Yes | T | 4 |
| SMU Group 2649 | Yes | Yes | T | Yes | Yes | 4 |
| SMU Group 5067 | Yes | Yes | T | Yes | Yes | 4 |
| SMU Group 6891 | Yes | Yes | I, T | I, T | I, T | 2 |
| SMU Group 92115 | Yes | T | I, T | I, T | I, T | 1 |
| Total Yes: | 18 | 14 | 7 | 4 | 3 | 46 |

Notes:

- Yes - Deposit groups and depths with sufficient data to perform a time trend analysis.
- I - Insufficient data (fewer than 10 observations).
- N - No observations.
- T - No time variation. Need at least two measured PCB concentrations (not below detection limits) at each of two distinct times.

4.2 Geographic Groups for Time Trend Analysis

As noted earlier, we regrouped the data into more compact geographic deposits (deposit groups, noted in Table 1 through Table 4). The majority of the original deposit designations transferred primarily, but not always wholly, into one of our time trend deposit groups. The exceptions, where a geographically extensive original deposit was broken into a number of separate groups for analysis, included Little Lake Butte des Morts Deposit E (which became our Little Lake Butte des Morts deposit groups E, F, and GH) and Little Rapids Deposit EE (which became our Little Rapids deposit groups Upper EE, Lower EE, FF and

GGHH). In addition, a number of observations in the database supplied to us had no deposit designation in the database supplied to us (e.g., noted as “No Designation,” Table 1 through Table 4), and were allocated to one of our deposit groups based on location. As noted in Table 1 through Table 4, we were able to include a substantial number of observations in the time trends analysis by forming new deposit groups. For example, in the De Pere Reach, we analyzed 731 observations (Table 4) that had no deposit designation in the FRDB. The result of our grouping for time trend analysis is captured by Figure 5 through Figure 12. As can be seen from the plot, the deposit groups are fairly compact.

The data analyzed included diverse spatial configurations. An illustration of the variety of geographic configurations can be found on Figures A-1 through A-43 (see Appendix), an example of which can be found on Figure 19. The description and interpretation of the plot were presented earlier. The plot demonstrates how the geographic configuration is not necessarily the same for the two time periods, illustrating the importance of controlling for geography in analyzing time trends. By failing to control for sample geography, an apparent time trend could simply be due to sampling from, for example, a high concentration area in an earlier period and a lower concentration area in a later period without any real shift in concentration in either area over time. The figures show measured concentrations and concentrations below detection limits as circles and squares, respectively, with the magnitude of the PCB concentration indicated by the size of the square or circle.

4.3 Time Trends in Sediment Concentrations

Time trends in PCB concentrations differ both by depth and by deposit group. Appendix Table A-1 presents detailed numerical results, sections of which are reproduced here in Table 9 for 46 different analyses, representing different deposit groups and depths. The key results from the table are:

- Coefficient of the time term (this parameter represents the slope estimate on a \log_{10} scale as rate of change in \log_{10} PCB concentration per year),
- Standard error of the time coefficient based on the window subsampling empirical variance (WSEV) method,
- The annual percentage rate of change (compounded), and
- The p -value for the null hypothesis that the true slope is zero ($b_t = 0$ in Equation 2, Section 2.6). The “statistically significant” slopes are also designated by asterisk(s) in the table. The deposit group and depth combinations that are “statistically significant” will very likely have true non-zero rates of change over time.

Statistical significance plays an important role in interpreting Table 9 and other tables presenting rates of change. The p -value column in this and other tables shows the degree of statistical significance of the calculated rate of change of log PCB concentration versus time. The p -value, which constitutes the numeric statement of statistical significance, quantifies the strength of the evidence against the null hypothesis that the true rate of change is zero. The closer the p -value is to zero, the more confidence we have that the true rate of change is not zero. Formally, the p -value is defined as the probability of observing a result as or more extreme than that actually observed if the null were, in fact, true. More explicitly, the p -value can be interpreted as the outcome of the following hypothetical experiment. We can imagine taking samples from a deposit group whose **true** rate of change is zero and repeating this operation many times. For example, Little Lake Butte des Morts Deposit Group AB has $n = 67$ samples at 0 to 10 cm depth. We would take many samples of size $n = 67$ from the deposit group and analyze them as we have here, yielding one slope for each set of 67 samples. Due to random variation in sampling, each calculated slope would differ to a greater or lesser extent from the true slope. If the true slope were really zero, then these random slopes would have some distribution around zero. For any slope value that we choose or observe, we can look at the distribution and determine what fraction of our random slopes are as large or larger than a given slope. Usually, we take the fraction of slopes that are larger in either the positive or negative direction from the value. For example, for the slope of -0.097 , we would look at the fraction of random slopes smaller than -0.097 and larger than $+0.097$, because random variation can take us either in a positive or negative direction away from zero. The key concept is that if the true slope is really zero, the observed slope should not stray too far from zero. Traditionally (but with no other basis than that), $p < 0.05$ has been used to designate statistical significance. This p -value means that there are fewer than 5 chances in 100 that a slope as large or larger than that observed could have been generated by chance, if the **true** slope is zero. We adopt this definition and also designate $p < 0.05$ as “statistically significant.” In the tables, we note this with one asterisk and also use the following conventions: ** $p < 0.01$, *** $p < 0.001$.

In reality, one need not compute the hypothetical experiment to get the p -value. In fact, the p -values computed in Table 9 use the very standard t -test. As a conservative measure, we have chosen the degrees of freedom for the t -test as the number of grid cells with at least one sample, determined in the WSEV method described earlier. The number of non-empty grid cells is included in an appendix table.

We have also included in Table 9 a 95 percent confidence interval for the percent rate of change of the PCB concentration over time (derived from the slope and its standard error using the t -distribution with the same degrees of freedom as in the calculation of the p -value). We can state with 95 percent confidence that the true rate of change lies in this interval. If this interval is especially narrow, we have a very precise idea of the true rate of change. A particularly wide interval casts much doubt on the true rate of change.

Appendix Table A-1 presents the form of the linear regression model—either linear or quadratic, fitted to the data. “Linear” indicates that depth, easting, and northing are used as linear terms in the regression model. “Quadratic” indicates that these terms plus squared terms for easting and northing are also used. Time is always introduced as a linear term, in years, and all models include an intercept.

Table 9 Sediment Time Trend Parameters by Depth and Deposit Group

| Deposit Group | Depth Range (cm) | Log ₁₀ (PCB) Time Trend Slope Estimate | WSEV Standard Error | WSEV <i>p</i> -value | Statistically Significant Slopes | Est. Annual Compound Percent Increase in PCB Level | Estimated Annual Compound Percent Increase in PCB Level | |
|------------------------------------|------------------|---|---------------------|----------------------|----------------------------------|--|---|----------------------------|
| | | | | | | | 95% Conf. Int. Lower-bound | 95% Conf. Int. Upper-bound |
| <i>Little Lake Butte des Morts</i> | | | | | | | | |
| AB | 0–10 | -0.0970 | 0.0348 | 0.0131 | * | -20.0 | -32.5 | -5.2 |
| | 10–30 | -0.0213 | 0.0647 | 0.7535 | | -4.8 | -33.9 | 37.1 |
| | 30–50 | -0.0144 | 0.1113 | 0.8995 | | -3.3 | -45.0 | 70.0 |
| C | 0–10 | -0.0612 | 0.0342 | 0.1481 | | -13.2 | -30.2 | 8.1 |
| | 10–30 | 0.0317 | 0.0770 | 0.7018 | | 7.6 | -34.2 | 76.0 |
| POG | 0–10 | -0.0893 | 0.0567 | 0.1900 | | -18.6 | -43.3 | 16.9 |
| D | 0–10 | -0.0755 | 0.0317 | 0.0307 | * | -16.0 | -28.1 | -1.8 |
| | 10–30 | 0.3168 | 0.0454 | 0.0009 | *** | 107.4 | 58.5 | 171.3 |
| F | 0–10 | -0.0373 | 0.0136 | 0.0252 | * | -8.2 | -14.6 | -1.4 |
| | 10–30 | -0.0760 | 0.0749 | 0.3246 | | -16.1 | -41.7 | 20.8 |
| GH | 0–10 | -0.1244 | 0.0541 | 0.0443 | * | -24.9 | -43.1 | -0.9 |
| <i>Appleton</i> | | | | | | | | |
| IMOR | 0–10 | 0.0412 | 0.0255 | 0.1810 | | 9.9 | -6.6 | 29.4 |
| N Pre-dredge | 0–10 | -0.0281 | 0.0065 | 0.0233 | * | -6.3 | -10.6 | -1.7 |
| | 10–30 | 0.0572 | 0.0440 | 0.2061 | | 14.1 | -7.5 | 40.7 |
| | 30–50 | 0.0846 | 0.0932 | 0.3877 | | 21.5 | -25.2 | 97.4 |
| VCC | 0–10 | -0.0582 | 0.0275 | 0.0878 | | -12.5 | -25.7 | 2.9 |
| | 10–30 | -0.1537 | 0.0164 | 0.0000 | *** | -29.8 | -35.4 | -23.7 |
| | 30–50 | -0.0060 | 0.0151 | 0.6984 | | -1.4 | -8.7 | 6.6 |
| <i>Little Rapids</i> | | | | | | | | |
| Upper EE | 0–10 | -0.0447 | 0.0435 | 0.3618 | | -9.8 | -31.7 | 19.1 |
| | 10–30 | -0.0944 | 0.0429 | 0.0554 | | -19.5 | -35.6 | 0.6 |
| | 30–50 | -0.0712 | 0.0536 | 0.2173 | | -15.1 | -35.8 | 12.2 |
| Lower EE | 0–10 | -0.0682 | 0.0193 | 0.0387 | * | -14.5 | -25.8 | -1.5 |
| | 10–30 | -0.0759 | 0.0390 | 0.0695 | | -16.0 | -30.6 | 1.6 |
| | 30–50 | 0.0900 | 0.0330 | 0.0213 | * | 23.0 | 3.9 | 45.7 |
| FF | 0–10 | -0.0549 | 0.0557 | 0.3400 | | -11.9 | -32.9 | 15.8 |
| | 10–30 | -0.0962 | 0.0390 | 0.0389 | * | -19.9 | -34.9 | -1.4 |
| GGHH | 0–10 | -0.0394 | 0.0231 | 0.1643 | | -8.7 | -21.2 | 5.9 |
| | 10–30 | -0.0182 | 0.0596 | 0.7631 | | -4.1 | -27.7 | 27.3 |
| | 30–50 | 0.1762 | 0.1008 | 0.1188 | | 50.0 | -12.5 | 156.3 |
| | 50–100 | 0.1012 | 0.0700 | 0.1586 | | 26.2 | -9.2 | 75.4 |
| | 100+ | 0.0365 | 0.0249 | 0.1587 | | 8.8 | -3.5 | 22.6 |

Table 9 Sediment Time Trend Parameters by Depth and Deposit Group

| Deposit Group | Depth Range (cm) | Log ₁₀ (PCB) Time Trend Slope Estimate | WSEV Standard Error | WSEV <i>p</i> -value | Statistically Significant Slopes | Est. Annual Compound Percent Increase in PCB Level | Estimated Annual Compound Percent Increase in PCB Level | |
|----------------|------------------|---|---------------------|----------------------|----------------------------------|--|---|----------------------------|
| | | | | | | | 95% Conf. Int. Lower-bound | 95% Conf. Int. Upper-bound |
| <i>De Pere</i> | | | | | | | | |
| SMU Group 2025 | 0-10 | -0.0528 | 0.0231 | 0.0838 | | -11.4 | -23.6 | 2.6 |
| | 10-30 | -0.0556 | 0.0750 | 0.4796 | | -12.0 | -40.9 | 31.0 |
| | 30-50 | -0.0580 | 0.0322 | 0.1016 | | -12.5 | -25.8 | 3.2 |
| | 50-100 | -0.0847 | 0.1058 | 0.4306 | | -17.7 | -50.2 | 35.9 |
| 2649 | 0-10 | -0.0608 | 0.0109 | <0.0001 | *** | -13.1 | -17.4 | -8.5 |
| | 10-30 | -0.2882 | 0.1440 | 0.0764 | | -48.5 | -75.7 | 9.0 |
| | 50-100 | 0.1957 | 0.1419 | 0.2399 | | 56.9 | -36.6 | 288.7 |
| | 100+ | 0.0177 | 0.1548 | 0.9146 | | 4.2 | -61.3 | 180.3 |
| 5067 | 0-10 | -0.0998 | 0.0345 | 0.0136 | * | -20.5 | -33.2 | -5.5 |
| | 10-30 | 0.0912 | 0.0649 | 0.1800 | | 23.4 | -10.3 | 69.6 |
| | 50-100 | 0.3677 | 0.0684 | 0.0030 | ** | 133.2 | 55.5 | 249.5 |
| | 100+ | -0.1963 | 0.2223 | 0.4112 | | -36.4 | -81.8 | 122.6 |
| 6891 | 0-10 | -0.2208 | 0.0944 | 0.1013 | | -39.9 | -69.9 | 20.1 |
| | 10-30 | -0.1685 | 0.0765 | 0.0550 | | -32.2 | -54.4 | 1.0 |
| 92115 | 0-10 | 0.0413 | 0.0426 | 0.3493 | | 10.0 | -10.9 | 35.8 |

Notes:

- * $p < 0.05$
- ** $p < 0.01$
- *** $p < 0.001$

The annual percentage rate of change corresponding to a given slope, b_t , is calculated as

Equation 12

$$\text{Percentage} = 100\% * (10^{b_t} - 1).$$

The halving time is $\frac{\log_{10}(0.5)}{b_t}$ if b_t is negative (decrease over time). If b_t is positive, the doubling time is $\frac{-\log_{10}(0.5)}{b_t}$. The 95 percent confidence interval for the slope, b_t , is given by:

Equation 13

$$[b_t - t_{0.025, df} * SE(b_t), b_t + t_{0.025, df} * SE(b_t)]$$

where

$SE(b_t)$ = the WSEV standard error of b_t , and

$t_{0.025, df}$ = from the t -distribution, 0.025 tail area, with degrees of freedom = df = number of non-empty grid cells, noted in Table A-1.

The 95 percent confidence interval for the percent rate of change is calculated by first deriving the confidence interval for the slope and then using Equation 12 to convert the upper and lower bounds for the slope to upper and lower bounds for the percentage.

The percent increase and the 95 percent confidence interval for the percent increase/decrease (along with the scale for the doubling time or halving time) are presented on Figure 20 through Figure 28. The figures show a number of statistically significant trends. Apparent from Table 9 and the figures is a tendency for more negative slopes to occur at shallower depths and more positive slopes to occur at greater depths. For example, in our Little Lake Butte des Morts Deposit Group D, the slope in the upper 10 cm of sediment is -0.0755 per year, implying a rate of decrease of 16 percent compounded per year; and in the 10- to 30-cm stratum, the slope is 0.317 per year, indicating a rate of increase of 107 percent, compounded annually with trends in both depths being statistically significant ($p = 0.03$ for 0 to 10 cm, $p = 0.0009$ for 10 to 30 cm).

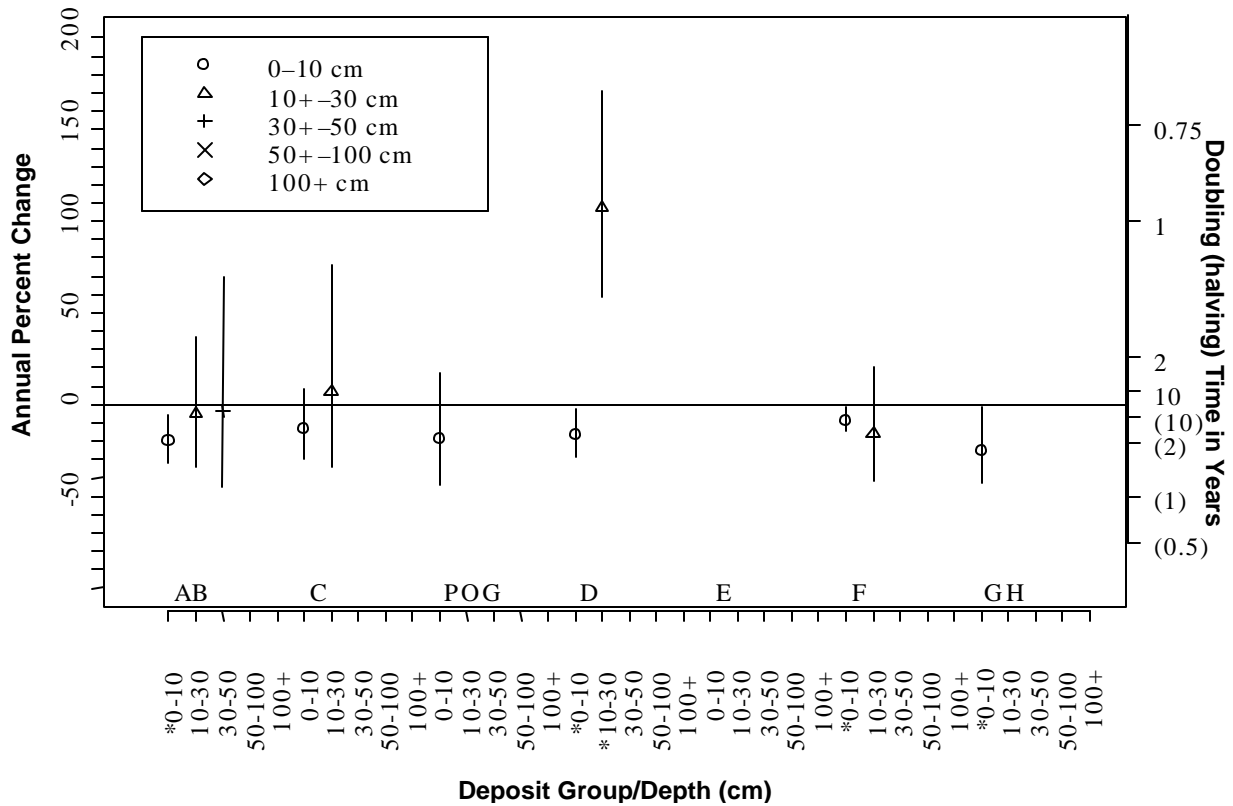


Figure 20 95 Percent Confidence Intervals Showing Annual Percent Rate of Change (Left Vertical Axis) in PCB Concentration

for Little Lake Butte des Morts Deposit Group and Depth Strata

An asterisk (*) below the depth label indicates that a rate of change differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times. Confidence intervals are shown for all deposit groups and depths with sufficient data to perform an analysis of time trend.

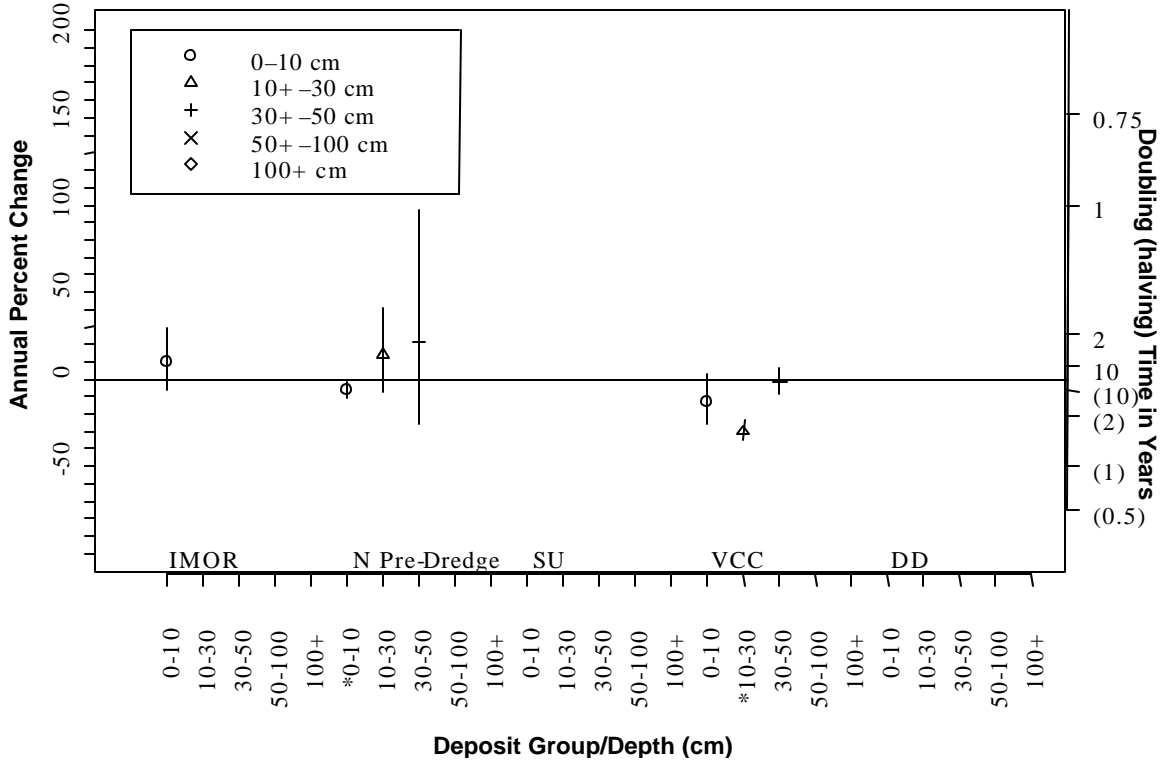


Figure 21 95 Percent Confidence Intervals Showing Annual Percent Rate of Change (Left Vertical Axis) in PCB Concentration for Appleton Deposit Group and Depth Strata

An asterisk (*) below the depth label indicates that a rate of change differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times.

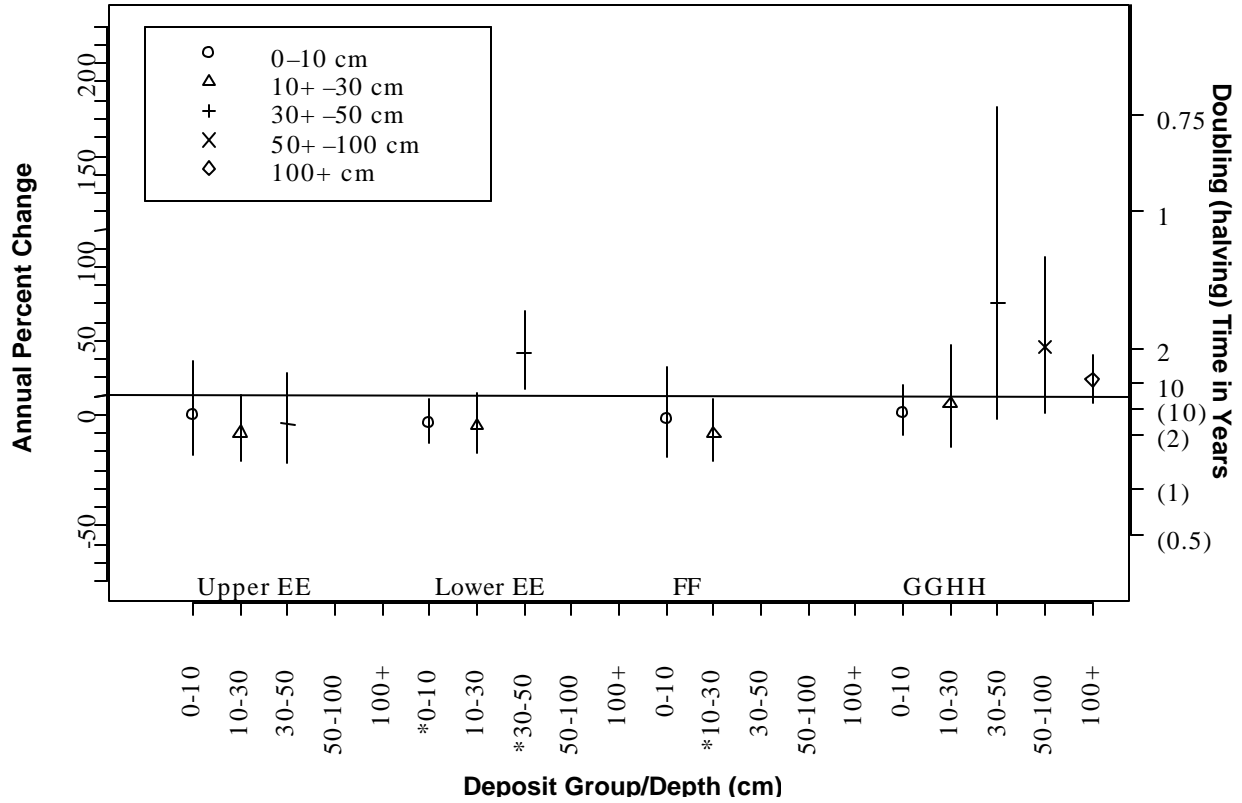


Figure 22 95 Percent Confidence Intervals Showing Annual Percent Rate of Change (Left Vertical Axis) in PCB Concentration for Little Rapids Deposit Groups and Depth Strata

An asterisk (*) below the depth label indicates that a rate of change differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times.

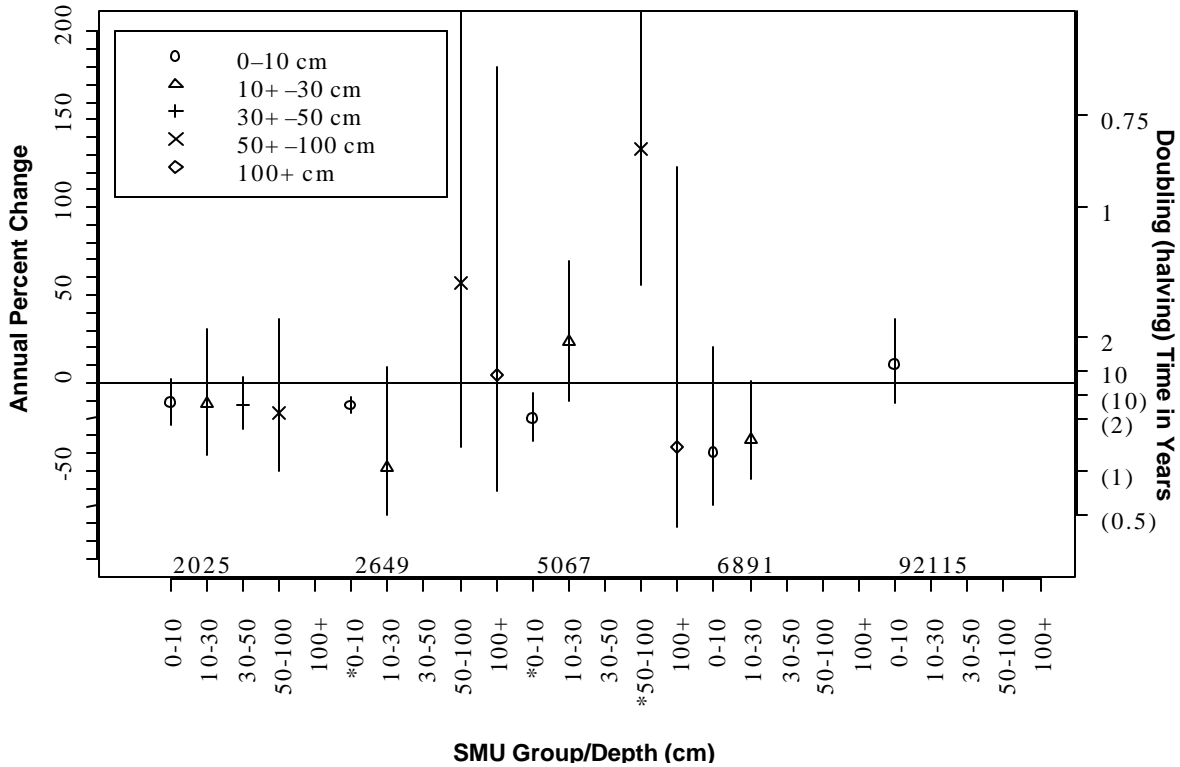


Figure 23 95 Percent Confidence Intervals Showing Annual Percent Rate of Change (Left Vertical Axis) in PCB Concentration for De Pere SMU Groups and Depth Strata

An asterisk (*) below the depth label indicates that a rate of change differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times.

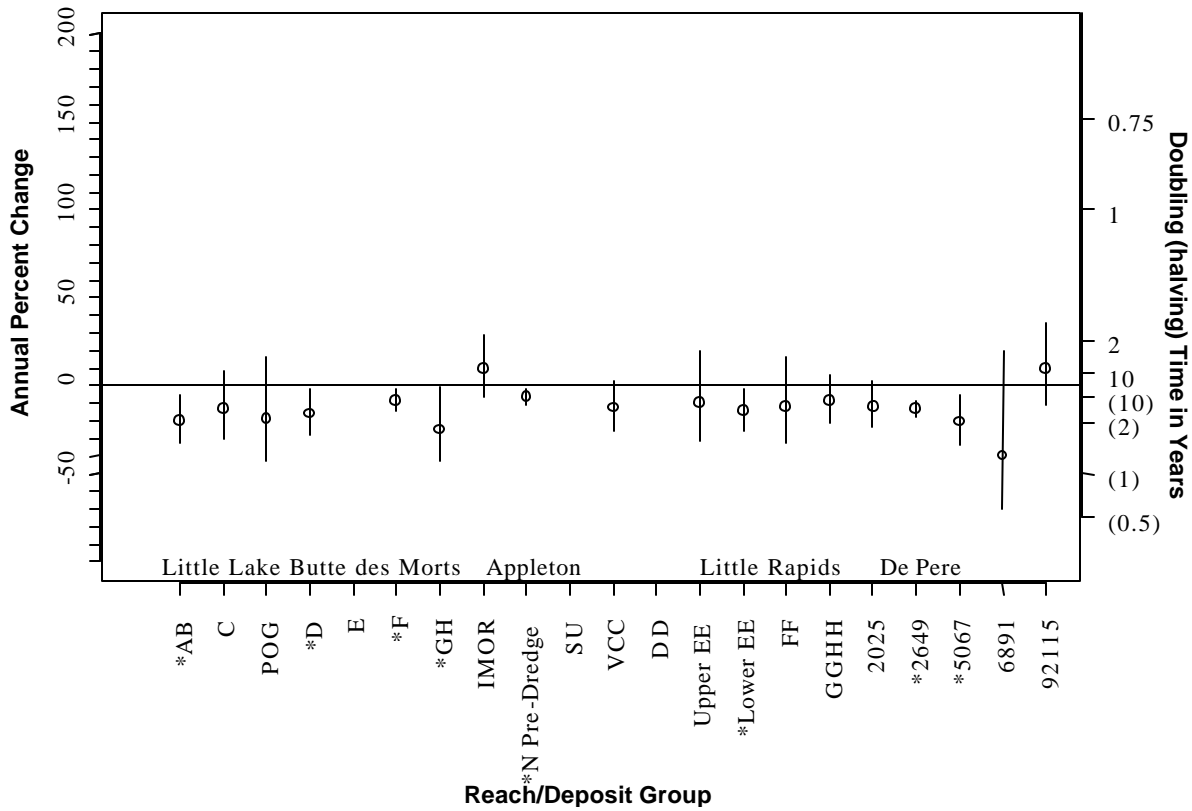


Figure 24 95 Percent Confidence Intervals for Annual Percent Rate of Change at Depth 0 to 10 cm

An asterisk (*) below the depth label indicates that a rate of change differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times.

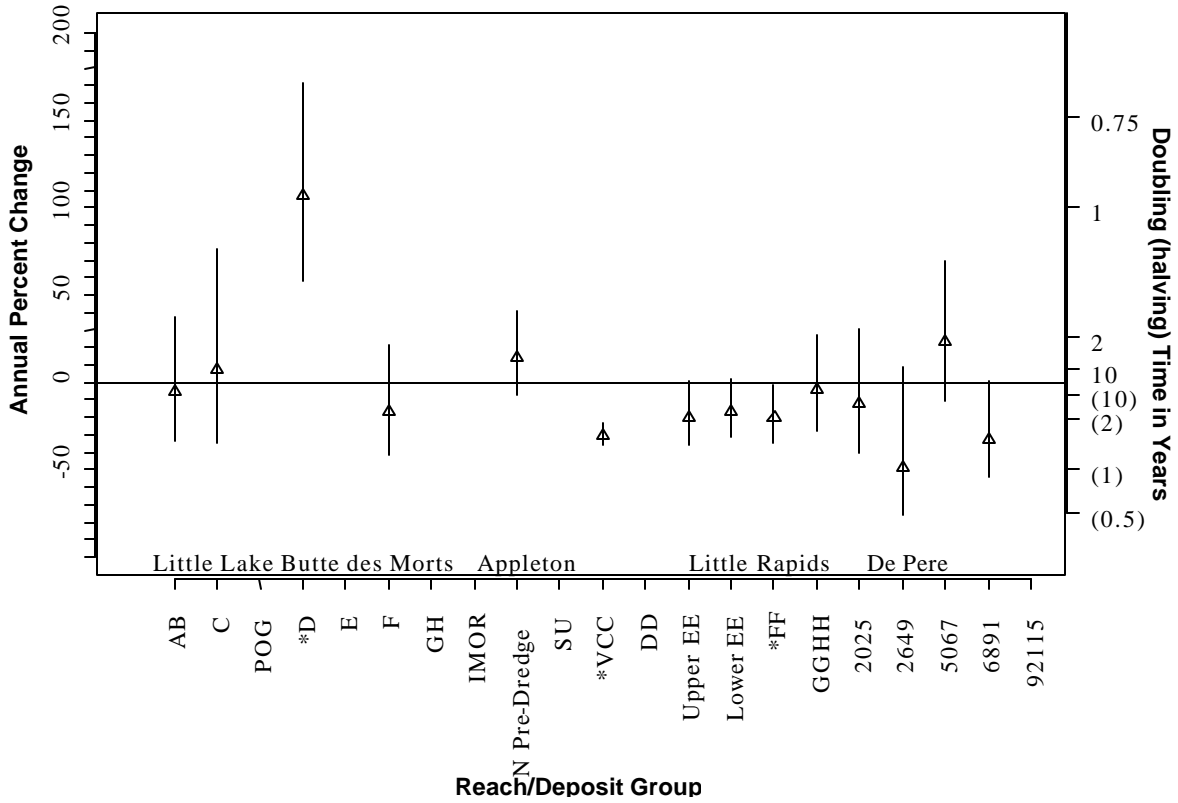


Figure 25 95 Percent Confidence Intervals for Annual Percent Rate of Change at Depth 10+ to 30 cm

An asterisk (*) below the depth label indicates that a rate of change differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times.

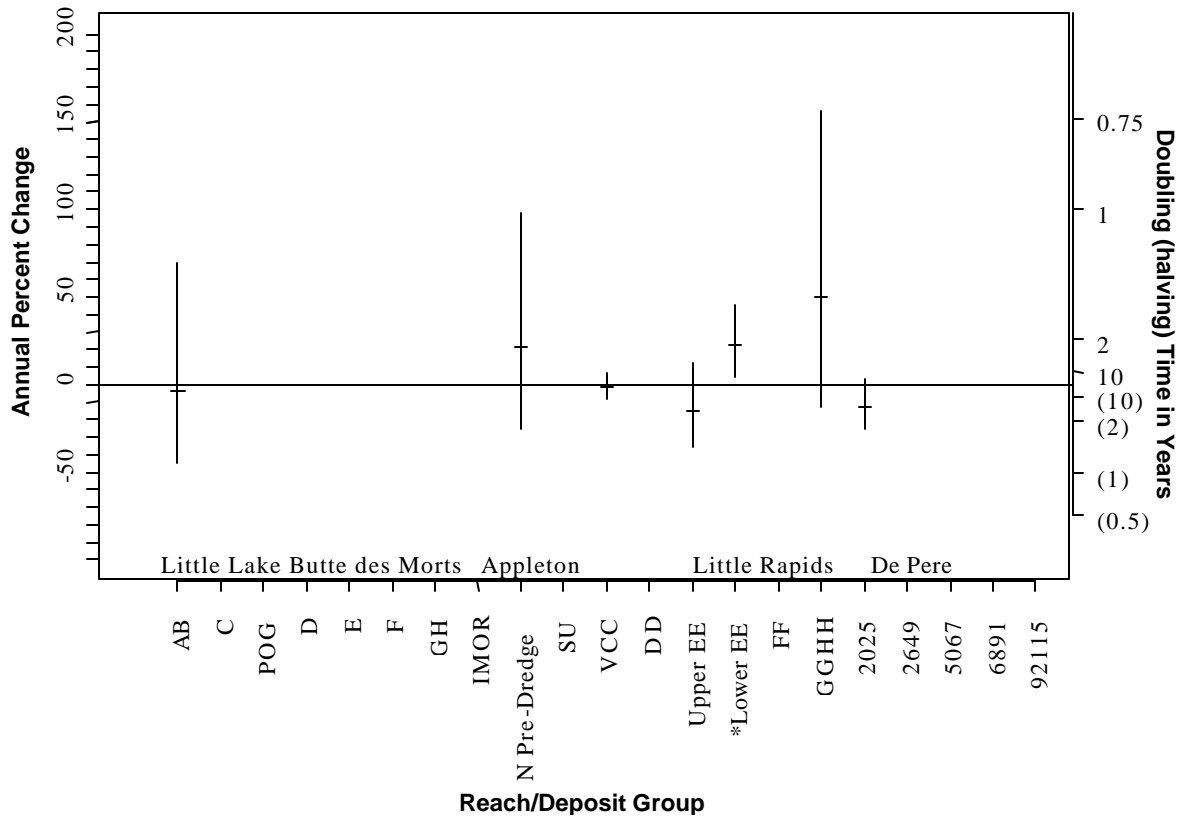


Figure 26 95 Percent Confidence Intervals for Annual Percent Rate of Change at Depth 30+ to 50 cm

An asterisk (*) below the depth label indicates that a rate of change differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times.

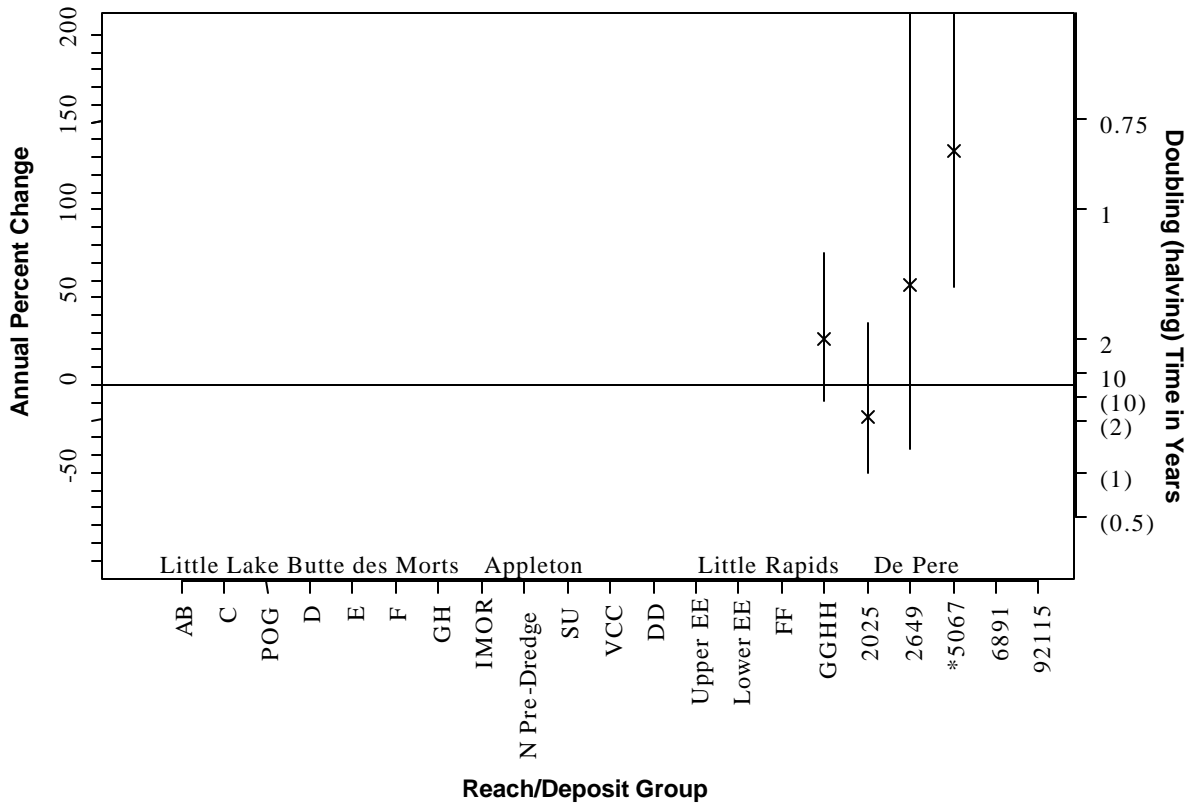


Figure 27 95 Percent Confidence Intervals for Annual Percent Rate of Change at Depth 50+ to 100 cm

An asterisk (*) below the depth label indicates that a rate of change differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times.

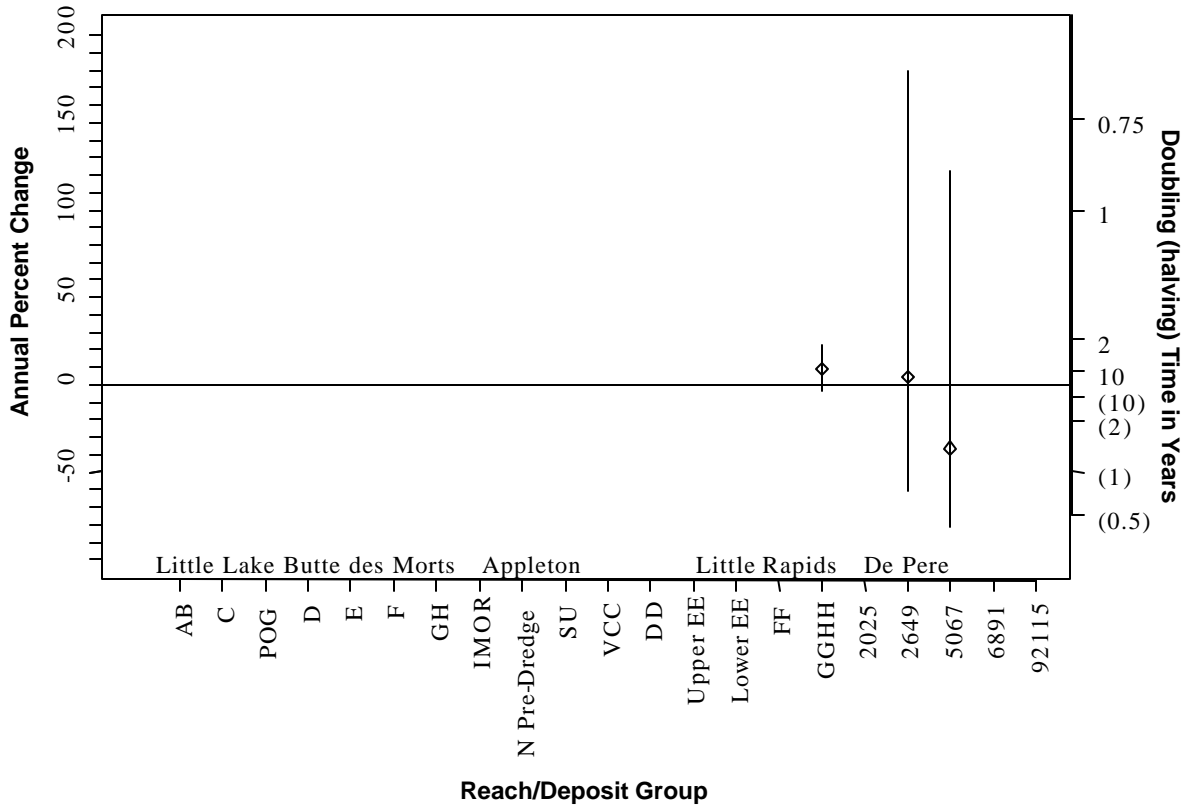


Figure 28 95 Percent Confidence Intervals for Annual Percent Rate of Change at Depth 100+ cm

Right vertical axis expresses time trend change in terms of doubling and halving times.

We note that negative slopes are 89 percent of the calculated slopes from 0 to 10 cm, 71 percent (10/14) of the slopes (16/18) at 10 to 30 cm, 57 percent (4/7) at 30 to 50 cm, 25 percent (1/4) at 50 to 100 cm, and 33 percent (1/3) at 100 cm and over. This indicates a powerful trend toward fewer or weaker negative slopes and more or stronger positive slopes at greater depths. This suggests either that some of the PCBs may transfer out of the river and into Green Bay, instead moving to greater depths, or that attrition of PCBs slows at greater depths, or even that both mechanisms are occurring. These findings can be compared with mass balance studies discussed in the Remedial Investigation for the Lower Fox River and Green Bay.

4.4 Time Trends by Reach

4.4.1 Little Lake Butte des Morts

With the exception of two strata at 10 to 30 cm in two separate deposit groups, slopes are negative (9 out of 11 analyses). Statistically significant negative slopes

(decreasing PCB concentration over time) occur in surface sediments (0 to 10 cm) of four deposit groups (AB, D, F, GH) with estimated rates of decrease ranging from 8 to 24 percent per year (Table 9 and Figure 24). The only statistically significant increasing trend of PCB concentrations occurs at 10 to 30 cm in Deposit Group D, where the rate of increase is 108 percent per year. The confidence intervals for these rates of change are quite wide. For the significantly decreasing slopes in the surface 0- to 10-cm stratum, the confidence intervals indicate a rate of decrease of as little as 1 to 5 percent and as much as 15 to 43 percent per year. The confidence interval for the significantly increasing slope at 10 to 30 cm in Deposit Group D indicates a rate as low as 59 percent and as high as 171 percent per year. This must represent a temporary positive trend because a projection of the PCB concentration even at the minimum of 59 percent per year yields an absurd 10,000-fold increase in PCB concentration after 20 years. Again, the negative slopes also refer to the period of data collection, and one cannot guarantee that such negative slopes would continue indefinitely into the future.

An additional calculation for the surface strata of this reach yields an average slope. This average slope is a weighted mean, where the weights are estimated PCB masses for our deposit groups using mass estimation methods developed in other Fox River studies (WDNR, 1999b). The mass estimates for surface deposits (0 to 10 cm) refer to the boundaries noted on Figure 5 through Figure 8. Because new boundaries have been drawn for these deposit groups, the masses here differ from the masses quoted in other documents for the original deposit designations. Using the estimated PCB mass in the surface sediments (0 to 10 cm) as a relative weight, the weighted mean slope is $-0.071 \pm 0.018 \log_{10}$ PCB concentration per year (*mean* \pm *SE*, Table 11) with $p = 0.0001$ for the null hypothesis of zero slope (i.e., the weighted mean slope is significantly negative and corresponds to an 18 percent rate of decrease of PCB concentration per year). The weighted mean slope is calculated as:

Equation 14

$$b_{wt} = \frac{\left(\sum_{i=1}^K b_i \cdot w_i \right)}{\left(\sum_{i=1}^K w_i \right)}$$

where the b_i are the slopes of the individual deposit groups, $i = 1, \dots, K$, from Table 9 and the w_i are the PCB masses in the strata (see Table 10). The standard error of b_{wt} is calculated as:

Equation 15

$$SE(b_{wt}) = \left[\sum_{i=1}^K (SE \cdot (b_i))^2 (w_i^*)^2 \right]^{0.5}$$

where the $SE(b_i)$ are the standard errors of the individual b values and $w_i^* = \frac{w_i}{\sum_{i=1}^K w_i}$. The statistical significance of the weighted slope is based on a two-sided, single-sample Z-test (twice the tail area of the normal distribution lying beyond $Z = \frac{b_{wt}}{SE(b_{wt})}$).

Table 10 Mass-weighted Combined Time Trend for 0 to 10 cm Depth by Reach

| Deposit Group | Log ₁₀ (PCB) Time Trend Slope Est. | WSEV Standard Error | PCB Mass (kg) | p-value | Annual Percent Change in PCB Conc. | Percent Change 95% Lower-bound | Percent Change 95% Upper-bound |
|------------------------------------|---|---------------------|---------------|----------------------|------------------------------------|--------------------------------|--------------------------------|
| <i>Little Lake Butte des Morts</i> | | | | | | | |
| AB | -0.09705 | 0.034798 | 71.7 | | | | |
| C | -0.06124 | 0.03423 | 25.4 | | | | |
| POG | -0.08935 | 0.056669 | 113.5 | | | | |
| D | -0.07554 | 0.031669 | 32.1 | | | | |
| F | -0.0373 | 0.013582 | 142.5 | | | | |
| GH | -0.12443 | 0.054119 | 15.7 | | | | |
| Reach, Combined | -0.07071 | 0.01831 | 400.9 | 0.0001*** | -15.0 | -21.8 | -7.7 |
| <i>Appleton</i> | | | | | | | |
| IMOR | 0.041186 | 0.025457 | 13.7 | | | | |
| N Pre-dredge | -0.02805 | 0.006544 | 6.9 | | | | |
| VCC | -0.05816 | 0.02746 | 5.2 | | | | |
| Reach, Combined | 0.0025 | 0.01469 | 25.9 | 0.9 | 0.6 | -5.9 | 7.5 |
| <i>Little Rapids</i> | | | | | | | |
| Upper EE | -0.04473 | 0.043487 | 85.0 | | | | |
| Lower EE | -0.06819 | 0.019322 | 25.4 | | | | |
| FF | -0.05486 | 0.055669 | 36.7 | | | | |
| GGHH | -0.03936 | 0.023149 | 131.6 | | | | |
| Reach, Combined | -0.04567 | 0.018764 | 278.7 | 0.01* | -10.0 | -17.3 | -2.0 |
| <i>De Pere</i> | | | | | | | |
| SMU Group 2025 | -0.05279 | 0.02305 | 225.6 | | | | |
| 2649 | -0.06078 | 0.010894 | 356.8 | | | | |
| 5067 | -0.09978 | 0.034549 | 92.4 | | | | |
| 6891 | -0.22081 | 0.094396 | 72.1 | | | | |
| 92115 | 0.041293 | 0.042639 | 37.1 | | | | |
| Reach, Combined | -0.07296 | 0.012829 | 784.0 | <0.0001*** | -15.5 | -20.2 | -10.4 |

Notes:

- * $p < 0.05$
- ** $p < 0.01$
- *** $p < 0.001$

Table 10 provides the weighted slope of surface sediment for each reach. One should interpret the weighted mean slope carefully. This descriptive statistic shows how rapidly the PCB mass is changing at the particular reference date for

the mass estimations (1989–1990), assuming that the rates of change in Table 9 correctly reflect the rates of change at the reference date. The weighted mean slope itself has a straightforward interpretation: it is the rate at which mass is decreasing from the 0- to 10-cm stratum of the collection of deposit groups in the reach. We caution readers when comparing the statistical significance of trends in individual deposit groups (Table 9) to the significance of the weighted mean pooled across all deposit groups in the reach (Table 10). One can calculate a non-significant weighted mean slope although one of the slopes shows, for example, a significant or even highly significant decrease in PCBs over time in the specific deposit group. This can arise where considerable uncertainty exists in some of the slopes being weighted, and when combined, overwhelms the relative certainty of one or two highly significant individual slopes. Thus, one can clearly interpret the value of the slope as representing the rate of decline of the PCB mass at the reference date used for total PCB mass evaluation. One must interpret statistical significance, however, as the likelihood that the observed weighted mean slope could arise, differing from zero, given *within-deposit* sampling variation. It could happen that one sees a significantly negative slope for an individual deposit group with a non-significant overall weighted mean slope. This could occur if, among other deposit groups, the slopes have values close to zero and large enough standard errors such that the mass could conceivably be increasing in these deposit groups. Hence, individual deposit groups with statistically significant slopes alongside a non-significant overall weighted slope should not alarm the reader. In fact, we face just such a contradiction in Appleton, the next reach considered.

The weighted mean slope should not be used for projection of PCB concentrations for the entire reach, because deposit groups with the lowest rate of decrease will in the future dominate the decay of PCB mass over time. The weighted mean slope serves as a summary descriptive value representing average change during the period of data collection and, also, as a statistic used to derive a significance level (*p*-value) for the hypothesis of no change. The PCB mass remaining in the future, *w*, can be estimated as:

Equation 16

$$w = \sum_{i=1}^K w_i * 10^{b_i t}$$

where time, *t*, is measured in decimal years since January 1, 1989, *w_i* are the PCB masses in the *K* strata, and *b_i* is the coefficient of time term in the model for the *i*th stratum. The equation works for any collection of *K* strata.

4.4.2 Appleton Reach

Two strata have statistically significant slopes. The, 0 to 10 cm in the Deposit Group N (pre-dredge) has a statistically significant negative slope of *b* = --0.028 (log₁₀ PCB concentration per year). This slope translates into a rate of decrease of 6 percent per year with a 95 percent confidence interval of 2 percent to 11 percent decrease per year (Table 9). The 10- to 30-cm stratum of Deposit Group VCC has

a statistically significant decrease of -0.154 (\log_{10} PCB concentration per year), implying a 30 percent rate of decrease per year with a 95 percent confidence interval of -35 to -24 percent (Table 9).

The weighted slope for surface strata is 0.003 per year, implying a rate of increase of 0.6 percent per year with a 95 percent confidence interval of -6 to $+7$ percent per year. This mass-weighted mean slope of -0.011 per year is not statistically significant ($p = 0.4$). Even though the N Pre-dredge Deposit Group has a significantly decreasing slope in the 0- to 10-cm stratum (equivalent to a 2.6% decrease per year), the total PCB mass in surface sediments in the entire Appleton Reach may be either increasing, decreasing, or remaining constant over time. The reach includes the one statistically significant negative slope for surface sediments, as well as an additional positive and negative slope. Thus, while it is likely that one surface deposit is, indeed, decreasing in PCB concentration, the combination of positive and negative slopes convey a state of uncertainty as to the trends in total PCB mass in the combined surface deposits in the reach.

4.4.3 Little Rapids to De Pere Reach

This reach has a majority of negative slopes (change in \log_{10} [PCB concentration] per year). Two of the three significant slopes are negative and occur in the 0- to 10-cm and 10- to 30-cm depth strata. One large positive statistically significant slope occurs at the 30- to 50-cm depth (Table 9).

The surface sediment (0 to 10 cm) in the Lower EE Deposit Group has a significantly negative slope (-0.068 per year), implying a rate of decrease of 15 percent per year with a 95 percent confidence interval of 2 to 26 percent rate of decrease per year. In the same deposit group, the deeper 30- to 50-cm stratum shows a significantly positive slope, indicating a rate of increase of 23 percent per year and a 95 percent confidence interval of 4 to 46 percent per year. In Deposit Group FF, the 10 to 30 cm layer has a significantly negative slope with a rate of PCB concentration decrease of 20 percent per year with a 95 percent confidence interval of 1 to 35 percent. Again, while the estimates speak to significant decreasing or increasing PCB concentrations over time in these strata and deposit group combinations, we still encounter notably wide confidence intervals.

Although only one surface sediment has a statistically significant decline, we nonetheless find an overall statistically significant combination of declining PCB concentrations in the reach, with a slope of -0.046 per year ($p = 0.01$), implying a 10 percent per year rate of decrease (95 percent confidence interval: -17 to -2 percent). While some uncertainty may persist in the individual surface deposits, the PCB mass in the surface of this reach appears to be generally declining as of the mass estimation date, 1989 through 1990.

4.4.4 De Pere to Green Bay Reach

This reach, again, has primarily negative slopes (Table 9). Statistically significant negative slopes occur in three combinations of deposit group and depth. Our SMU Group 2649 has a significantly negative slope in the surface deposit (0 to 10 cm), with a rate of decrease of 13 percent per year (95 percent confidence interval of 8 to 17 percent per year) and $p < 0.0001$. SMU Group 5067, 0 to 10 cm, also has a significantly negative slope implying an annual rate of decrease of 21 percent (95 percent confidence interval of 5 to 33 percent) and $p = 0.01$. In the same SMU group (5067), at a greater depth of 50 to 100 cm, we observe a statistically significant and large positive slope with a rate of increase of 133 percent per year (95 percent confidence interval of 56 to 250 percent) and $p = 0.003$.

We noted earlier (Section 2.6 and Table 5) an exceptional value of PCB concentration in SMU Group 5067. Sample A3_0-4 had a concentration of 99,000 ppb, whereas all other samples in the 0- to 10-cm stratum in this deposit ranged from 400 to 7,800 ppb. In a statistical sense, the sample is an “outlier,” but that does not imply error in the value of 99,000. We have no reason to suspect invalidity of the concentration of 99,000 ppb for sample A3_0-4, especially given internal evidence in the deposit corroborating it (see below). However, the sample is a statistical outlier to the spatial relationships of PCB concentrations in the deposit, as we shall show. The spatial layout of the samples in the 0- to 10-cm stratum of SMU Group 5067 is shown on Figure 29. The samples occur in an intensively sampled area (see Figure 8). Sample A3_0-4 lies close to the shore of the river, and we have been informed that this sample was located in the vicinity of direct deposition of PCBs. The more immediate vicinity of the sample is shown on Figure 30, which includes 34 out of the 58 samples in the 0- to 10-cm layer of the deposit. Figure 30 also designates the exceptional sample A3_0-4 (#1 in the plot) and the six samples closest to it.

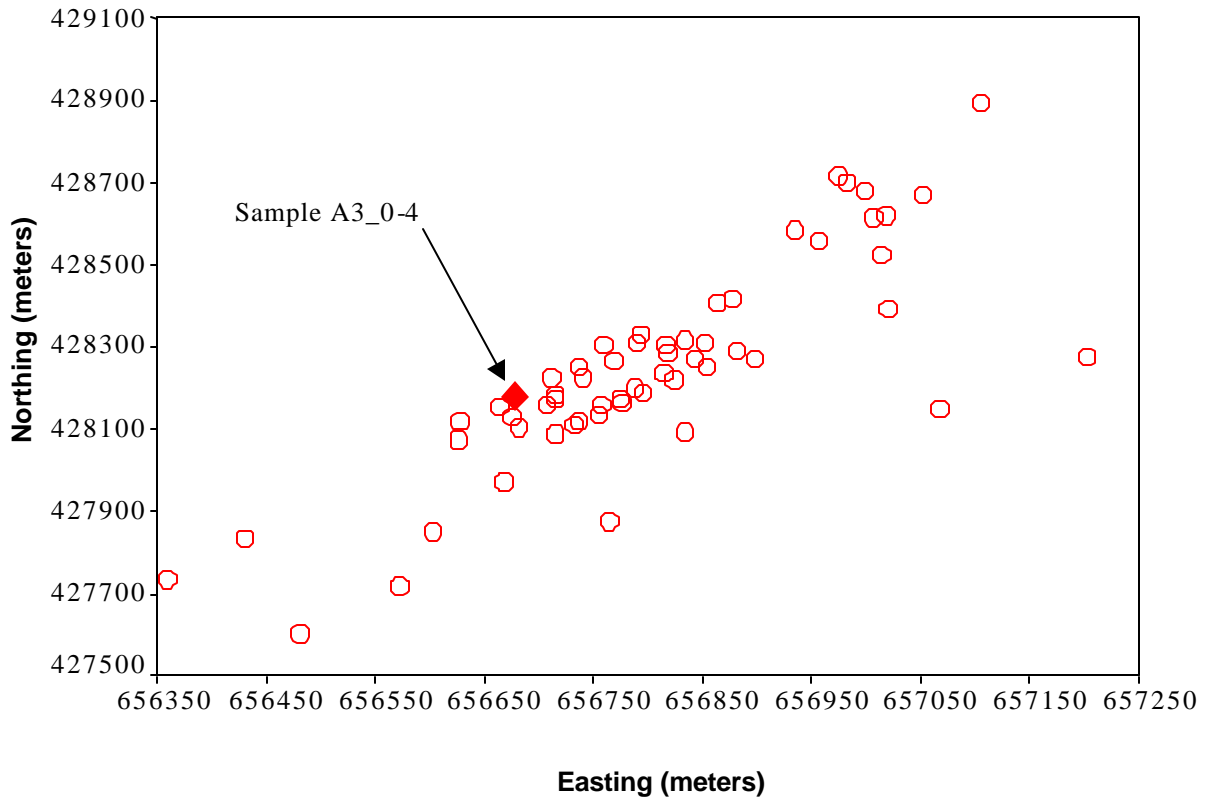


Figure 29 De Pere SMU Group 5067: Location of 0 to 10 cm Core-averaged Samples with Sample A3_0-4 Identified

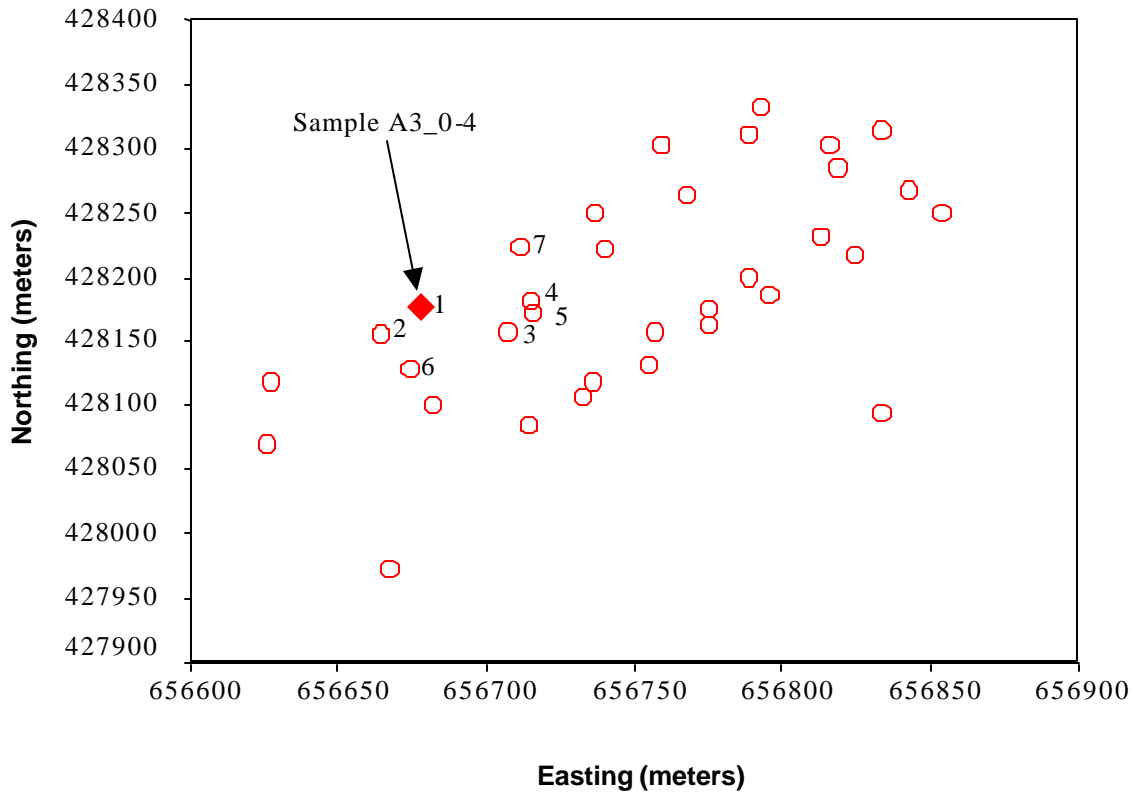


Figure 30 Sample Locations for SMU Group 5067, 0 to 10 cm Depth, Samples Closest to Sample A3_0-4 (Less than 208 meters Distance)

A3_0-4 and the six samples closest to it are labeled.

The specific concentrations of the samples near sample A3_0-4 are shown in Table 11, which includes not only concentrations for the 0- to 10-cm layer, but concentrations in lower sediment layers in precisely the same locations. (The samples have the same northing and easting coordinates down through the layers, presumably because multilayer samples were collected in a single coring operation.) These seven samples all occur within a radius of less than 60 meters from the location of A3_0-4. We note that in the 0- to 10-cm layer, all of these nearby samples are in the 2,000 to 3,000 ppb range, less than one-twentieth of the concentration of sample A3_0-4. In the next layer down, 10 to 30 cm, the highest concentration by a wide margin occurs at the same location as sample A3_0-4, suggesting that this sample location does, indeed, have a high concentration of PCBs and that the location differs from immediately neighboring sediment. We excluded the layer 30 to 50 cm from our time trends analysis (due to lack of time variation of samples) and, therefore, it does not appear in the table. The 50- to 100-cm layer shows a high concentration at the location of sample A3_0-4, but the other samples near it also show high

concentrations. At 100+ cm, the PCB concentration no longer stands out at the location of sample A3_0-4.

Table 11 PCB Concentrations at Various Depths and Distances from Sample A3_0-4

| Sample | 0–10 cm | 10–30 cm | 50–100 cm | 100+ cm | Year | Easting (meters) | Northing (meters) | Distance (meters) |
|--------|---------|----------|-----------|---------|------|------------------|-------------------|-------------------|
| 1 | 99,000 | 150,000 | 150,000 | 53,122 | 1997 | 656678 | 428177 | 0 |
| 2 | 2,000 | 2,200 | 34,000 | 18,128 | 1997 | 656664 | 428155 | 26 |
| 3 | 2,100 | 3,100 | 7,000 | 61,094 | 1997 | 656707 | 428158 | 35 |
| 4 | 1,900 | 4,300 | 170,000 | 66,106 | 1997 | 656715 | 428182 | 37 |
| 5 | 2,700 | 4,800 | 13,000 | 30,948 | 1995 | 656716 | 428172 | 38 |
| 6 | 2,000 | 4,500 | 7,800 | 41,729 | 1997 | 656675 | 428128 | 49 |
| 7 | 3,000 | 9,900 | 120,000 | 6,665 | 1997 | 656711 | 428224 | 58 |

The value of 99,000 ppb stands out as considerably larger than nearby samples, which have quite uniform concentrations of PCBs and thereby heighten the contrast. We do not imply that the value of 99,000 is artificial, but it cannot readily be included in a regression analysis for the deposit. A valid regression analysis depends upon the included concentrations approximately following a normal (bell-shaped) distribution around the fitted regression model. A model fitted to the log concentrations in the 0- to 10-cm layer with sample A3_0-4 included shows that the sample is 5.5 standard deviations away from the model-fitted value, whereas all other samples are at most 2.6 standard deviations from their model-fitted values. With a sample of this size ($n = 58$, including A3_0-4), the occurrence of observations lying three or more standard deviations from the model questions the accuracy of the model. The deviation of 5.5 is exceptionally large. Even ignoring the modeling process, the log concentration of A3_0-4 is 7.4 standard deviations above the mean of the balance of observations, and the next largest observation is only 2.5 standard deviations above the same mean.

Thus, sample A3_0-4 appears to represent a real but exceptional concentration in the 0- to 10-cm layer. The regression model excluding it thus covers all of the 0- to 10-cm layer in the deposit except the immediate vicinity of this sample. The statistically significant decline in PCBs noted for this layer in Table 9 does not, then, necessarily apply to this small area. It is impossible to develop an estimate of the time trend for this “hotspot” alone. Of the nearby samples (Table 11), all except one occur at the same time as sample A3_0-4—1997. The lack of time variation of samples in the vicinity of A3_0-4 precludes a separate regression analysis for this sub-area.

The large concentration at the same location as A3_0-4, but one layer down—the 150,000 ppb concentration at 10 to 30 cm, is not an outlier to its layer. Its nearby samples vary considerably more among themselves relative to the variability observed in the corresponding samples from the 0- to 10-cm layer. Thus, the 150,000 value does not stand out with nearly as much contrast relative to the 99,000 value among its neighbors. The residuals from the regression

analysis of the PCB concentrations in the 10- to 30-cm layer also show no statistical outliers. One reason that the concentration at this location in the 10- to 30-cm layer is so large may be that a hotspot extends from the 0- to 10-cm layer into at least part of the 10- to 30-cm layer.

In summary, the 0- to 10-cm layer of the deposit, outside of the vicinity of A3_0-4, shows a statistically significant decline in PCB concentration over time. The vicinity of A3_0-4 encompassed an area of exceptionally high concentrations with an unknown time trend. The exceptional vicinity of A3_0-4 is a small fraction of the total deposit area. A circle centered on A3_0-4 and bounded by the nearest sample (which has a typical concentration), 26 meters away, would have an area of 2,100 square meters, or approximately 0.3 percent of the 840,000-square meter total area covered by all samples of SMU Group 5067 in the 0- to 10-cm layer.

The mean slope for surface sediments in this reach, weighted by PCB mass, is -0.073 ± 0.013 and highly significant ($p < 0.0001$, Table 10). The negative slope implies a rate of decrease of 15 percent per year (95 percent confidence interval: -20 to -10 percent per year).

4.5 Comments on Combined Reaches

There may be some concern about the many analyses carried out and the possibility that some of the trends, both positive and negative, are statistically significantly different from zero by chance alone. We carried out a formal test for the hypothesis that the slopes, positive and negative, are simply randomly distributed around zero (i.e., the statistically significant differences from zero result from the large number of analyses carried out). Under the null hypothesis that the true slopes are all 0, the p -values should be uniformly distributed between 0 and 1.0, and minus twice the sum of the natural log of the p -values will yield a chi-squared variable with degrees of freedom equal to twice the total number of analyses (p -values) included. Carrying out this operation and obtaining a p -value for this null hypothesis of all zero true slopes yields $p < 0.0001$ for depth 0 to 10 cm, $p < 0.0001$ for depth 10 to 30 cm, $p = 0.07$ for 30 to 50 cm, $p = 0.01$ for 50 to 100 cm, and $p = 0.46$ for 100+ cm. Thus, it appears clear that there exist non-random changes in slope, both positive and negative, for all depths, except, possibly, 30 to 50 cm and 100+ cm. We conclude that real changes in concentrations are taking place over time in the Lower Fox River.

5 Fish Results

5.1 Number of Observations

A total of 1,742 fish samples were available for analysis, including sample types of fillet without skin, fillet with skin, and whole body. We excluded samples of eggs, stomach, carcass, and other miscellaneous sample types, as well as those for which percent lipid was unknown. As a criterion for analysis, we included only unique combinations of species and sample type for a given reach with at least 14 observations. In general, our largest model included seven parameters to be estimated. Thus, the minimum of 14 observations ensures at least twice as many observations as parameters. As some statistical “rules of thumb” require at least four or five times as many observations as parameters, our rule might strike many as rather generous. Nevertheless, we decided to err on the side of inclusiveness and to interpret with some caution analyses with a small number of observations. As an important additional condition, we required sufficient variation in time to provide a meaningful estimate of a time trend. The data provided 108 combinations of reach, species, and sample type with at least one observation, but only 19 of these had sufficient numbers of samples and an adequate time spread for analysis (see Table 12). In Little Lake Butte des Morts, 6 out of 23 combinations could be analyzed. For the other reaches, corresponding numbers are 1 of 20 for Appleton Reach, 0 of 16 for Little Rapids Reach, 7 of 24 for De Pere Reach, and 5 of 25 for Green Bay Zone 2. The 19 combinations that could be analyzed for time trends represent 868 samples—over half of all samples of whole body, fillet with skin, and fillet without skin. Carp and walleye provided the largest number of observations. None of the observations of fillet without skin would be analyzed due to either inadequate sample size or inadequate time variation. One outlier was detected and removed (see Appendix Table A-2).

Table 12 Sample Sizes for Total PCB Time Trend Analyses by Reach, Species, and Sample Type

| | Fillet/ No Skin | Fillet/ Skin-on Fillet | Whole Fish, Whole Body, Whole Body Composite | Eggs, Stomach, Carcass, Other | Total Sample Size: Fillet, Fillet No Skin, Skin-on Fillet, Whole Fish, Whole Body, Whole Body Composite |
|------------------------------------|--------------------|------------------------------|---|--|---|
| <i>Little Lake Butte des Morts</i> | | | | | |
| Brown Bullhead | 4 | 8 | 6 | | 18 |
| Carp | 20 | 55* | 40* | | 115 |
| Gizzard Shad | | | 4 | | 4 |
| Northern Pike | | 19* | 5 | | 24 |
| Smallmouth Bass | | 7 | 2 | | 9 |
| Walleye | 7 | 63* | 18* | | 88 |
| White Bass | | 26 | | 2 | 26 |
| White Sucker | 10 | 19 | 8 | | 37 |
| Yellow Perch | | 34* | 7 | 1 | 41 |
| Other | 2 | 10 | 5 | 1 | 17 |
| <i>Appleton to Little Rapids</i> | | | | | |
| Brown Bullhead | 1 | 2 | | | 3 |
| Carp | | 24 | 13 | | 37 |
| Channel Catfish | 6 | | | | 6 |
| Northern Pike | | 7 | 4 | | 11 |
| Smallmouth Bass | | 5 | 4 | | 9 |
| Walleye | | 30* | 4 | | 34 |
| White Bass | | 8 | 2 | | 10 |
| White Sucker | | 17 | 6 | | 23 |
| Yellow Perch | | 2 | 7 | | 9 |
| Other | 1 | 10 | 3 | | 14 |
| <i>Little Rapids to De Pere</i> | | | | | |
| Carp | | 2 | 22 | | 24 |
| Channel Catfish | 3 | | | | 3 |
| Gizzard Shad | | | 3 | | 3 |
| Northern Pike | | 3 | 1 | | 4 |
| Smallmouth Bass | | 16 | 2 | | 18 |
| Walleye | | 48 | 4 | | 52 |
| White Bass | | 14 | | | 14 |
| Yellow Perch | | 3 | 2 | | 5 |
| Other | 4 | 6 | 8 | | 18 |

Table 12 Sample Sizes for Total PCB Time Trend Analyses by Reach, Species, and Sample Type

| | Fillet/Fillet No Skin | Skin-on Fillet | Whole Fish, Whole Body Composite | Eggs, Stomach, Carcass, Other | Total Sample Size: Fillet, Fillet No Skin, Skin-on Fillet, Whole Fish, Whole Body, Whole Body Composite |
|-------------------------------------|-----------------------|----------------|----------------------------------|-------------------------------|---|
| <i>De Pere to Green Bay</i> | | | | | |
| Alewife | | | 15 | | 15 |
| Brown Bullhead | | | 2 | | 2 |
| Carp | | 12 | 90* | 13 | 102 |
| Channel Catfish | 17 | | | | 17 |
| Gizzard Shad | | 2 | 19* | | 21 |
| Northern Pike | | 40* | 6 | | 46 |
| Smallmouth Bass | | 15 | 4 | | 19 |
| Walleye | 14 | 120* | 58* | 8 | 192 |
| White Bass | 3 | 58* | 9 | 8 | 70 |
| White Sucker | | 44* | 22 | 2 | 66 |
| Yellow Perch | | 11 | 9 | | 20 |
| Other | 6 | 36 | 42 | 1 | 84 |
| <i>Green Bay Zone 2 (2A and 2B)</i> | | | | | |
| Alewife | | 3 | 44* | | 47 |
| Brown Bullhead | 6 | 2 | 1 | | 9 |
| Carp | | 28* | 57* | 28 | 85 |
| Channel Catfish | 5 | | | | 5 |
| Gizzard Shad | | 1 | 32* | | 33 |
| Northern Pike | | 7 | 1 | | 8 |
| Rainbow Smelt | | 2 | 33 | | 35 |
| Smallmouth Bass | | | 2 | | 2 |
| Walleye | | 17 | 34 | | 51 |
| White Bass | | 3 | | | 3 |
| White Sucker | | 7 | 1 | | 8 |
| Yellow Perch | | 19* | 5 | | 24 |
| Other | 3 | 33 | 2 | | 38 |
| Total (all reaches): | | | | | 1,678 |

Note:

* Included in time trends analysis. Total $n = 868$.

While inadequate sample size for some species from some reaches presented the greatest obstacle to analysis, several cases with substantial numbers of observations suffered from inadequate spread over time, such as whole body white sucker in the De Pere to Green Bay Reach, with 22 observations. Notably, Little Rapids to De Pere Reach had no groups with both sufficient sample size and time spread.

Overall, only a small fraction of the observations had values below detection limit (BDL). Among the 19 combinations with a total of 868 samples, only $n = 28$ (3%)

were BDL. Several combinations had no BDL concentrations (0%), and BDL observations occurred mainly in four combinations, which had 13 to 29 percent BDL values. All observations, both above and below detection limits, in the selected combinations of reach, species, and sample type were used in the time trends analysis. Appendix Table A-3 indicates the number of observations below detection limits.

5.2 Time Trends in PCB Concentrations in Fish

We organize results in three major sections:

First, we introduce some ancillary results relevant to the process of model fitting, such as identifying the optimal location of the breakpoint and coefficients on percent lipids and seasonality (Section 5.2.1).

Then we turn to the main results, concerning rates of decline of PCB concentrations. The time trends for each species and sample type, within each reach, can be found in this section (Section 5.2.2).

Finally, we consider alternative models, such as those with a common breakpoint at 1985 for all fish categories and curvilinear (quadratic) models to test whether trends are constant or changing over time (Section 5.2.3).

5.2.1 Testing Spline Model versus Simple Linear Model

Table 13 shows results of testing the null hypothesis of a linear relationship between log of PCB concentration and time over the entire time period of the data versus the alternative hypothesis of a spline: two linear segments joined at a breakpoint. The year of the best-fitting spline model is shown in Table 13, and the p -value indicates whether the spline model significantly improves the fit to the data. With one exception (yellow perch, skin-on fillet in Green Bay Zone 2), the spline model has been used if $p < 0.05$ in Table 13; this means that a spline model fits significantly better than a simple, single-slope linear model.

Table 14 through Table 16 provide a description, reach by reach, of the final slopes from the fitted models (or the only slope, if there is no breakpoint) and Table 18 provides other model parameters discussed in this section. One can find the complete model in Appendix Table A-3 or A-6.

Table 13 Testing the Null Hypothesis that a Straight Line Fits As Well As a Spline Model with a Breakpoint

| Reach and Species | Sample Type | Year of Best-fitting Breakpoint | Sample Size (n) | Breakpoint | |
|-------------------------------------|-------------------------------|---------------------------------|-----------------|-------------------|---------------------------|
| | | | | p-value | Statistically Significant |
| <i>Little Lake Butte des Morts</i> | | | | | |
| Carp | skin-on fillet | 1979 | 55 | 0.0347 | * |
| Carp | whole fish ⁺ | 1987 | 40 | 0.0263 | * |
| Northern Pike | skin-on fillet | 1996 | 19 | 0.2723 | |
| Walleye | skin-on fillet | 1990 | 63 | 0.0423 | * |
| Walleye | whole fish ⁺ | 1987 | 18 | 0.0088 | ** |
| Yellow Perch | skin-on fillet | 1981 | 34 | 0.0062 | ** |
| Combined⁺⁺ | | | 229 | <0.0001 | *** |
| <i>Appleton</i> | | | | | |
| Walleye | skin-on fillet | 1983 | 30 | 0.4526 | |
| <i>De Pere</i> | | | | | |
| Carp | whole fish ⁺ | 1995 | 90 | 0.0087 | ** |
| Gizzard Shad | whole fish ⁺ | 1990 | 19 | 0.4672 | |
| Northern Pike | skin-on fillet | 1996 | 40 | 0.1421 | |
| Walleye | skin-on fillet | 1993 | 120 | 0.5680 | |
| Walleye | whole fish ⁺ | 1996 | 58 | 0.5550 | |
| White Bass | skin-on fillet | 1996 | 58 | 0.6059 | |
| White Sucker | skin-on fillet | 1990 | 44 | 0.1986 | |
| Combined⁺⁺ | | | 429 | 0.0906 | |
| <i>Green Bay Zone 2 (2A and 2B)</i> | | | | | |
| Alewife | whole fish ⁺ | 1986 | 44 | 0.0863 | |
| Carp | skin-on fillet | 1985 | 28 | 0.1811 | |
| Carp | whole fish ⁺ | 1983 | 57 | 0.0001 | *** |
| Gizzard Shad | whole fish ⁺ | 1996 | 32 | 0.6655 | |
| Yellow Perch | skin-on fillet ⁺⁺⁺ | 1986 | 19 | 0.0008 | *** |
| Combined⁺⁺ | | | 180 | <0.0001 | *** |

Notes:* $p < 0.05$ ** $p < 0.01$ *** $p < 0.001$

+ Whole fish, or whole body, or whole body composite.

++ Indicates p -value for testing the null hypothesis that all fish categories in a reach do not have a breakpoint.

+++ A model with a breakpoint was rejected. See text.

Reach 1 — Little Lake Butte des Morts

In the first reach, for five of the six fish categories, the spline model fit significantly better than the linear model. In all cases, the initial slope decreased more steeply than the final slope, as seen by the negative coefficient for the slope difference. Figure 31 for carp fillet with skin in Little Lake Butte des Morts shows an example of an initial steep slope until 1979, followed by a continuing decline, but at a slower rate. Similar plots for all analyses are found in the Appendix. Figure 32 shows an example of initial decline until 1990, followed by a virtually flat line implying no further decline in PCBs. Figure 33 shows an example in which PCB concentration actually increases after the breakpoint in 1987. With

only 18 data points and 9 distinct time points, one should interpret this result cautiously. Of note, the fitted line appears to fit poorly prior to 1987 because all of the observations lie above the fitted line. The fitted line, however, represents the prediction for fish with percent lipid equal to the mean, sampled on July 1. For this fish category, samples were taken prior to 1987 in late August and early September, and after 1987 mainly in July and some as early as April. This discrepancy, plus evidence for a significant seasonal effect for this fish category, explains the poor visual fit on the plot. The row at the bottom of the panel for Little Lake Butte des Morts in Table 14 reports the p -value from a meta-analysis for this reach. This meta-analysis combines the results from all species within this reach to test the global null hypothesis that a linear model fits well for **all** species/types versus the alternative that a spline model with a breakpoint gives a better fit for at least one species. The highly significant p -value provides strong evidence to reject the null hypothesis that every species has a constant rate of decline over the entire time frame in Little Lake Butte des Morts.

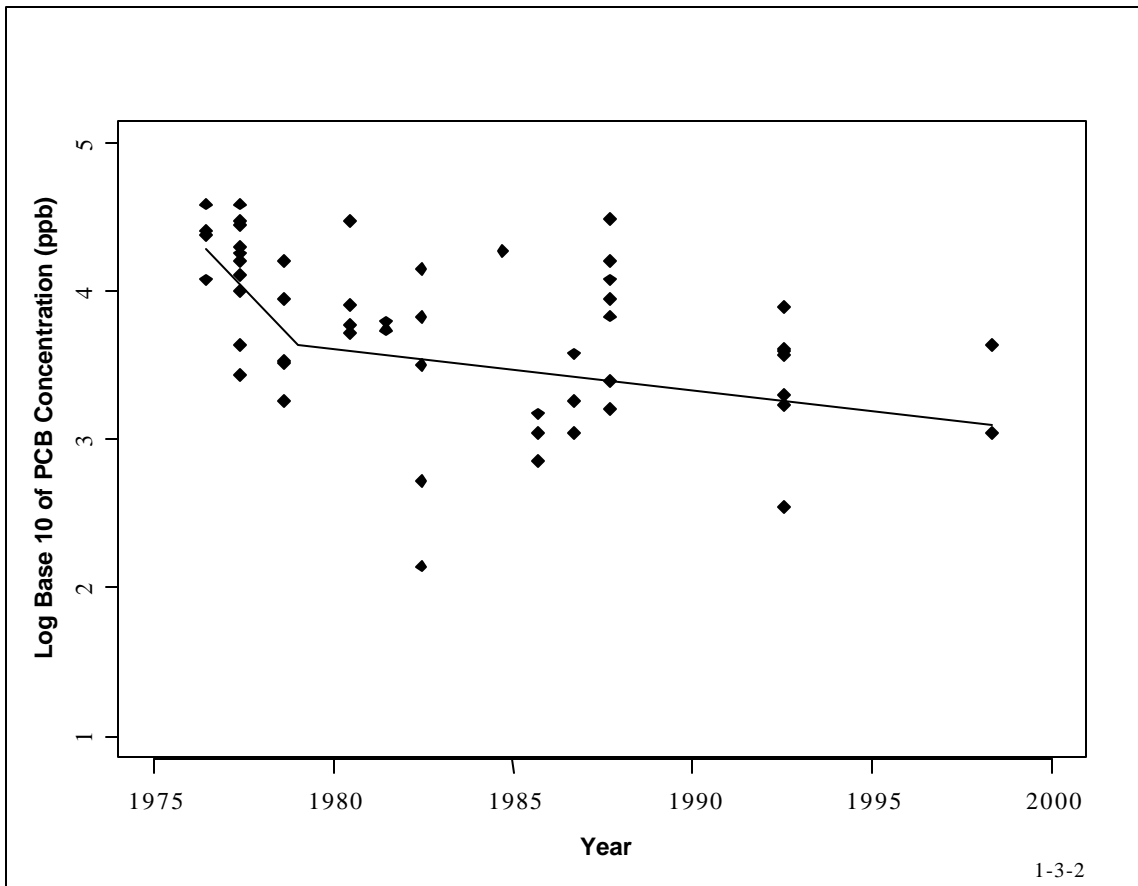


Figure 31 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Carp, Skin-on Fillet, versus Time

Breakpoint = 1979 ($p = 0.03$), Final Slope (\log_{10} PCB versus time) = -0.028 ($p = 0.02$), Rate of Change of PCB Concentration During Period of Final Slope = -6.1% (95% confidence interval: -10.9% to -1.1%).

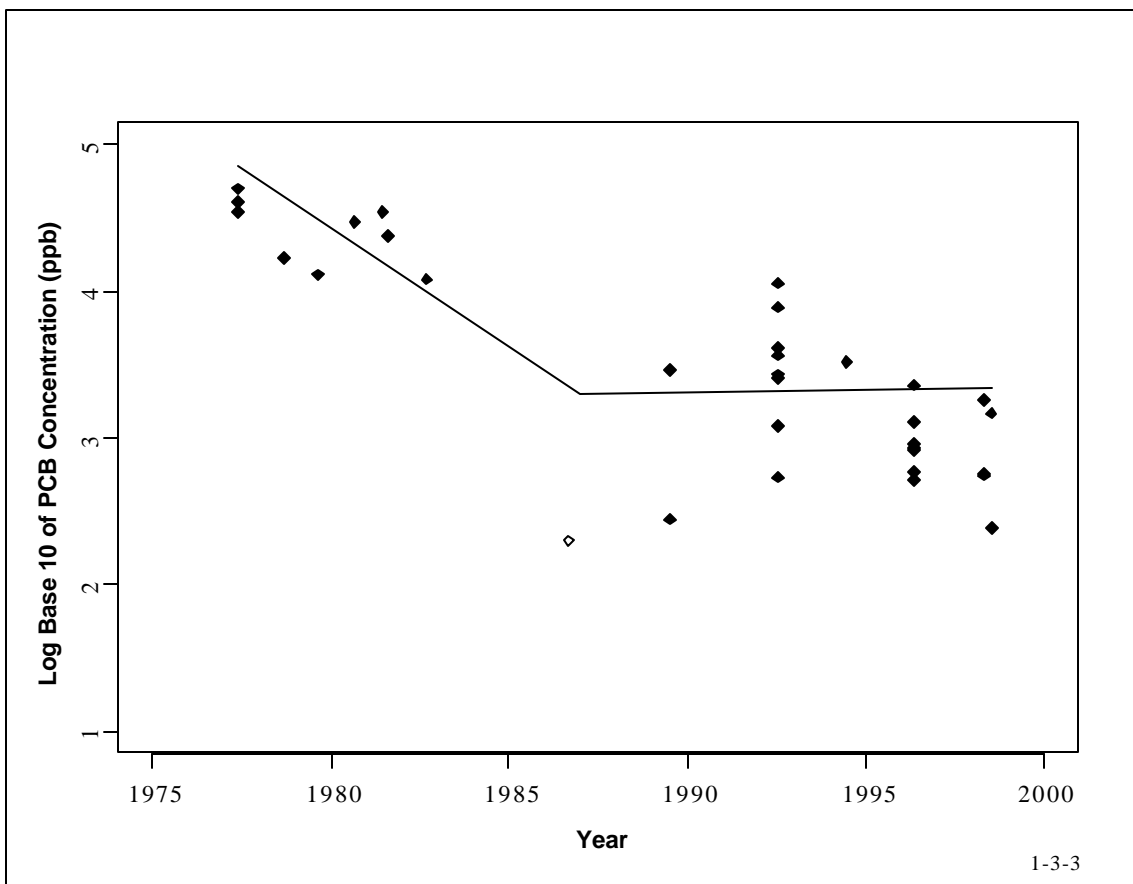


Figure 32 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Carp, Whole Body, versus Time

Breakpoint = 1987 ($p = 0.03$), Final Slope (\log_{10} PCB versus time) = 0.003 ($p = 0.9$), Rate of Change of PCB Concentration During Period of Final Slope = 0.7% (95% confidence interval: -12.3% to 15.6%). Any values below detection limit are depicted as \diamond .

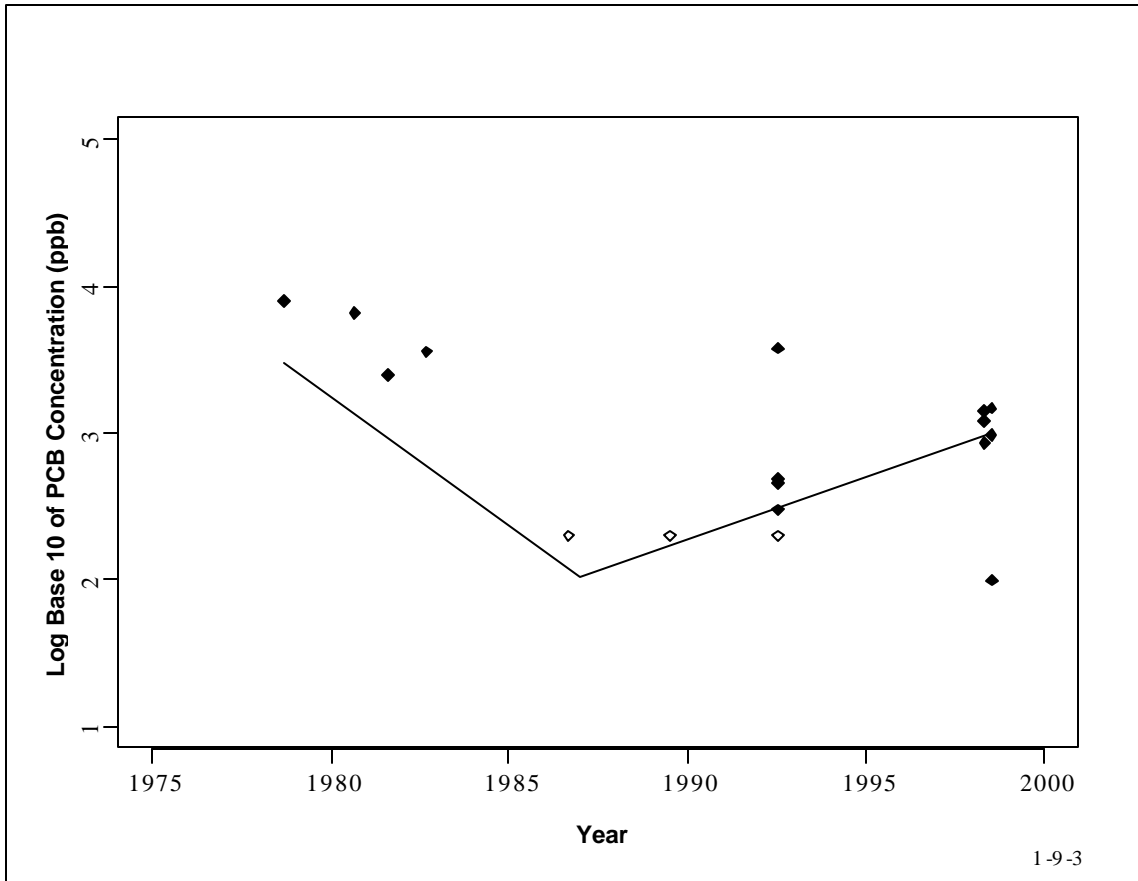


Figure 33 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Walleye, Whole Body, versus Time

Breakpoint = 1987 ($p = 0.009$), Final Slope (\log_{10} PCB versus time) = 0.084 ($p = 0.09$), Rate of Change of PCB Concentration During Period of Final Slope = 21.5% (95% confidence interval: -3.5% to 52.9%). Any values below detection limit are depicted as \diamond .

Table 14 PCB Time Trend Results for Fish Samples in Little Lake Butte des Morts Reach

| Species | Sample Type | Year of Break-point | n | Final (post-break) Slope | | | | 95% Confidence Interval | |
|---------------|----------------|---------------------|----|--------------------------|-------|-----------|------------|-------------------------|------|
| | | | | Final Slope | SE | p-value | % per Year | LCL | UCL |
| Carp | skin-on fillet | 1979 | 55 | -0.028 | 0.011 | 0.0177* | -6.1 | -10.9 | -1.1 |
| | whole body | 1987 | 40 | 0.003 | 0.030 | 0.9172 | 0.7 | -12.3 | 15.6 |
| Northern Pike | skin-on fillet | N/A | 19 | 0.055 | 0.011 | 0.0003*** | -11.8 | -16.7 | -6.7 |
| Walleye | skin-on fillet | 1990 | 63 | 0.015 | 0.025 | 0.5576 | 3.4 | -7.8 | 16.0 |
| | whole body | 1987 | 18 | 0.084 | 0.045 | 0.0874 | 21.5 | -3.5 | 52.9 |
| Yellow Perch | skin-on fillet | 1981 | 34 | 0.003 | 0.012 | 0.8025 | 0.7 | -5.0 | 6.8 |

Notes:

N/A – Not applicable; no breakpoint.

* $p < 0.05$

** $p < 0.01$

*** $p < 0.001$

Reach 2 — Appleton to Little Rapids

Only data for walleye can be analyzed for this reach. The data provide no evidence to reject the null hypothesis of a constant rate of decline over the time span of observation. $P = 0.5$ for the spline model versus the simple linear model (Table 13).

Table 15 PCB Time Trend Results for Fish Samples in Appleton to Little Rapids Reach

| Species | Sample Type | Year of Break-point | n | Final (post-break) Slope | | | | 95% Confidence Interval | |
|---------|----------------|---------------------|----|--------------------------|-------|----------|------------|-------------------------|------|
| | | | | Final Slope | SE | p-value | % per Year | LCL | UCL |
| Walleye | skin-on fillet | N/A | 30 | -0.046 | 0.014 | 0.0028** | -10.0 | -15.7 | -3.9 |

Notes:

N/A – Not applicable; no breakpoint.

* $p < 0.05$

** $p < 0.01$

*** $p < 0.001$

Reach 3 — Little Rapids to De Pere

No fish species with both an adequate sample size and sufficient spread of samples over time for analysis occurred in this reach.

Reach 4 — De Pere to Green Bay

In this reach, six of the seven fish categories show no significant improvement in fit of the spline model over the linear model. Figure 34 shows an example where the linear model fits quite well. For one species, though, a model with a change point in 1995 fits significantly better than the linear model (De Pere to Green Bay, carp, whole body). As seen in Figure 35, this model shows a large increase in log PCB concentration between 1997 and 1999. The substantial number of samples at these two time points may in fact represent a real increase in PCB concentration.

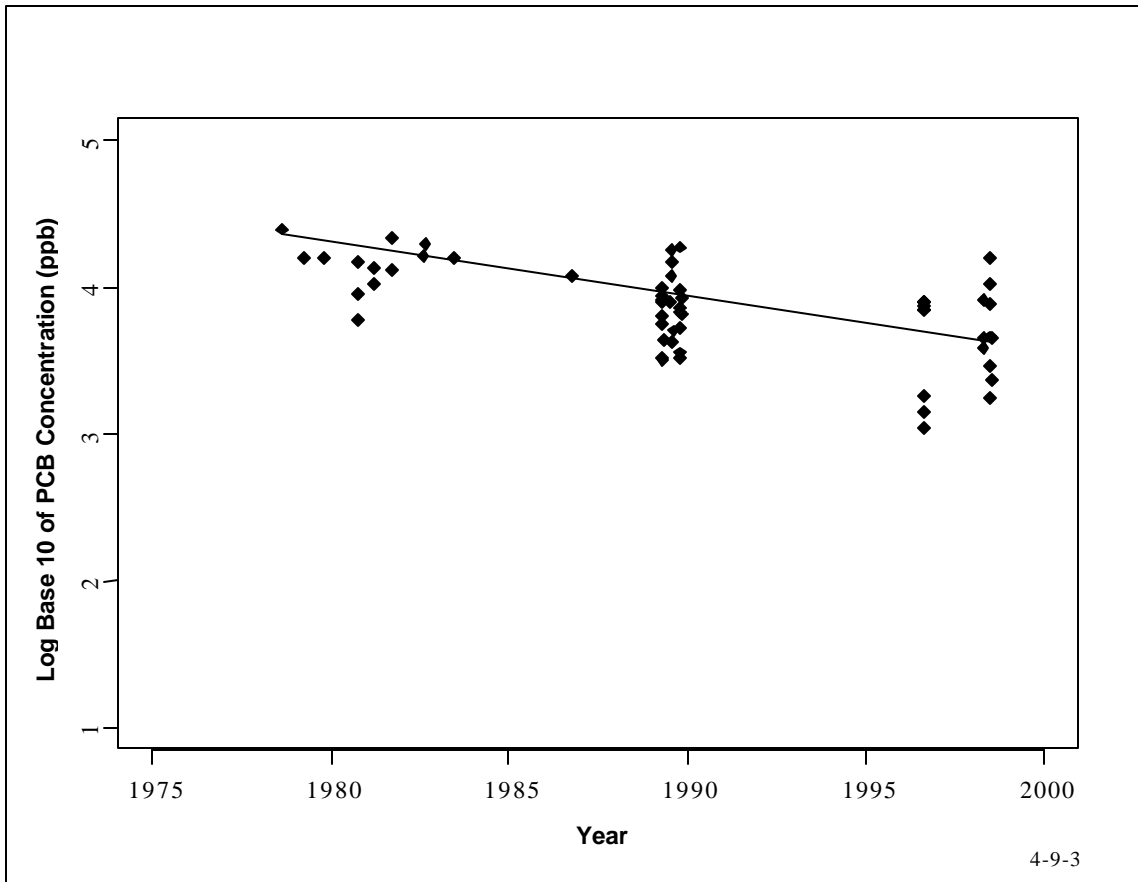


Figure 34 Log₁₀ PCB Concentration (ppb) in De Pere to Green Bay Walleye, Whole Body, versus Time

No Breakpoint, Final Slope (log₁₀ PCB versus time) = -0.037 ($p < 0.0001$), Rate of Change of PCB Concentration During Period of Final Slope = -8.1% (95% confidence interval: -10.4% to -5.8%).

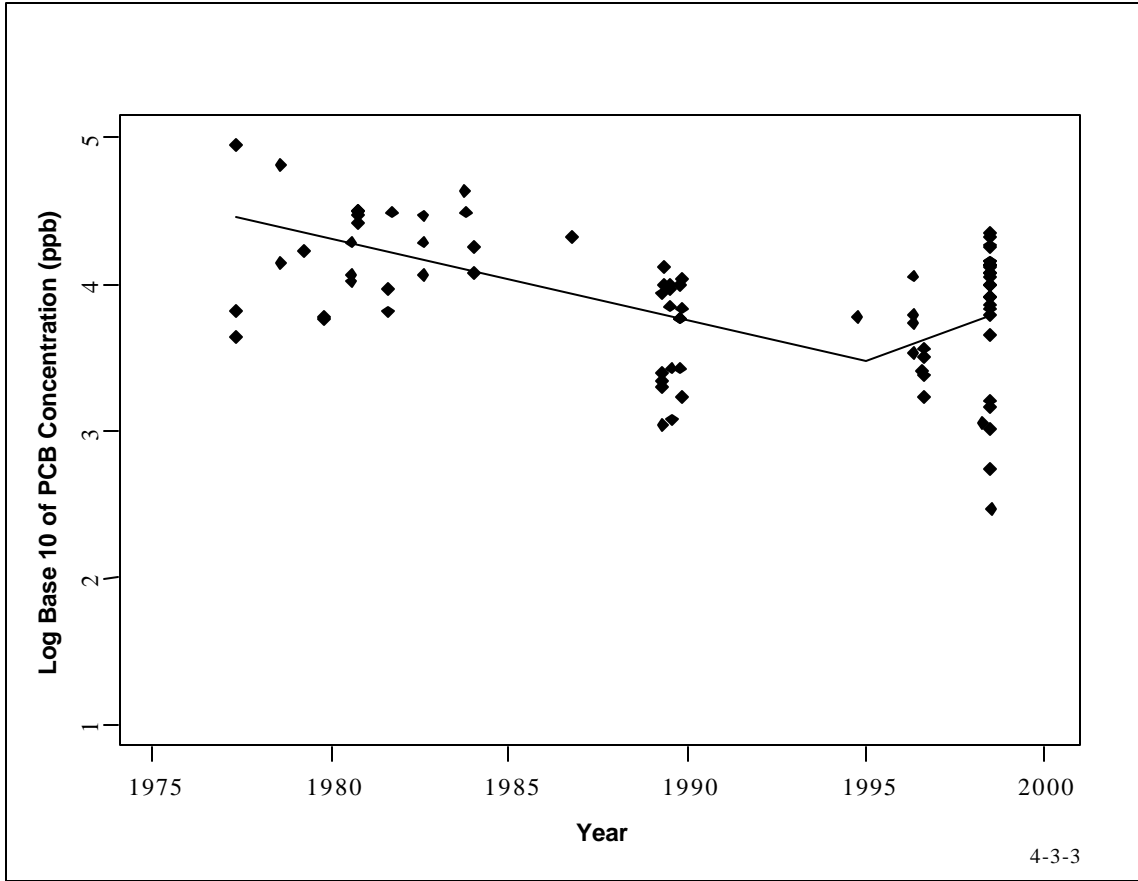


Figure 35 Log₁₀ PCB Concentration (ppb) in De Pere to Green Bay Carp, Whole Body, versus Time

Breakpoint = 1995 ($p = 0.009$), Final Slope (\log_{10} PCB versus time) = 0.086 ($p = 0.03$), Rate of Change of PCB Concentration During Period of Final Slope = 21.8% (95% confidence interval: 2.2% to 45.0%).

The non-significant ($p = 0.09$) meta-analysis for this reach indicates only weak evidence to reject the overall null hypothesis of a constant rate of change for all species within this reach over the time span of observation (Table 13). The meta-analysis partially remedies the problem of multiple comparisons. That is, if one conducts seven independent hypothesis tests and uses the standard criterion $p < 0.05$ to designate statistical significance, the probability of finding at least one significant p -value out of these seven tests, when the null hypothesis is really true, approaches 30 percent. This considerably exceeds the 5 percent false positives behind “ $p < 0.05$.” Thus, the single significant breakpoint for this reach in Table 16 with $p = 0.009$ may have occurred by chance.

Table 16 PCB Time Trend Results for Fish Samples in De Pere to Green Bay Reach

| Species | Sample Type | Year of Break-point | n | Final (post-break) Slope | | | | 95% Confidence Interval | |
|---------------|----------------|---------------------|-----|--------------------------|-------|------------|------------|-------------------------|------|
| | | | | Final Slope | SE | p-value | % per Year | LCL | UCL |
| Carp | whole body | 1995 | 90 | 0.086 | 0.038 | 0.0277* | 21.8 | 2.2 | 45.0 |
| Gizzard Shad | whole body | N/A | 19 | -0.023 | 0.005 | 0.0002*** | -5.1 | -7.2 | -2.9 |
| Northern Pike | skin-on fillet | N/A | 40 | -0.046 | 0.007 | <0.0001*** | -10.0 | -13.0 | -6.8 |
| Walleye | skin-on fillet | N/A | 120 | -0.032 | 0.004 | <0.0001*** | -7.2 | -8.7 | -5.6 |
| | whole body | N/A | 58 | -0.037 | 0.005 | <0.0001*** | -8.1 | -10.4 | -5.8 |
| White Bass | skin-on fillet | N/A | 58 | -0.021 | 0.006 | 0.0020** | -4.7 | -7.5 | -1.8 |
| White Sucker | skin-on fillet | N/A | 44 | -0.036 | 0.006 | <0.0001*** | -7.9 | -10.3 | -5.5 |

Notes:

N/A – Not applicable; no breakpoint.

* $p < 0.05$

** $p < 0.01$

*** $p < 0.001$

Reach 5 — Green Bay Zone 2

In the final reach considered, two of the fish categories show a highly significant improvement of the change point model over the linear model. For carp whole body samples, PCB concentration rises sharply until 1983 and then drops. Prior to 1983, there were samples for only five fish at two distinct time points. A similar pattern holds for carp fillet with skin samples, though the spline is statistically non-significant compared to the linear model.

For yellow perch skin-on fillet, we rejected the spline model, even though it formally provided a “better” fit, which can be seen in Figure 36. In this model, one finds a very steep fitted decrease until 1986, followed by a fitted step increase. The huge amplitude of the estimated seasonal effect, however, exceeds by five- or ten-fold that for other fish categories. These strange results raised the concern that we may have over-fit the model for this species. The spline model for Figure 36 relied on 19 samples collected at seven distinct time points. There are six parameters in time (intercept, final slope, initial slope difference, location of breakpoint, sine and cosine of time of year) and only seven distinct time points. Introducing as many parameters in time as time points risks over-fitting and uncertain or erroneous estimates.

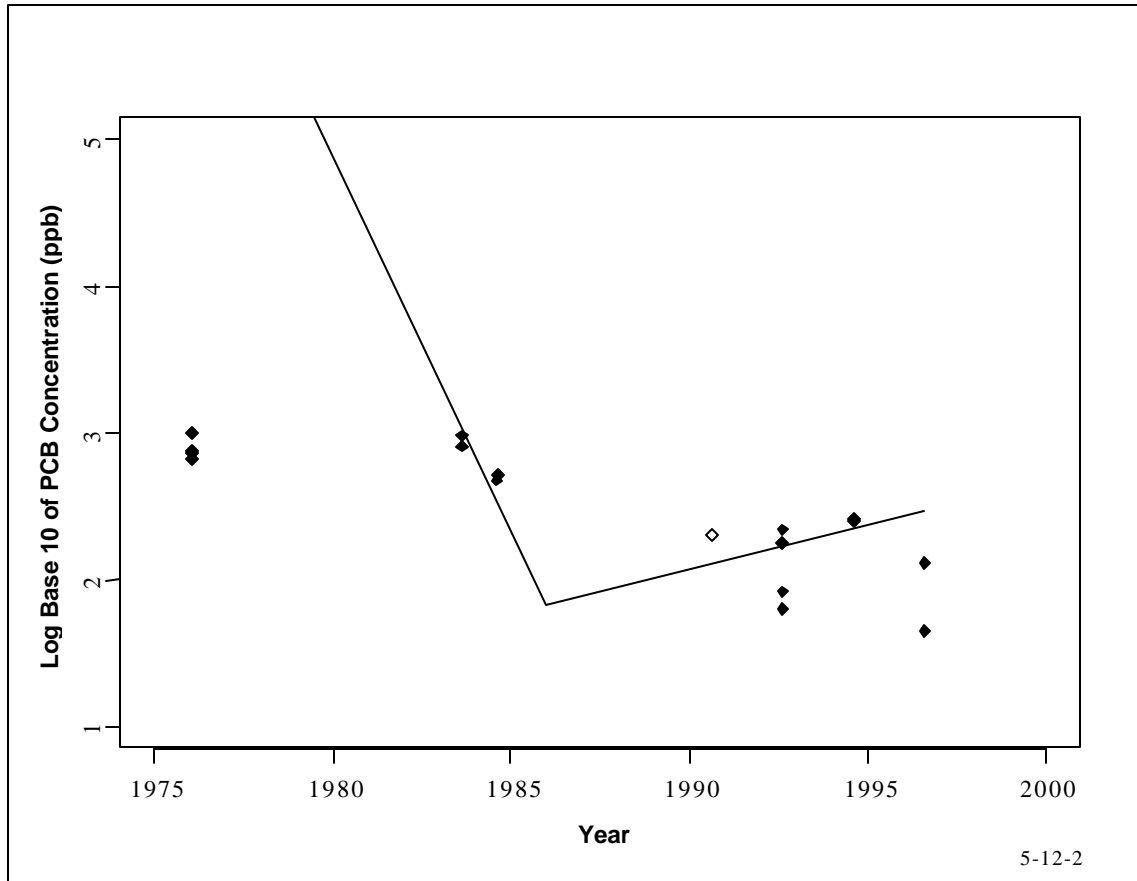


Figure 36 Rejected Spline Model for Green Bay Zone 2 Yellow Perch, Skin-on Fillet

Let us explain over-fitting by analogy. Suppose we choose six distinct time points. At each time point we randomly generate 10 values for $\log(PCB)$ as if 10 fish were sampled at that time point, for a total of 60 values. Then we fit a polynomial with six parameters (powers of time = X , from constant— X^0 —through X^5) and plot the raw data and fitted line on a scatter plot. This polynomial will fit perfectly in time—it will go exactly through the mean value at each time point. Of course, it will probably generate an implausible curve that varies drastically, perhaps with extremely large peaks or valleys between time points. This hypothetical example speaks to our situation. Fitting our model with six parameters in time mirrors fitting a polynomial with six parameters and, therefore, may give ridiculous results. In the example of yellow perch fillet with skin, we encounter only one additional, distinct time point (seven time points instead of six), which reduces but does not eliminate the risk of over-fitting. We recommend discarding the fitted model with a breakpoint at 1986 for yellow perch fillet with skin in this reach, as it exemplifies over-fitting. Therefore, we will use the simple linear model as the best-fitting model for these data (Figure 37). The model provides not only a more plausible fit, but a visually acceptable fit as well.

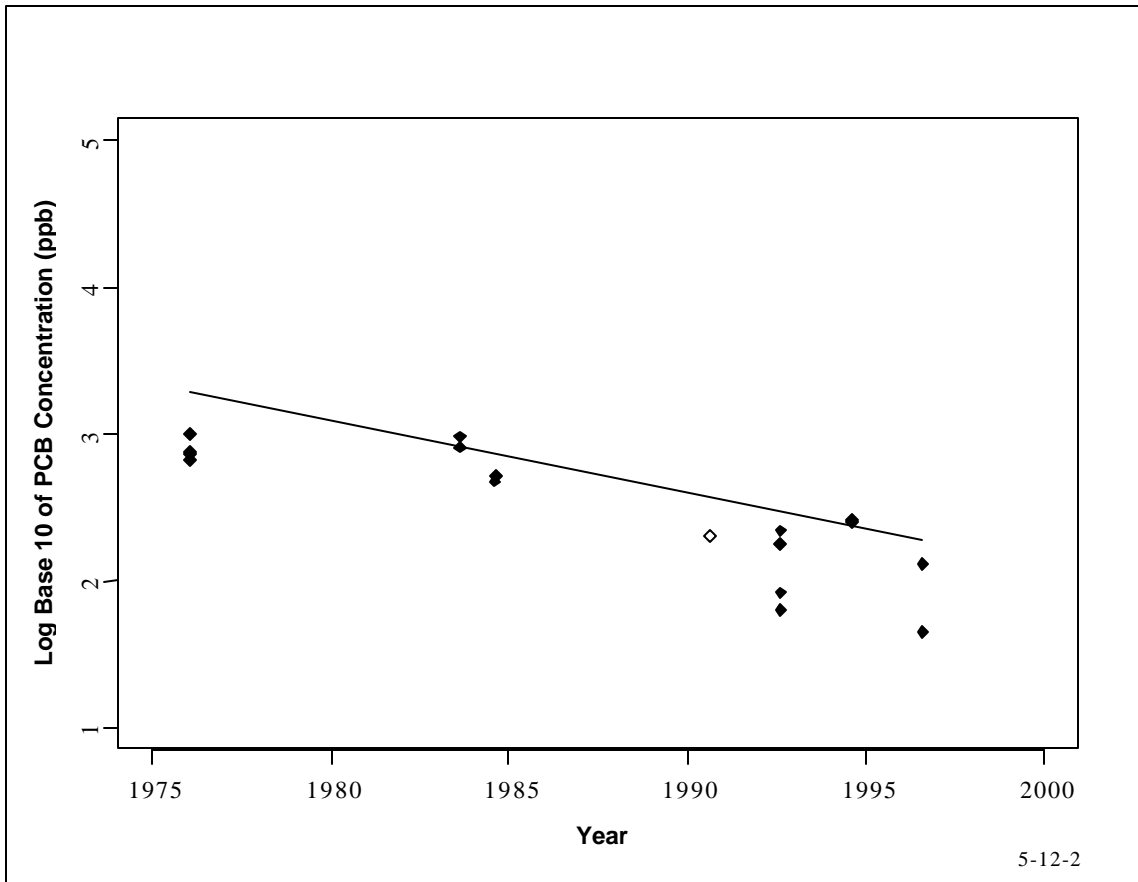


Figure 37 Log₁₀ PCB Concentration in Green Bay Zone 2 Yellow Perch, Skin-on Fillet, versus Time

No Breakpoint, Final Slope (log₁₀ PCB versus time) = -0.049 ($p = 0.004$), Rate of Change of PCB Concentration During Period of Final Slope = -10.7% (95% confidence interval: -16.8% to -4.2%). Any values below detection limit are depicted as \diamond .

Table 17 PCB Time Trend Results for Fish Samples in Green Bay Zone 2

| Species | Sample Type | Year of Break-point | n | Final (post-break) Slope | | | | 95% Confidence Interval | |
|--------------|----------------|---------------------|----|--------------------------|-------|------------|------------|-------------------------|-------|
| | | | | Final Slope | SE | p-value | % per Year | LCL | UCL |
| Alewife | whole body | N/A | 44 | -0.018 | 0.009 | 0.0497* | -4.0 | -7.8 | 0.0 |
| Carp | skin-on fillet | N/A | 28 | -0.023 | 0.015 | 0.1557 | -5.1 | -11.8 | 2.2 |
| | whole body | 1983 | 57 | -0.073 | 0.010 | <0.0001*** | -15.5 | -19.5 | -11.4 |
| Gizzard Shad | whole body | N/A | 32 | 0.025 | 0.010 | 0.0144* | 5.9 | 1.2 | 10.8 |
| Yellow Perch | skin-on fillet | N/A | 19 | -0.049 | 0.014 | 0.0038** | -10.7 | -16.8 | -4.2 |

Notes:

N/A – Not applicable; no breakpoint.

* $p < 0.05$ ** $p < 0.01$ *** $p < 0.001$ **Impact of Seasonality and Lipid Content on Best-fitting Model**

For each fish category (reach/species/type combination), we determined the best-fitting model, either the linear model or the spline model with one breakpoint, if that showed a significantly better fit than the linear model. Table 18 shows details of the fitted models.

From left to right, Table 18 shows the year of the breakpoint or “N/A” for no breakpoint) in units of \log_{10} (PCB concentration as ppb) per year and the standard error and p -value of the slope; the rate of change per year as a percentage along with a 95 percent confidence interval for the percentage; the difference between early and late slope, if applicable, in units of \log_{10} (PCB concentration as ppb) per year, along with the standard error and p -value for the difference between early and late slope; the coefficient of \log_{10} (lipid percent) and its standard error and p -value; and the month of the maximum amplitude of the seasonal effect and the amplitude (A) and the p -value for the seasonal effect. The quantities 10^A and 10^{-A} are multipliers that show the relative increase or decrease, respectively, of the seasonal maximum or minimum compared to the annual mean.

We note some interesting features about the covariates in Table 18. The coefficient of log of percent lipid departs significantly from zero for almost all fish categories. This coefficient approaches one for many fish categories, meaning that an analysis using the log of lipid-normalized PCB concentration as the outcome variable, without including percent lipids as a covariate, would be approximately correct. (As noted earlier, lipid normalization is usually calculated as PCB concentration divided by the percent lipid in the tissue.) Yet for several

species the coefficient fails to reach 1. This suggests that traditional lipid normalization alone does not control the lipid contribution adequately. The amplitude of the seasonal effect is significantly non-zero for the majority of fish categories, falling mainly in the 0.2 to 0.6 range. We define the amplitude as the height of the seasonal sine curve from zero to the maximum on the log scale, so the range from minimum to maximum is twice this value. On the log scale, the majority of species would fall between 0.4 and 1.2. Calculating the antilog of 0.4 and 1.2 (i.e., 10 raised to that power) tells us that the ratio of maximum to minimum over a year ranges from 2.5 to 16 for the majority of species. This represents substantial seasonal variation. The month in which the peak PCB concentration occurs varies quite a bit across fish categories. A footnote to Appendix Table A-3 explains how to calculate estimated PCB concentration for any time of year, taking account of seasonal variation.

As seen in the plots, we observe quite a bit of variation in log of PCB concentration around the fitted line. Even fish samples taken at the same time vary greatly in PCB concentration. The residual standard deviation (SD), after fitting the model, measures the magnitude of this variation. Using the approximation of plus or minus two SDs allows us to estimate the range, which covers most of the data (from low to high end), at about four SDs. From an appendix table, most of the standard deviation values (calculated as the square root of the mean squared error) fall between 0.15 and 0.35. Four SDs is thus between 0.60 and 1.40 for most species. Taking the antilog of 0.60 and 1.40 gives 4.0 and 25, respectively. This implies very high variation in PCB concentration for a particular reach/species/type: for species with the least variation, the values differ from the low end to the high end by roughly a factor of four, corresponding to an SD of 0.15. That is, for the species with an SD of 0.15, it would not be uncommon to find different samples with a fourfold difference in PCB concentration when sampled at the same time of year and with the same lipid content (e.g., whole body alewife, in Green Bay Zone 2 has an SD = 0.17, similar to 0.15). For species with an SD of 0.35 (such as carp fillet with skin, Little Lake Butte des Morts), it would not be uncommon to find samples differing by a factor of 25 in PCB concentration. Figure 31 shows just such variation and supports the notion of highly variable PCB concentrations within species.

Table 18 Model Parameters and Other Statistics for the Best-fitting Model

| Species | Sample Type | Year of Break-point | n | Final (post-break) Slope | | | | 95% Confidence Interval | | Pre-break Slope Minus Final Slope | SE | p-value Slope Change | Coef-ficient of Log (% lipid) | SE | p-value for Log (% lipid) | Seasonal Peak | | p-value for Seasonal Effect |
|------------------------------------|----------------|---------------------|-----|--------------------------|-------|------------|------------|-------------------------|-------|-----------------------------------|-------|----------------------|-------------------------------|--------|---------------------------|---------------|------------|-----------------------------|
| | | | | Final Slope | SE | p-value | % per Year | LCL | UCL | | | | | | | Mo. | Ampli-tude | |
| <i>Little Lake Butte des Morts</i> | | | | | | | | | | | | | | | | | | |
| Carp | skin-on fillet | 1979 | 55 | -0.028 | 0.011 | 0.0177* | -6.1 | -10.9 | -1.1 | -0.228 | 0.085 | 0.0102 | 0.87 | 0.15 | 0.0000 | 12.9 | 0.39 | 0.0078 |
| | whole body | 1987 | 40 | 0.003 | 0.30 | 0.9172 | 0.7 | -12.3 | 15.6 | -0.165 | 0.059 | 0.0084 | 0.86 | 0.33 | 0.0131 | 7.0 | 0.83 | 0.0025 |
| Northern Pike | skin-on fillet | N/A | 19 | -0.055 | 0.011 | 0.0003*** | -11.8 | -16.7 | -6.7 | | | 0.45 | 0.30 | 0.1554 | 1.3 | 0.67 | 0.1594 | |
| Walleye | skin-on fillet | 1990 | 63 | 0.015 | 0.025 | 0.5576 | 3.4 | -7.8 | 16.0 | -0.095 | 0.037 | 0.0140 | 0.50 | 0.15 | 0.0011 | 11.6 | 0.20 | 0.0273 |
| | whole body | 1987 | 18 | 0.084 | 0.045 | 0.0874 | 21.5 | -3.5 | 52.9 | -0.261 | 0.080 | 0.0069 | 0.99 | 0.36 | 0.0185 | 11.6 | 0.46 | 0.0040 |
| Yellow Perch | skin-on fillet | 1981 | 34 | 0.003 | 0.012 | 0.8025 | 0.7 | -5.0 | 6.8 | -0.247 | 0.077 | 0.0034 | 0.49 | 0.21 | 0.0236 | 7.0 | 0.22 | 0.0007 |
| <i>Appleton</i> | | | | | | | | | | | | | | | | | | |
| Walleye | skin-on fillet | N/A | 30 | -0.046 | 0.014 | 0.0028** | -10.0 | -15.7 | -3.9 | | | 1.08 | 0.16 | 0.0000 | 8.1 | 0.43 | 0.0010 | |
| <i>De Pere</i> | | | | | | | | | | | | | | | | | | |
| Carp | whole body | 1995 | 90 | 0.086 | 0.038 | 0.0277* | 21.8 | 2.2 | 45.0 | -0.141 | 0.044 | 0.0022 | 0.79 | 0.11 | 0.0000 | 6.7 | 0.06 | 0.0004 |
| Gizzard Shad | whole body | N/A | 19 | -0.023 | 0.005 | 0.0002*** | -5.1 | -7.2 | -2.9 | | | 0.51 | 0.09 | 0.001 | 8.6 | 0.58 | 0.0000 | |
| Northern Pike | skin-on fillet | N/A | 40 | -0.046 | 0.007 | <0.0001*** | -10.0 | -13.0 | -6.8 | | | 0.72 | 0.17 | 0.0001 | 10.1 | 0.17 | 0.3531 | |
| Walleye | skin-on fillet | N/A | 120 | -0.032 | 0.004 | <0.0001*** | -7.2 | -8.7 | -5.6 | | | 0.85 | 0.07 | 0.0000 | 9.5 | 0.02 | 0.7566 | |
| | whole body | N/A | 58 | -0.037 | 0.005 | <0.0001*** | -8.1 | -10.4 | -5.8 | | | 0.44 | 0.12 | 0.0007 | 7.0 | 0.12 | 0.2038 | |
| White Bass | skin-on fillet | N/A | 58 | -0.021 | 0.006 | 0.0020** | -4.7 | -7.5 | -1.8 | | | 0.82 | 0.11 | 0.0000 | 6.7 | 0.33 | 0.1043 | |
| White Sucker | skin-on fillet | N/A | 44 | -0.036 | 0.006 | <0.0001*** | -7.9 | -10.3 | -5.5 | | | 0.43 | 0.15 | 0.0071 | 6.9 | 0.08 | 0.5528 | |
| <i>Green Bay Zone 2</i> | | | | | | | | | | | | | | | | | | |
| Alewife | whole body | N/A | 44 | -0.018 | 0.009 | 0.0497* | -4.0 | -7.8 | 0.0 | | | 0.91 | 0.14 | 0.0000 | 6.1 | 0.17 | 0.0335 | |
| Carp | skin-on fillet | N/A | 28 | -0.023 | 0.015 | 0.1557 | -5.1 | -11.8 | 2.2 | | | 0.76 | 0.15 | 0.0000 | 3.9 | 0.24 | 0.0288 | |
| | whole body | 1983 | 57 | -0.073 | 0.010 | <0.0001*** | -15.5 | -19.5 | -11.4 | 0.266 | 0.059 | 0.0000 | 0.90 | 0.10 | 0.0000 | 6.9 | 0.24 | 0.0000 |
| Gizzard Shad | whole body | N/A | 32 | 0.025 | 0.010 | 0.0144* | 5.9 | 1.2 | 10.8 | | | -0.13 | 0.12 | 0.2811 | 2.6 | 0.34 | 0.0300 | |

Table 18 Model Parameters and Other Statistics for the Best-fitting Model

| Species | Sample Type | Year of Break-point | n | Final (post-break) Slope | | | | 95% Confidence Interval | | Pre-break Slope Minus Final Slope | SE | p-value Slope Change | Coef-ficient of Log (% lipid) | SE | p-value for Log (% lipid) | Seasonal Peak | | p-value for Seasonal Effect |
|--------------|----------------|---------------------|----|--------------------------|-------|----------|------------|-------------------------|------|-----------------------------------|----|----------------------|-------------------------------|-------|---------------------------|---------------|------------|-----------------------------|
| | | | | Final Slope | SE | p-value | % per Year | LCL | UCL | | | | | | | Mo. | Ampli-tude | |
| Yellow Perch | skin-on fillet | N/A | 19 | -0.049 | 0.014 | 0.0038** | -10.7 | -16.8 | -4.2 | | | 1.09 | 0.47 | 0.353 | 4.7 | 0.45 | 0.5489 | |

Notes:

N/A – Not applicable; no breakpoint.

* $p < 0.05$

** $p < 0.01$

*** $p < 0.001$

5.2.2 Best-fitting Model, Meta-analysis, Sensitivity Analysis, and Future Projections

In the preceding section, Table 13 and the related discussion presented decisions for each reach, species, and sample type on the choice between a model including a breakpoint in the time trend and a model without a time trend. Accepting that decision, Table 14 through Table 17 presented the final slopes for the best-fitted models. Table 18 presented additional parameters for each best-fitted model. The Appendix includes the full set of parameters for each model.

Table 19 shows the results of meta-analyses for each reach. The final row in each reach gives a combined species analysis. The combined post-breakpoint slope is a weighted average of all the slopes within this reach, weighted by the inverse of the standard error squared. The inverse standard error squared provides weights leading to a minimum variance of the weighted mean estimate in many common sampling situations. Unlike the meta-analysis of surface sediments introduced in Section 4.4.1 (Table 10), where PCB mass provided a natural set of weights, there is no *a priori* set of weights available to use with fish. Thus, weights with good statistical properties have been chosen for the fish meta-analysis. This weighting gives high weights to more precise estimates, usually based on a large sample size, and low weights to imprecise estimates, usually derived from small sample sizes. The *p*-value (based on the normal distribution) tests whether this summary slope differs significantly from zero.

The fish species included in the meta-analysis have diverse habitats, lifecycles, and feeding patterns. Nevertheless, the PCB concentration in each species serves as a sentinel of PCBs in their environment. Just as the economic growth rate of each unique industrial sector of a nation can combine into a single growth rate for a national economy, the time trends of diverse species can combine into a meaningful descriptive statistical time trend for fish species in a reach. This summary rate of change cannot replace the individual species' rates of change. It means only what its definition implies: weighting more heavily on species with more precise slope estimates and less heavily on species with less precise slope estimates provides a reach mean slope which can be compared to zero. An individual species may possibly have a real slope that differs substantially from the combined reach slope. While the combined slope is a summary, the individual slopes cannot be ignored. Also, as noted in Section 4.4.1 in reference to sediment, the combined slope should not be used to project PCB concentrations for all species in the reach.

In addition to the combined reach slope in Table 19, the percent rate of change of PCB concentration implied by the combined slope, *b*, is also presented, using the following equation:

Equation 17

$$\text{percent change} = 100 * (10^b - 1)$$

The 95 percent confidence interval for the percent change is also shown in the table (calculated by deriving the 95 percent confidence interval for the slope of log PCB concentration versus time—using the normal distribution and converting the upper and lower confidence bounds to percentages by Equation 17).

In this section, we also address an issue of uncertainty associated with the breakpoint. As mentioned in the methods section, the standard errors for time trend slopes and *p*-values for the best-fitting model do not incorporate the variation due to estimating the location of the breakpoint. They therefore underestimate the uncertainty in the time trend slope. The standard errors shown in the table are too small for those species where the model has a breakpoint. We addressed this problem by performing a sensitivity analysis for each of the seven reach/species/type combinations with a breakpoint model. We identified the earliest and the latest breakpoints that were “plausible,” as described in the methods section. Table 20 shows results for these “earliest” and “latest” models, when there is a breakpoint.

Table 19 Meta-analysis of Fish Time Trends

| Species | Sample Type | Log ₁₀ (PCB) Time Trend Final Slope Estimate | Standard Error | Statistical Weight ⁺ | <i>p</i> -value | Annual % Change in PCB Concen- tration | % Change 95% Lower Bound | % Change 95% Upper Bound |
|------------------------------------|----------------|--|-------------------|------------------------------------|-------------------|---|-----------------------------------|-----------------------------------|
| <i>Little Lake Butte des Morts</i> | | | | | | | | |
| Carp | skin-on fillet | -0.028 | 0.011 | 0.31 | | | | |
| | whole body | 0.003 | 0.30 | 0.05 | | | | |
| Northern Pike | skin-on fillet | -0.055 | 0.011 | 0.30 | | | | |
| Walleye | skin-on fillet | 0.015 | 0.025 | 0.06 | | | | |
| | whole body | 0.084 | 0.045 | 0.02 | | | | |
| Yellow Perch | skin-on fillet | 0.003 | 0.012 | 0.26 | | | | |
| Combined | | -0.022 | 0.006 | 1.00 | 0.0006 | -4.9 | -7.5 | -2.1 |
| <i>Appleton</i> | | | | | | | | |
| Walleye | skin-on fillet | -0.056 | 0.016 | | 0.003 | -10.0 | -17.9 | -5.6 |
| <i>De Pere</i> | | | | | | | | |
| Carp | whole body | 0.086 | 0.038 | 0.00 | | | | |
| Gizzard Shad | whole body | -0.023 | 0.005 | 0.21 | | | | |
| Northern Pike | skin-on fillet | -0.046 | 0.007 | 0.08 | | | | |
| Walleye | skin-on fillet | -0.032 | 0.004 | 0.32 | | | | |
| | whole body | -0.037 | 0.005 | 0.15 | | | | |
| White Bass | skin-on fillet | -0.021 | 0.006 | 0.10 | | | | |
| White Sucker | skin-on fillet | -0.036 | 0.006 | 0.14 | | | | |
| Combined | | -0.031 | 0.002 | 1.00 | <0.0001 | -6.9 | -7.8 | -6.0 |

Table 19 Meta-analysis of Fish Time Trends

| Species | Sample Type | Log ₁₀ (PCB) Time Trend Final Slope Estimate | Standard Error | Statistical Weight ⁺ | p-value | Annual % Change in PCB Concen- tration | % Change 95% Lower Bound | % Change 95% Upper Bound |
|-------------------------|----------------|--|-------------------|------------------------------------|-------------------|---|-----------------------------------|-----------------------------------|
| <i>Green Bay Zone 2</i> | | | | | | | | |
| Alewife | whole body | -0.018 | 0.009 | 0.31 | | | | |
| Carp | skin-on fillet | -0.023 | 0.015 | 0.10 | | | | |
| | whole body | -0.073 | 0.010 | 0.22 | | | | |
| Gizzard Shad | whole body | 0.025 | 0.010 | 0.26 | | | | |
| Yellow Perch | skin-on fillet | -0.049 | 0.014 | 0.12 | | | | |
| Combined | | -0.033 | 0.007 | 1.00 | <0.0001 | -5.1 | -7.2 | -3.0 |

Note:

- + Statistical weight is proportional to the inverse of the squared standard error. Weights sum to 1.0 within each reach.

Figure 38 captures the estimated percent change per year for the best-fitting model for each fish category. The confidence intervals shown in these plots obtain from the results of the best-fitting model and do not incorporate the extra uncertainty due to estimating the location of the breakpoint. Therefore, the reader must remember that the plotted confidence intervals are too narrow for the seven analyses with a breakpoint.

Table 20 Final Slope and Percent Change per Year for Best-fitting Model and Sensitivity Analysis

| Species | Sample | | Best-fitting Model | | | Earliest Breakpoint | | | Latest Breakpoint | | |
|------------------------------------|----------------|-----|--------------------|-------------------|-----------------|---------------------|--------------------------------|-----------------|-------------------|--------------------------------|-----------------|
| | Type | n | Break point Year | % Change per Year | p-value (% = 0) | Year | Final Slope: % Change per Year | p-value (% = 0) | Year | Final Slope: % Change per Year | p-value (% = 0) |
| <i>Little Lake Butte des Morts</i> | | | | | | | | | | | |
| Carp | skin-on fillet | 55 | 1979 | -6.15 | 0.0177 | 1979 | -6.15 | 0.0177 | 1985 | -1.56 | 0.7419 |
| | whole body | 40 | 1987 | 0.71 | 0.9172 | 1985 | -4.04 | 0.5264 | 1990 | -0.25 | 0.9765 |
| Northern Pike | skin-on fillet | 19 | N/A | -11.83 | 0.0003 | | | | | | |
| Walleye | skin-on fillet | 63 | 1990 | 3.44 | 0.5576 | 1979 | -8.37 | 0.0000 | 1994 | 8.82 | 0.4482 |
| | whole body | 18 | 1987 | 21.47 | 0.0874 | 1984 | 15.10 | 0.2024 | 1990 | 21.11 | 0.1324 |
| Yellow Perch | skin-on fillet | 34 | 1981 | 0.73 | 0.8025 | 1979 | 0.27 | 0.9252 | 1996 | 333.61 | 0.0122 |
| <i>Appleton</i> | | | | | | | | | | | |
| Walleye | skin-on fillet | 30 | N/A | -9.97 | 0.0028 | | | | | | |
| <i>De Pere</i> | | | | | | | | | | | |
| Carp | whole body | 90 | 1995 | 21.76 | 0.0277 | 1990 | -0.69 | 0.8232 | 1996 | 29.80 | 0.0191 |
| Gizzard Shad | whole body | 19 | N/A | -5.07 | 0.0002 | | | | | | |
| Northern Pike | skin-on fillet | 40 | N/A | -9.95 | 0.0000 | | | | | | |
| Walleye | skin-on fillet | 120 | N/A | -7.19 | 0.0000 | | | | | | |
| | whole body | 58 | N/A | -8.11 | 0.0000 | | | | | | |
| White Bass | skin-on fillet | 58 | N/A | -4.72 | 0.0020 | | | | | | |
| White Sucker | skin-on fillet | 44 | N/A | -7.90 | 0.0000 | | | | | | |
| <i>Green Bay Zone 2</i> | | | | | | | | | | | |
| Alewife | whole body | 44 | N/A | -3.96 | 0.0497 | | | | | | |
| Carp | skin-on fillet | 28 | N/A | -5.06 | 0.1557 | | | | | | |
| | whole body | 57 | 1983 | -15.54 | 0.0000 | 1983 | -15.54 | 0.0000 | 1984 | -16.15 | 0.0000 |
| Gizzard Shad | whole body | 32 | N/A | 5.91 | 0.0144 | | | | | | |
| Yellow Perch | skin-on fillet | 19 | N/A | -10.75 | 0.0038 | | | | | | |

Note:

N/A – Not applicable; no breakpoint.

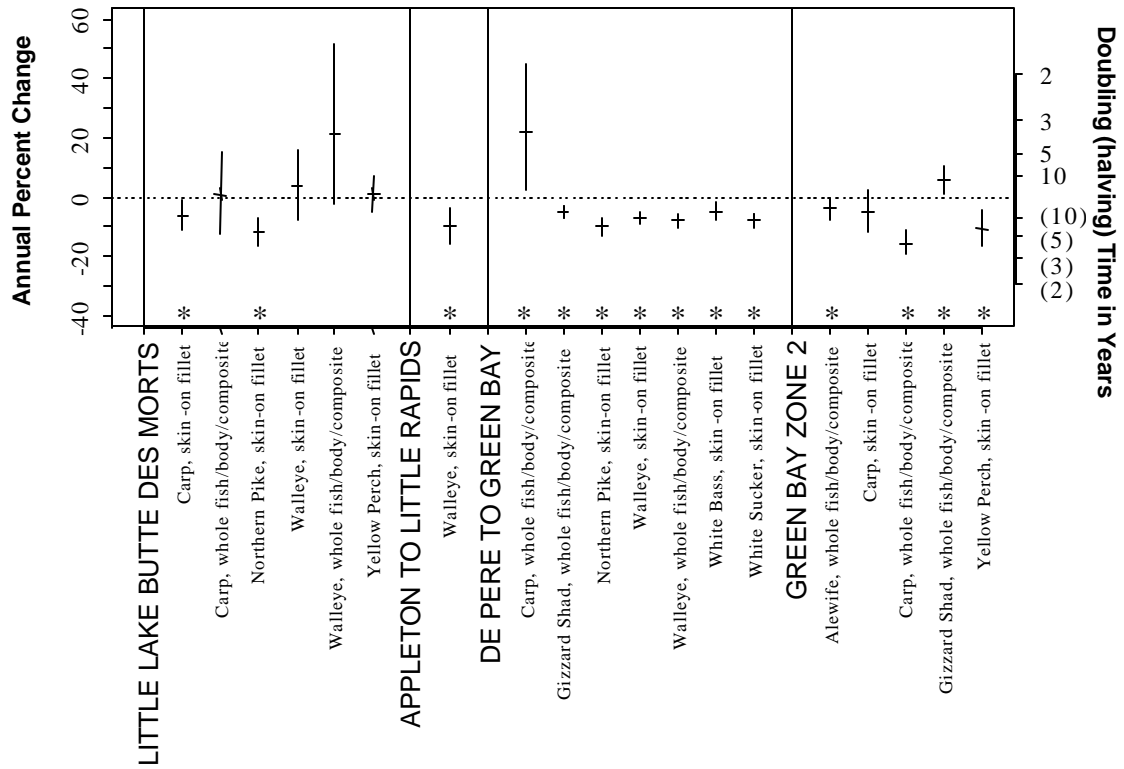


Figure 38 95 Percent Confidence Intervals Showing Annual Percent Rate of Change (Left Vertical Axis) in PCB Concentrations by Reach, Species, and Sample Type

An asterisk (*) indicates a rate of change that differs significantly from zero. Right vertical axis expresses time trend change in terms of doubling and halving times.

Table 21 shows projections into the future based on the best-fitting model, spline, or simple linear trend. We present the estimated mean PCB concentration in the years 1999 and 2020, with 95 percent confidence intervals for the concentration at each year. For fish categories with a negative final slope (the post-breakpoint slope for the spline models), the table also shows estimated times until PCB concentration drops below specified concentrations. The methods section provided the formulae for computing these quantities.

Table 21 Projecting into the Future—Predicted Mean PCB Concentration (ppb) in 1999 and 2020 and Time When Specified PCB Concentrations Will Be Reached

| Species | Sample Type | Year of Break point | Estimate of Mean PCB Concentration in 1999 | | | Estimate of Mean PCB Concentration in 2020 | | | Year in Which Specified PCB Concentration (ppb) Is Reached | | | | | | | | | |
|------------------------------------|----------------|---------------------|--|--------------|--------------|--|--------------|--------------|--|------|------|------|------|------|------|------|------|--|
| | | | Mean PCB (ppb) | Lower 95% CI | Upper 95% CI | Mean PCB (ppb) | Lower 95% CI | Upper 95% CI | 1,400 | 240 | 220 | 140 | 63 | 38 | 20 | 5 | 0.5 | |
| <i>Little Lake Butte des Morts</i> | | | | | | | | | | | | | | | | | | |
| Carp | skin-on fillet | 1979 | 1,399 | 589 | 3,319 | 369 | 56 | 2,429 | 1999 | 2027 | 2028 | 2035 | 2048 | 2056 | 2066 | 2088 | 2124 | |
| | whole body | 1987 | 2,506 | 1,055 | 5,954 | 2,910 | 78 | 109,080 | | | | | | | | | | |
| Northern Pike | skin-on fillet | N/A | 149 | 59 | 375 | 11 | 2 | 73 | 1981 | 1995 | 1996 | 1999 | 2006 | 2010 | 2015 | 2026 | 2044 | |
| Walleye | skin-on fillet | 1990 | 251 | 131 | 483 | 511 | 27 | 9,824 | | | | | | | | | | |
| | whole body | 1987 | 1,266 | 515 | 3,113 | 75,208 | 534 | 10,591,388 | | | | | | | | | | |
| Yellow Perch | skin-on fillet | 1981 | 255 | 110 | 590 | 296 | 40 | 2,173 | | | | | | | | | | |
| <i>Appleton</i> | | | | | | | | | | | | | | | | | | |
| Walleye | skin-on fillet | N/A | 376 | 117 | 1,212 | 41 | 3 | 496 | 1986 | 2003 | 2004 | 2008 | 2016 | 2021 | 2027 | 2040 | 2062 | |
| <i>De Pere</i> | | | | | | | | | | | | | | | | | | |
| Carp | whole body | 1995 | 7,526 | 5,439 | 10,414 | 470,285 | 9,207 | 24,021,513 | | | | | | | | | | |
| Gizzard Shad | whole body | N/A | 1,709 | 1,463 | 1,995 | 573 | 329 | 1,000 | 2003 | 2037 | 2038 | 2047 | 2062 | 2072 | 2085 | 2111 | 2156 | |
| Northern Pike | skin-on fillet | N/A | 542 | 364 | 807 | 60 | 25 | 145 | 1990 | 2007 | 2008 | 2012 | 2020 | 2024 | 2030 | 2044 | 2066 | |
| Walleye | skin-on fillet | N/A | 781 | 647 | 941 | 163 | 103 | 257 | 1991 | 2015 | 2016 | 2022 | 2033 | 2039 | 2048 | 2067 | 2098 | |
| | whole body | N/A | 4,343 | 3,384 | 5,575 | 736 | 374 | 1,449 | 2012 | 2033 | 2034 | 2040 | 2049 | 2055 | 2063 | 2079 | 2106 | |
| White Bass | skin-on fillet | N/A | 2,693 | 1,659 | 4,370 | 975 | 342 | 2,781 | 2013 | 2049 | 2051 | 2060 | 2077 | 2087 | 2100 | 2129 | 2177 | |
| White Sucker | skin-on fillet | N/A | 637 | 414 | 981 | 113 | 48 | 268 | 1989 | 2011 | 2012 | 2017 | 2027 | 2033 | 2041 | 2058 | 2086 | |
| <i>Green Bay Zone 2</i> | | | | | | | | | | | | | | | | | | |
| Alewife | whole body | N/A | 2,106 | 1,378 | 3,219 | 901 | 269 | 3,022 | 2009 | 2053 | 2055 | 2066 | 2086 | 2098 | 2114 | 2148 | 2205 | |

Table 21 Projecting into the Future—Predicted Mean PCB Concentration (ppb) in 1999 and 2020 and Time When Specified PCB Concentrations Will Be Reached

| Species | Sample Type | Year of Break point | Estimate of Mean PCB Concentration in 1999 | | | Estimate of Mean PCB Concentration in 2020 | | | Year in Which Specified PCB Concentration (ppb) Is Reached | | | | | | | | |
|--------------|----------------|---------------------|--|--------------|--------------|--|--------------|--------------|--|------|------|------|------|------|------|------|------|
| | | | Mean PCB (ppb) | Lower 95% CI | Upper 95% CI | Mean PCB (ppb) | Lower 95% CI | Upper 95% CI | 1,400 | 240 | 220 | 140 | 63 | 38 | 20 | 5 | 0.5 |
| Carp | skin-on fillet | N/A | 4,852 | 2,224 | 10,587 | 1,630 | 180 | 14,784 | 2023 | 2057 | 2059 | 2067 | 2083 | 2092 | 2105 | 2131 | 2176 |
| | whole body | 1983 | 1,468 | 935 | 2,305 | 42 | 10 | 175 | 1999 | 2010 | 2010 | 2013 | 2018 | 2021 | 2024 | 2033 | 2046 |
| Gizzard Shad | whole body | N/A | 3,159 | 2,129 | 4,687 | 10,549 | 2,965 | 37,524 | | | | | | | | | |
| Yellow Perch | skin-on fillet | N/A | 150 | 23 | 997 | 14 | 1 | 143 | 1979 | 1995 | 1996 | 2000 | 2007 | 2011 | 2017 | 2029 | 2049 |

Note:

N/A – Not applicable; no breakpoint.

All of the estimated times to reach specified concentrations in Table 21, as well as the estimated concentrations for 1999 and 2020, require extremely careful interpretation. We have based all of these estimates on the untestable assumption that the PCB concentration will continue to change in the future at the same rate as during the post-breakpoint period. In addition, as noted repeatedly, the confidence intervals for models that include a breakpoint do not incorporate the extra uncertainty related to breakpoint estimation and are too narrow.

A striking feature of the table is that most of the confidence intervals are very wide. For instance, for carp whole body in Little Lake Butte des Morts, the expected mean concentration in the year 2020 is 2,910 ppb, but the range is huge: 78 to 109,080 ppb. For those cases with a wide confidence interval in 2020 (or 1999), the time to reach specified concentrations (in the right half of the table) can also be expected to have a wide confidence interval.

We now discuss these tables for each reach. The appendix contains plots of observed values and fitted time trends for every fish category referred to below. Remember that the fitted values represent fish sampled on July 1 of the given year and with mean log lipid content as observed in the samples used to build the model. The values of mean log percent lipid are shown in Table 22. Thus, the fitted trend lines may differ from a best visual fit that does not account for lipids or season. This apparent lack of correspondence occurs in several plots.

Table 22 Mean Log₁₀ Percent Lipid in Fish Tissue

| Reach | Species | Type | Mean Log Percent Lipid |
|------------------------------------|----------------|----------------|------------------------|
| <i>Little Lake Butte des Morts</i> | Carp | skin-on fillet | 0.68 |
| | | whole body | 0.90 |
| | Northern Pike | skin-on fillet | 0.00 |
| | Walleye | skin-on fillet | 0.11 |
| | | whole body | 0.73 |
| | Yellow Perch | skin-on fillet | -0.01 |
| <i>Appleton</i> | Walleye | skin-on fillet | -0.03 |
| <i>De Pere</i> | Carp | whole body | 0.88 |
| | Gizzard Shad | whole body | 0.82 |
| | Northern Pike | skin-on fillet | 0.07 |
| | | Walleye | skin-on fillet |
| | | whole body | 0.97 |
| | White Bass | skin-on fillet | 0.60 |
| White Sucker | skin-on fillet | 0.23 | |
| <i>Green Bay Zone 2</i> | Alewife | whole body | 0.97 |
| | Carp | skin-on fillet | 0.82 |
| | | whole body | 0.98 |
| | Gizzard Shad | whole body | 0.77 |
| Yellow Perch | skin-on fillet | -0.29 | |

Reach 1 — Little Lake Butte des Morts

Carp, Skin-on Fillet

After the breakpoint in 1979, PCB concentration declines at a rate of 6 percent per year ($p = 0.02$, Table 14) down to about 1,400 ppb by 1999. Projecting the same rate of decline out to the year 2020 gives an estimated mean PCB concentration of 370 ppb (Table 21), but with a very wide 95 percent confidence interval. Note in particular that the 2,400 ppb upper-bound on the confidence interval for the concentration in 2020 is higher than the estimated concentration 21 years earlier in 1999. Sensitivity analysis (Table 20) shows that a later breakpoint, at 1985, agrees with the data and gives a lower estimate of the post-breakpoint rate of decline, namely, 1.6 percent per year.

The significant negative slope from the best model (Table 14) and the negative slopes from both the earliest and latest breakpoints in the sensitivity analysis (Table 20) consistently suggest that PCBs are decreasing in this species/type in Little Lake Butte des Morts.

Carp, Whole Body

After the breakpoint at 1987, PCB concentration stays almost constant at a level of about 2,500 ppb (0.7% per year, $p = 0.9$). Figure 39 identifies two rather low values in 1987 and 1990. These values do not warrant rejection from the analysis, and the slope calculated with them is appropriate. As a learning exercise, on the other hand, one can illustrate the strong influence of individual observations by omitting these values. A calculation of slope without these two samples would show a continuing decline in PCB concentration to less than 1,000 ppb by 1999.

The barely positive and non-significant slope from the best model versus the negative and barely negative slopes from the earliest and latest breakpoint models, respectively, show no clear evidence of a slope differing from zero for carp whole body samples in Little Lake Butte des Morts.

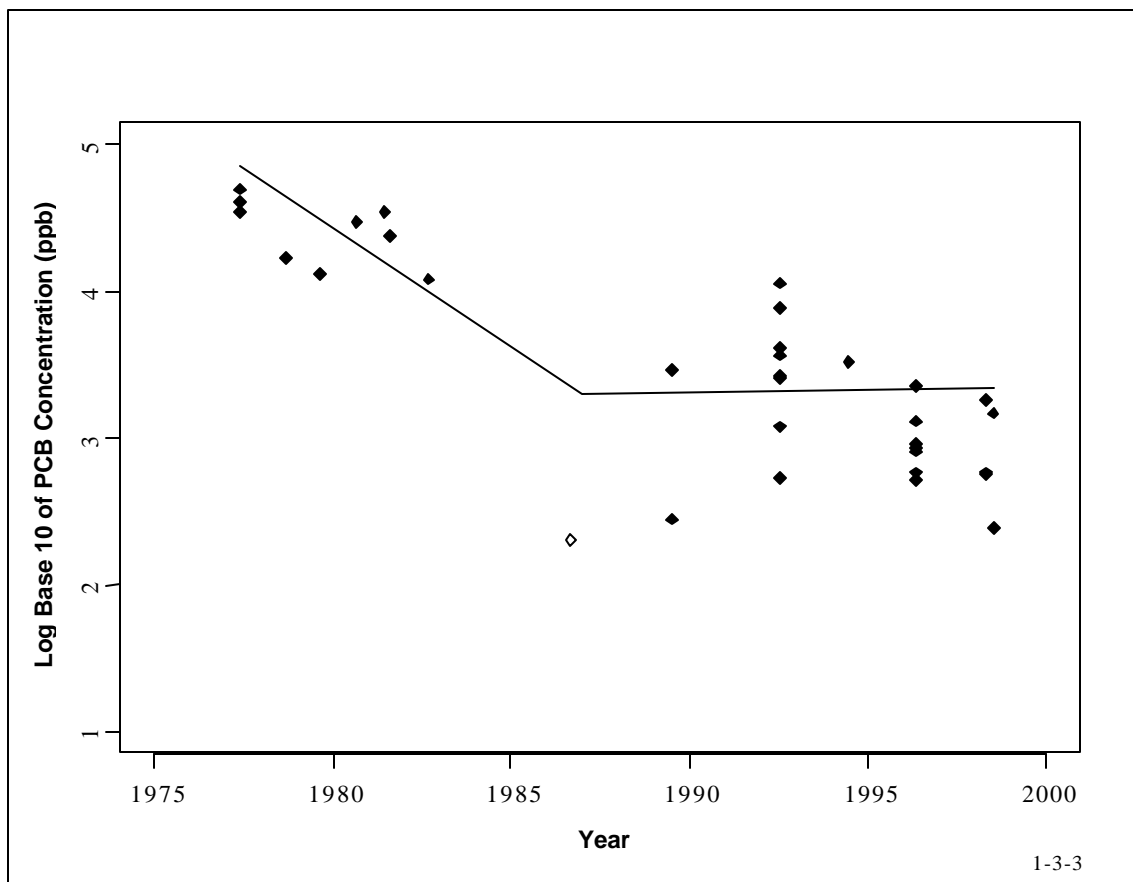


Figure 39 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Carp, Whole Body, versus Time

Breakpoint = 1987 ($p = 0.03$), Final Slope (\log_{10} PCB versus time) = 0.003 ($p = 0.09$), Rate of Change of PCB Concentration During Period of Final Slope = 0.7% (95% confidence interval: -12.3% to 15.6%). Any values below detection limit are depicted by \diamond .

Northern Pike, Skin-on Fillet

The best-fitting model has no breakpoint, but rather a constant rate of decline of 12 percent per year ($p = 0.0003$) yielding a concentration of about 150 ppb by 1999, with a projected mean in the year 2020 of 10 ppb. This is a case of a clear decline during the observation period.

Walleye, Skin-on Fillet

After the breakpoint in 1990, we view a barely increasing PCB concentration hovering around 250 ppb (3.4% per year, $p = 0.6$). The sensitivity analysis (Table 20) shows that a model with an earlier breakpoint, in 1979, also suits the data, producing a post-breakpoint decline of 8 percent per year, and the late 1994 breakpoint produces an increase of 9 percent per year. There is no strong evidence of a slope differing from zero.

Walleye, Whole Fish

The best-fitting model shows a decline in PCB concentration to about 100 ppb in 1987, then a sharp increase at 21 percent per year up to a level of 1,300 ppb by 1999. These parameter estimates are rather imprecise since this model relied upon only 18 samples. The estimated final slope of a 21 percent increase per year is not significantly different from zero, and its confidence interval is very wide: – 4 to 53 percent.

Yellow Perch, Skin-on Fillet

PCB concentration declines sharply until 1981 at 43 percent per year and stays fairly constant thereafter at a level of about 250 ppb (+0.7% per year, $p = 0.8$). There is no evidence of a decreasing late trend.

Summary of Results for Reach 1 — Little Lake Butte des Morts

For most of the fish categories in this reach, we observe an early rapid decline followed by either a slower decline or a flattening without further decline. We find strong evidence against the rate of decline being constant over the whole time range.

On Figure 38, we notice narrow confidence intervals for three fish categories (carp, skin-on fillet; northern pike, skin-on fillet; yellow perch, skin-on fillet). The confidence intervals are much wider for the other three categories, which indicates that the data from these categories do not provide sufficient information to accurately estimate the final slope. The meta-analysis that combines all six results assigns almost all the weight to the three with narrow confidence intervals. Two of these show a negative final slope while one shows a final slope of virtually zero. The combined analysis gives an estimated post-breakpoint rate of decline of 4.9 percent per year—significantly different from zero ($p = 0.0006$). This combined analysis leads us to conclude that PCB concentrations were declining, on the average, at a slow rate during the data collection period. During future periods, species with lower rates of decline would gradually dominate the average rate of decline across species. As noted earlier, the combined rate of change cannot be used for forward projection.

Reach 2 — Appleton to Little Rapids

Walleye, Skin-on Fillet

PCB concentration declines at a constant rate of 10 percent per year over the whole time period ($p = 0.003$), down to an estimated mean of 380 ppb in 1999 and a projected mean of 40 ppb by the year 2020. The sensitivity analysis also shows a negative slope for both the earliest (1982) and latest (1994) breakpoints.

Reach 4 — De Pere to Green Bay

Carp, Whole Fish

This model shows decline in PCB concentration to a minimum of about 3,200 ppb in 1995 (the breakpoint), followed by a sharp increase of 22 percent per year ($p = 0.03$) up to a mean of 7,500 ppb by 1999. We find a rather wide confidence interval for this rate of increase, but it does not quite include zero. The sensitivity analysis, on the other hand, shows that the data are also consistent with an earlier breakpoint in 1990, followed by a slightly negative slope, close to zero. Thus, despite the p -value of 0.03 for the post-breakpoint negative slope, when we add in the uncertainty due to the breakpoint, the final slope is not convincingly different than zero.

Gizzard Shad, Whole Fish

PCB concentration declines at a constant rate of 5 percent per year ($p = 0.0002$) to a mean of 1,700 ppb in 1999 and a projected mean of 570 ppb in 2020.

Northern Pike, Skin-on Fillet

PCB concentration declines at a constant rate of 10 percent per year ($p < 0.0001$) to a mean of 540 ppb in 1999 and projected mean of 60 ppb in 2020.

Walleye, Skin-on Fillet

PCB concentration declines at a constant rate of 7 percent per year ($p < 0.0001$) to a mean of 780 ppb in 1999 and projected mean of 160 ppb in 2020. The spread of observations (more than 20 years) in this analysis, and in the preceding analysis for northern pike, helps to considerably improve the precision of the combined slope estimates for this reach (see below).

Walleye, Whole Fish

PCB concentration declines at a constant rate of 8 percent per year ($p < 0.0001$) to a mean of 4,300 ppb in 1999 and projected mean of 740 ppb in 2020.

White Bass, Skin-on Fillet

PCB concentration declines at a constant rate of 5 percent per year ($p = 0.002$), to a mean of 2,700 ppb in 1999 and projected mean of 980 ppb in 2020. Sensitivity analysis shows the data are consistent with a late breakpoint at 1996 followed by a slope that slightly increases.

White Sucker, Skin-on Fillet

PCB concentration declines at a constant rate of 8 percent per year ($p < 0.0001$), to a mean of 640 ppb in 1999 and projected mean of 110 ppb in 2020.

Summary of Results for Reach 4 — De Pere to Green Bay

All but one of the fish categories show a decline in PCB concentration at a constant rate. The meta-analysis results reflect this, with an estimated rate of decline of 7 percent per year, highly significantly different from zero

($p < 0.0001$). Whole body carp, with a breakpoint in 1995, emerges as the only exception to this pattern of monotonically decreasing PCB concentration, occurring in six out of seven of the analyses. These slopes have relatively tight confidence intervals. One can explain the large increase after 1995 in carp due to high PCB concentrations observed in a large number of fish sampled on July 2 and July 6, 1998. Such a phenomenon might reflect a scouring event that exposed buried sediment with a high PCB concentration. Or the large positive slope for the carp may be random, given that the sensitivity analysis accords with a slightly negative to a large positive slope for this reach/species/type combination, as discussed earlier.

Reach 5 — Green Bay Zone 2

Alewife, Whole Body

PCB concentration declined at a constant rate of 4 percent per year ($p = 0.05$) to a mean of 2,100 ppb in 1999 and a projected mean of 900 ppb in 2020.

Carp, Skin-on Fillet

PCB concentration declines at a constant rate of 5 percent per year ($p = 0.16$, not significantly different from zero) to a mean of 4,900 ppb in 1999 and projected mean of 1,630 ppb in 2020.

Carp, Whole Fish

PCB concentration increases to a maximum of about 25,000 ppb in 1983, and then declines at a rate of 16 percent per year ($p < 0.0001$) down to a mean of 1,500 ppb in 1999 and projected mean of 40 ppb in 2020. An informal sensitivity analysis does not alter the combination of an initially positive and final negative slope. However, we are concerned about having potentially over-fit the model. We have only 5 years during which data were collected over a period covering about 20 years. Given that five parameters in the model relate to time (breakpoint, final slope, slope difference [early minus late], and two season parameters), it is possible to fit a spline model “too well” to the limited number of years with observations. In any case, the final slope does appear firmly negative, though it may be less negative than the 16 percent. A model fitted without a breakpoint yields a single negative slope with a rate of decline of 9 percent per year.

Gizzard Shad, Whole Fish

Samples were only taken over a relatively short time period from 1989 to 1999. PCB concentration appears to increase over this time period at a rate of 6 percent per year ($p = 0.01$) to a mean of 3,200 in 1999.

Yellow Perch, Skin-on Fillet

PCB concentration declines at a constant rate of 11 percent per year ($p = 0.004$) to a mean of 150 ppb in 1999 and projected mean of 14 ppb in 2020.

We have rejected a model with a breakpoint at 1986, even though the breakpoint is, formally, highly significant ($p = 0.0008$). The breakpoint model yields a final rate of change of plus 15 percent per year and a pre-break rate of minus 69 percent per year. We regard this implausible combination as due to over-fitting (mentioned earlier) and accept, instead, the single-slope model noted in the figure and table.

Summary of Results for Reach 5 — Green Bay Zone 2

Four out of the five fish categories for this reach show a continuing decline in PCB concentration. The meta-analysis results reflect this, yielding a combined estimate of final rate of decline of 5 percent per year ($p < 0.0001$).

5.2.3 Additional Analysis of Alternative Models

Results for Fitting Models with Breakpoint at 1985

In addition to showing results for the best-fitting model, we fit models to the 19 fish categories using a single common breakpoint. The best year for this breakpoint is 1985. A breakpoint at 1985 fits nearly as well as the optimal breakpoint for almost all fish categories. Table 23 shows results of fitting this model to every fish category.

Testing for a Non-constant Final Slope

Projection of PCB concentrations presumes some kind of steady or predictable state. In this section, we consider the “steadiness” of time trends. In order to test the assumption of a constant linear slope in the time period after the breakpoint, we fit models including a quadratic term for that time period. Table 24 shows the results of these analyses for the best-fitting model.

Table 23 Details of Fitting Models with a Breakpoint at 1985 for Every Fish Category

| Species | Type | Model | Break-point Year | n | Intercept | SE Int | Estimate of Final (post-1985) Slope | | | Early Slope Difference | | | Coefficient of Log of Percent Lipids | | | Peak of Seasonal Variation | | |
|------------------------------------|----------------|-------|------------------|-----|-----------|--------|-------------------------------------|-------|---------|------------------------|-------|---------|--------------------------------------|------|---------|----------------------------|------|---------|
| | | | | | | | Slope | SE | p-value | Difference | SE | p-value | Log ₁₀ | SE | p-value | Mo. | Amp. | p-value |
| <i>Little Lake Butte des Morts</i> | | | | | | | | | | | | | | | | | | |
| Carp | skin-on fillet | 2 | 1985 | 55 | 3.23 | 0.12 | -0.007 | 0.021 | 0.7419 | -0.090 | 0.043 | 0.0403 | 0.86 | 0.16 | 0.0000 | 12.9 | 0.59 | 0.0268 |
| | whole body | 2 | 1985 | 40 | 3.41 | 0.16 | -0.018 | 0.028 | 0.5264 | -0.158 | 0.072 | 0.0360 | 0.87 | 0.34 | 0.0148 | 7.0 | 0.69 | 0.0099 |
| Northern Pike | skin-on fillet | 2 | 1985 | 19 | 2.84 | 0.19 | -0.079 | 0.024 | 0.0053 | 0.071 | 0.061 | 0.2663 | 0.57 | 0.31 | 0.0854 | 1.8 | 0.56 | 0.0829 |
| Walleye | skin-on fillet | 2 | 1985 | 63 | 2.46 | 0.09 | -0.026 | 0.012 | 0.0379 | -0.061 | 0.032 | 0.0570 | 0.43 | 0.14 | 0.0038 | 12.7 | 0.25 | 0.1026 |
| | whole body | 2 | 1985 | 18 | 2.22 | 0.39 | 0.074 | 0.045 | 0.1285 | -0.310 | 0.103 | 0.0106 | 0.97 | 0.36 | 0.0206 | 11.9 | 0.66 | 0.0077 |
| Yellow Perch | skin-on fillet | 2 | 1985 | 34 | 2.10 | 0.10 | 0.018 | 0.019 | 0.3297 | -0.133 | 0.049 | 0.0110 | 0.34 | 0.21 | 0.1144 | 10.7 | 0.11 | 0.0025 |
| <i>Appleton</i> | | | | | | | | | | | | | | | | | | |
| Walleye | skin-on fillet | 2 | 1985 | 30 | 3.20 | 0.20 | -0.065 | 0.022 | 0.0059 | 0.103 | 0.089 | 0.2574 | 1.23 | 0.20 | 0.0000 | 7.4 | 0.56 | 0.0005 |
| <i>De Pere</i> | | | | | | | | | | | | | | | | | | |
| Carp | whole body | 2 | 1985 | 90 | 3.94 | 0.09 | -0.025 | 0.011 | 0.0238 | -0.031 | 0.033 | 0.3508 | 0.82 | 0.12 | 0.0000 | 6.9 | 0.15 | 0.0304 |
| Northern Pike | skin-on fillet | 2 | 1985 | 40 | 3.13 | 0.11 | -0.039 | 0.010 | 0.0005 | -0.020 | 0.024 | 0.4111 | 0.71 | 0.17 | 0.0002 | 9.0 | 0.13 | 0.2505 |
| Walleye | skin-on fillet | 2 | 1985 | 120 | 3.21 | 0.05 | -0.035 | 0.005 | 0.0000 | 0.011 | 0.018 | 0.5282 | 0.86 | 0.07 | 0.0000 | 8.7 | 0.02 | 0.6196 |
| | whole body | 2 | 1985 | 58 | 4.00 | 0.07 | -0.039 | 0.009 | 0.0000 | 0.009 | 0.028 | 0.7440 | 0.45 | 0.12 | 0.0007 | 7.0 | 0.12 | 0.1931 |
| White Bass | skin-on fillet | 2 | 1985 | 58 | 3.61 | 0.07 | -0.019 | 0.007 | 0.0065 | -0.117 | 0.109 | 0.2897 | 0.83 | 0.11 | 0.0000 | 6.8 | 0.32 | 0.0592 |
| White Sucker | skin-on fillet | 2 | 1985 | 44 | 3.12 | 0.08 | -0.032 | 0.010 | 0.0020 | -0.013 | 0.025 | 0.6010 | 0.43 | 0.15 | 0.0065 | 7.3 | 0.08 | 0.4813 |
| <i>Green Bay Zone 2</i> | | | | | | | | | | | | | | | | | | |
| Alewife | whole body | 2 | 1985 | 44 | 3.42 | 0.06 | -0.002 | 0.011 | 0.8200 | -0.087 | 0.040 | 0.0341 | 0.90 | 0.13 | 0.0000 | 5.4 | 0.09 | 0.0034 |
| Carp | skin-on fillet | 2 | 1985 | 28 | 3.84 | 0.08 | -0.063 | 0.026 | 0.0226 | 0.105 | 0.055 | 0.0698 | 0.74 | 0.14 | 0.0000 | 3.0 | 0.41 | 0.0052 |
| | whole body | 2 | 1985 | 57 | 3.89 | 0.06 | -0.075 | 0.013 | 0.0000 | 0.135 | 0.040 | 0.0013 | 0.87 | 0.10 | 0.0000 | 6.6 | 0.23 | 0.0013 |
| Yellow Perch | skin-on fillet | 2 | 1985 | 19 | 2.60 | 0.35 | 0.015 | 0.018 | 0.4061 | -0.745 | 0.170 | 0.0007 | 1.54 | 0.35 | 0.0008 | 7.2 | 2.99 | 0.0008 |

Table 24 Test for Curvature in Final Slopes

| Species | Type | Coefficient of <i>t</i> -squared | SE of <i>t</i> -squared Coefficient | Tests for Curvature | | |
|------------------------------------|----------------|----------------------------------|-------------------------------------|--|--|---|
| | | | | <i>p</i> -value ⁺ (2-sided) | <i>p</i> -value ⁺ (1-sided, plus) | <i>p</i> -value ⁺ (1-sided, minus) |
| <i>Little Lake Butte des Morts</i> | | | | | | |
| Carp | skin-on fillet | -0.0014 | 0.0024 | 0.56 | 0.718 | 0.564 |
| | whole body | -0.0144 | 0.0067 | 0.04* | 0.981 | 0.039* |
| Northern Pike | skin-on fillet | -0.0033 | 0.0024 | 0.19 | 0.905 | 0.190 |
| Walleye | skin-on fillet | -0.0095 | 0.0094 | 0.32 | 0.842 | 0.317 |
| | whole body | -0.0202 | 0.0101 | 0.07 | 0.965 | 0.070 |
| Yellow Perch | skin-on fillet | -0.0021 | 0.0059 | 0.72 | 0.639 | 0.722 |
| <i>Appleton to Little Rapids</i> | | | | | | |
| Walleye | skin-on fillet | -0.0047 | 0.0041 | 0.26 | 0.872 | 0.255 |
| <i>De Pere to Green Bay</i> | | | | | | |
| Carp | whole body | 0.0168 | 0.0362 | 0.64 | 0.644 | 0.678 |
| Gizzard Shad | whole body | 0.0032 | 0.0029 | 0.29 | 0.290 | 0.855 |
| Northern Pike | skin-on fillet | 0.0009 | 0.0008 | 0.25 | 0.249 | 0.876 |
| Walleye | skin-on fillet | -0.0005 | 0.0006 | 0.42 | 0.791 | 0.418 |
| | whole body | 0.0000 | 0.0008 | 0.97 | 0.514 | 0.971 |
| White Bass | skin-on fillet | 0.0015 | 0.0018 | 0.41 | 0.410 | 0.795 |
| White Sucker | skin-on fillet | 0.0011 | 0.0010 | 0.30 | 0.300 | 0.850 |
| <i>Green Bay Zone 2</i> | | | | | | |
| Alewife | whole body | 0.0019 | 0.0011 | 0.10 | 0.099 | 0.950 |
| Carp | skin-on fillet | -0.0061 | 0.0035 | 0.10 | 0.952 | 0.096 |
| | whole body | 0.0034 | 0.0018 | 0.06 | 0.062 | 0.969 |
| Gizzard Shad | whole body | -0.0007 | 0.0032 | 0.82 | 0.591 | 0.818 |
| Yellow Perch | skin-on fillet | 0.0126 | 0.0034 | 0.003** | 0.003** | 0.999 |
| All | | | | 0.008** | 0.4 | 0.2 |

Notes:

⁺ The three *p*-values indicate the statistical significance of the *t*-squared (curvature) term in the regression model for time trends. In all three columns, the null hypothesis is no curvature (i.e., there is a straight-line constant slope after the breakpoint—or the whole period, if there is no breakpoint). In the first *p*-value column, the alternative hypothesis is that the final time period has some curvature (i.e., the slope is shifting **either** toward more positive **or** more negative values). In the second *p*-value column, the alternative hypothesis is that the slope is shifting toward more positive values (less decline in PCB concentrations). In the third column, the alternative hypothesis is that the slope is shifting toward more negative values (greater decline in PCB concentrations).

* $p < 0.05$

** $p < 0.01$

*** $p < 0.001$

This model introduces a time-squared term for the final period. It is an implausible model for projection of PCB concentration, but readily works to detect a non-constant rate of decline of PCBs during the final period. We refer to this as “curvature.” A positive sign for the time-squared term indicates a shifting slope over time toward either less reduction in PCBs or more accrual of PCBs. A negative sign indicates a shift toward more reduction or less accrual.

The results (Table 24) show two categories with significant curvature, discussed below. Overall, curvature may be a general phenomenon. A meta-analysis using

chi-squared calculated from the 19 p -values for curvature yields $X^2 = 61.3$, with 38 degrees of freedom and $p = 0.008$. Thus, we reject the null hypothesis that **all** of the final periods, after the breakpoints (including the entire period, if there is no breakpoint), have a simple linear trend (on the log scale). We note, also, that 6 out of the 19 p -values for curvature are less than 0.10, whereas only 2 would be expected by chance. This excess of small p -values suggests that “curvature,” or changing slopes over time, is common and not a feature confined to one or two of the categories analyzed here. Further, it appears that the curvature is a mixture of positive and negative changes (i.e., there are slopes that may shift toward either more negative or more positive rates of change as time passes). The evidence for a mixture of positive and negative changes is two-fold. First, there is both a positive and a negative curvature result among the two fish categories with $p < 0.05$ on the curvature test. In Green Bay Zone 2, yellow perch samples of skin-on fillet evidence that their rate of decline is decreasing (toward less reduction of PCBs) with $p = 0.002$, and in Little Lake Butte des Morts carp whole body samples evidence that their recent barely positive slope is changing toward an either flat or negative trend with more reduction of PCBs ($p = 0.04$). Among the six fish categories with $p < 0.10$ for curvature (marginally significant results) we again find quite an even mixture of positive and negative curvature—three of each. Overall, 9 categories with fitted curvature with a positive coefficient (rates of decline shifting toward slower reduction of PCBs over time) and 10 have negative curvature (rates of decline shifting toward faster reduction of PCBs over time).

There is a second reason we feel that slopes are shifting both positively and negatively. A meta-analysis using a one-sided test to detect an excess of fish categories with positive curvature (toward less reduction of PCBs) yields $p = 0.4$, and the p -value for an excess of negative curvature (toward more reduction of PCBs) yields $p = 0.2$. These two p -values indicate no significant excess of either only positively or only negatively curving slopes, but there is a significant excess of curving slopes in general (either positive or negative). Thus, we find evidence for changing slopes ($p = 0.008$, noted above), but of mixed direction among the fish categories. We can only be confident that there is change.

The generally non-significant p -values in the two-sided p -value column of Table 24, and in other p -value columns inspire confidence of curvature in very few cases. Except for the four p -values noted with asterisks (not including “All”), it is difficult to ascribe curvature to the specific combinations of species, reach, and sample type. However, the excess of relatively small two-sided p -values overall (even if individual p -values are not significant) does allow us to conclude with some confidence that there is changeability in the final slopes ($p = 0.008$). That is, we reject the notion that, during the period of final slopes, rates of changes were utterly constant for every combination of reach, species, and sample type. We accept the alternative that rates of change were shifting over time, both in a negative and positive direction for at least some combinations.

5.3 Conclusions about Trends over Time in PCB Concentration in Fish

The meta-analyses within three reaches with more than one fish category available for analysis show that PCB concentration was declining at a rate of 5 to 7 percent per year (Table 19, Little Lake Butte des Morts, De Pere, Green Bay Zone 2). The single fish category that could be analyzed for the Appleton Reach also shows a decline of 10 percent per year. Reach 1, Little Lake Butte des Morts, calls attention to a steeper decline in earlier years. All analyses with a breakpoint in this reach show a steeper decline before than after the breakpoint. But in the other reaches, except for 2 out of 13 categories, the data for each fish category considered individually are consistent with a constant rate of decline over the whole time period.

The majority of fish categories have data consistent with only a simple linear trend, and the balance of categories (with breakpoints) have post-break data fit well by a linear trend. Nevertheless, the collective evidence is that slopes (on the log scale) tend to be non-constant, as evidenced by the rejection of the hypothesis of no curvature in the final slopes based on the meta-analysis (Table 24).

We cannot project into the future with precision for several reasons. Many species suffer from rather sparse data with observations occurring at only a few time points. Models based on these data do not provide highly precise estimates. Incorporating the extra uncertainty due to estimating the breakpoint presents a challenge. We have done so in an informal fashion using a sensitivity analysis. The uncertainty in future projections would be greater if the uncertainty in the breakpoint were formally incorporated into calculations. Finally, some of the unusual changes in slope from before to after a breakpoint may be genuine, due to unpredictable events such as floods accompanied by scouring and deposition. If so, such events will continue adding variability to PCB concentration over time, making predictions based on the assumption of a future decline at a constant rate questionable. The presence of curvature (non-constant slopes) is consistent with the more dramatic changes represented by breakpoints and suggests a dynamic process, liable to change, rather than a steady state with constant rates of change.

5.4 Comparison of De Pere Reach to Green Bay Zone 2

We compared species and sample types between the De Pere to Green Bay Reach (equivalent to “Green Bay Zone 1” and so labeled in some reports) and Green Bay Zone 2. The two sets of observations from the two bodies of water are usually significantly different; either in the mean PCB concentration, the time trend of PCB concentration, or in the relationship of PCB concentration to lipid content of tissue.

We were able to carry out these analyses for some additional species and sample type combinations for which time trends could not be calculated by using a snapshot during a single year or short span of years. Table 25 shows which comparisons could be made. We carried out five analyses comparing De Pere Reach and Green Bay Zone 2 during a short “snapshot” cross-sectional period of years, and there were three analyses where time trends could be compared between reaches. In order to have a consistent period of years for the time trend comparison and to avoid differences between reaches arising from different sampling patterns over time, we limited the time trend analyses to a common period of years, 1989 through 1998.

We note that we limited our analysis and discussion to the data provided to us. A discussion of the biological and physical comparisons between the two bodies of water can be found in Technical Memorandum 7c (WDNR, 2001), the Remedial Investigation, and the Baseline Risk Assessment for the Lower Fox River and Green Bay (ThermoRetec, 2001a; ThermoRetec, 2001b).

Table 25 De Pere Reach and Green Bay Zone 2: Fish Types and Sample Types with Sufficient Data for PCB Comparisons

| Sample Type | Species | Type of Analysis | |
|---------------------------------|--------------|---|----------------------------------|
| | | Single Time Snapshot PCB Comparison - Years | Time Trend Analysis Across Years |
| Whole Fish/Whole Body/Composite | alewife | 1989 | 1989–1998 |
| Whole Fish/Whole Body/Composite | carp | 1989 | 1989–1998 |
| Whole Fish/Whole Body/Composite | gizzard shad | 1989 | 1989–1998 |
| Skin On Fillet | walleye | 1989–1991 | |
| Whole Fish/Whole Body/Composite | walleye | 1989 | |

The equation used to analyze the De Pere and Zone 2 reaches based on the snapshot data is:

Equation 18

$$\log_{10}(PCB) = b_0 + b_1L + b_2R + b_3L \cdot R + e$$

where

- PCB* = PCB concentration in units of ppb,
- L* = log₁₀(percent lipid content),
- R* = dichotomous indicator of Zone 2 versus De Pere Reach, and
- e* = random error.

For the comparison of time trends in the De Pere and Zone 2 reaches, the equation is extended to:

Equation 19

$$\log_{10}(PCB) = b_0 + b_1L + b_2R + b_3L \cdot R + b_4t + b_5t \cdot R + e$$

where

t = time in years since January 1, 1989.

In both the snapshot and time trends equations, all coefficients of terms involving R (reach) should be zero or close to zero if a given fish species takes in PCBs at a similar level and processes PCBs in a similar way in the two reaches. For example, in the snapshot equation if b_3 (the coefficient of $L \cdot R$) is zero, the increase in PCB concentration for a specified increase in fat content is the same in the two reaches. In addition, if b_2 (the coefficient of R) is also zero, then the mean PCB concentration is the same in the two reaches, given equal lipid content. As another example, if b_5 (coefficient of $t \cdot R$) is zero in the time trends model, then the rate of change of $\log_{10}(PCB)$ is the same in the two reaches. Thus, comparing the two reaches involves testing whether certain coefficients in regression models are significantly different from zero.

We detected one outlier, which was removed from the De Pere Reach versus Green Bay Zone 2 analysis. The outlier is noted in Table 27.

Table 26 Outlier from Analysis of De Pere Reach versus Green Bay Zone 2

| Database ID | Reach | Fish Type | Sample Type | Total PCBs |
|--------------|------------------|-----------|-------------|------------|
| WDF209006BC1 | Green Bay Zone 2 | alewife | whole body | 19,000 |

Reason:

Large outlier. Other PCB values range from 990 to 4,500.

5.4.1 De Pere Reach versus Green Bay Zone 2: “Snapshot” Analysis

Four out of five snapshot analyses (Figure 40 through Figure 44) showed statistically significant differences between the two reaches (Table 27). In two of the analyses, PCB concentrations varied with percent lipid in a different way in the two reaches, and in two analyses the mean log PCB concentration differed between the two reaches, controlling for lipid content.

The two species with different PCB-lipid relationships were carp and gizzard shad, both whole body samples. For carp (whole body) the coefficient of the log lipid term L , in the snapshot equation above, when, combined with the coefficient of $L \cdot R$, yields different rates of change of log PCB with changes in log lipid content ($p = 0.02$). The slope of log PCB versus log lipid is 0.68 and 1.01 in De Pere and Green Bay Zone 2 reaches, respectively. (In all De Pere versus Zone 2

analyses, reach was coded as “1” for De Pere to Green Bay and “2” for Green Bay Zone 2. Thus, based on the snapshot equation, the slope of log PCB versus log lipid in the De Pere Reach, coded as “1,” is $0.3426 + 0.3346 \times 1 = 0.6772$, and, in Green Bay Zone 2, coded as “2,” the slope is $0.3426 + 0.3346 \times 2 = 1.0118$.)

Table 27 Fitted Models for Log₁₀ (PCB Concentration) versus Log₁₀ (Percent Lipid) in De Pere Reach and Green Bay Zone 2 for Species with Sufficient Data During 1989

| Sample Type | Species | Single Time Snapshot PCB Comp: Years | De Pere Reach | | Green Bay Zone 2 | | Equal Slopes Likelihood Ratio <i>p</i> -value | Equal Intercepts Likelihood Ratio <i>p</i> -value |
|---|---------------------------|--------------------------------------|---------------|--------|------------------|-------|---|---|
| | | | Intercept | Slope | Intercept | Slope | | |
| Whole Fish/ Whole Body/ Composite | alewife ⁺ | 1989 | 2.943 | 0.663 | 2.668 | 0.663 | 0.32 | 0.00006*** |
| Whole Fish/ Whole Body/ Composite | carp ⁺ | 1989 | 3.092 | 0.677 | 2.675 | 1.012 | 0.016* | |
| Whole Fish/ Whole Body/ Composite | gizzard shad ⁺ | 1989 | 3.559 | -0.204 | 2.846 | 0.496 | 0.0009*** | |
| Skin On Fillet | walleye | 1989-1991 | 3.040 | 0.348 | 3.040 | 0.348 | 0.69 | 0.66 |
| Whole Fish/ Whole Body/ Composite | walleye ⁺ | 1989 | 3.390 | 0.501 | 3.269 | 0.501 | 0.17 | 0.0058** |

Notes:

- + Fish types significantly different between reaches at 5 percent significance level or better.
- * $p < 0.05$
- ** $p < 0.01$
- *** $p < 0.001$

To illustrate the implication of these coefficients, consider a doubling of lipid content (e.g., from 5 to 10 percent). It can be derived from Equation 18 that an increase in lipid content by any multiplicative factor F , such as $F = 2$, leads to an increase in PCB concentration by a multiplicative factor of $F^{b_1 + R \cdot b_3}$. Thus, a doubling of lipid content leads to an increase in PCB concentration (ppb, not log) by a factor of $2^{0.6772} = 1.60$ in the De Pere Reach, and in Green Bay Zone 2, by a factor of $2^{1.0118} = 2.02$. The increase in Green Bay Zone 2 is larger by 26 percent.

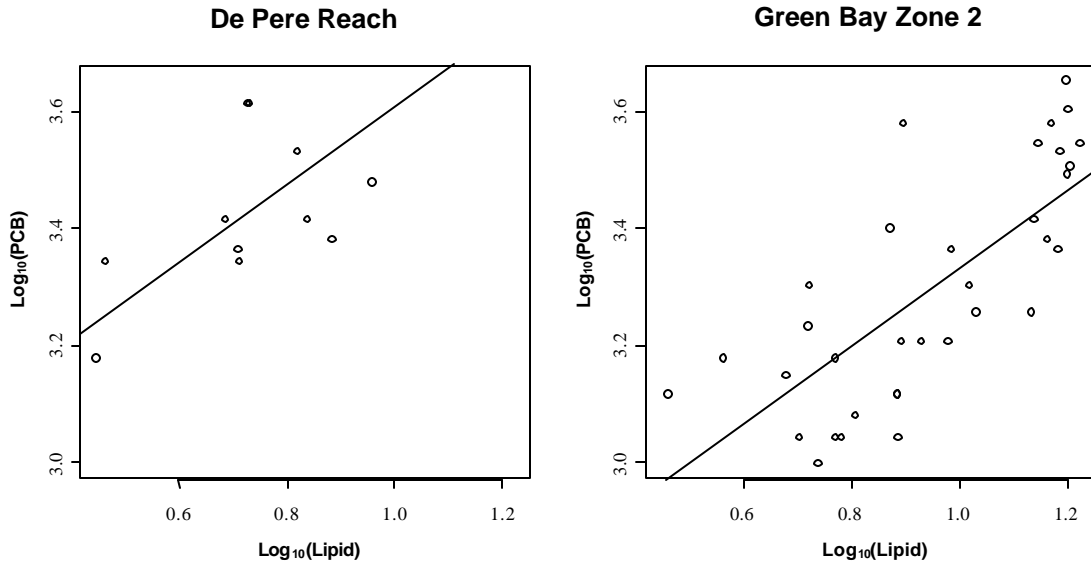


Figure 40 Log PCB Concentration versus Log Percent Lipid for De Pere Reach and Green Bay Zone 2 for Alewife, Whole Body, 1989

For the alewife species, there was no apparent difference in the slope of the relationship between \log_{10} (PCB) and \log_{10} (percent lipid) ($p = 0.3$, likelihood ratio test for slope differences). The intercepts were significantly different ($p = 0.00006$, likelihood ratio test). Thus, the mean PCB concentrations for alewife fish in the two zones are significantly different. Figure 40 shows that alewife in the De Pere Reach tend to have a higher PCB content at all lipid levels.

The carp whole body samples (Table 27, Figure 42) in Green Bay Zone 2 showed a greater rate of increase of PCBs with increasing lipid content than samples from the De Pere Reach. (See the steeper slope in Figure 42, right, than in the left panel.) The difference is statistically significant ($p = 0.02$).

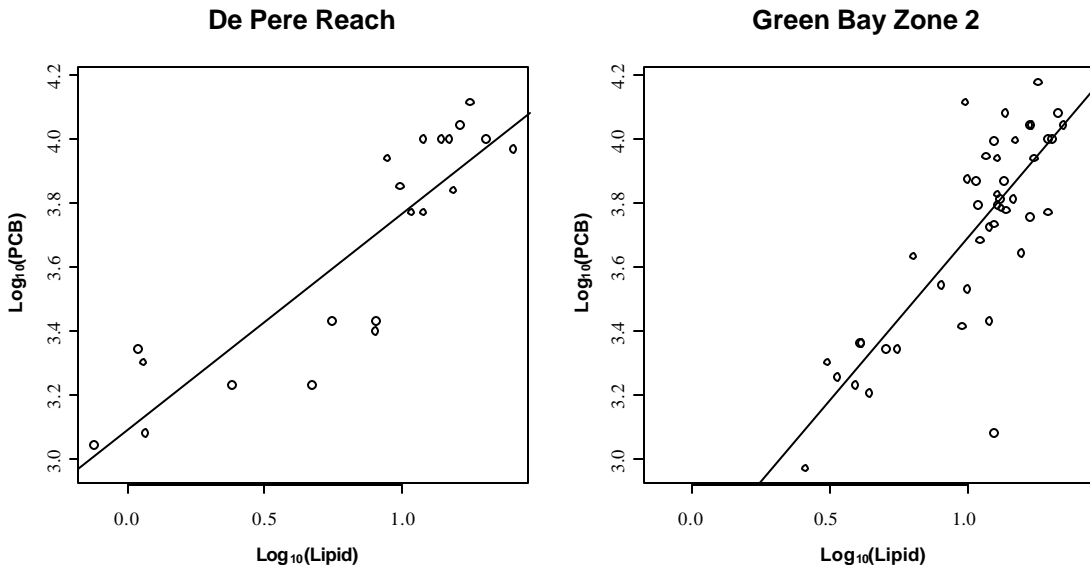


Figure 41 Log PCB Concentration versus Log Percent Lipid for De Pere Reach and Green Bay Zone 2 for Carp, Whole Body, 1989

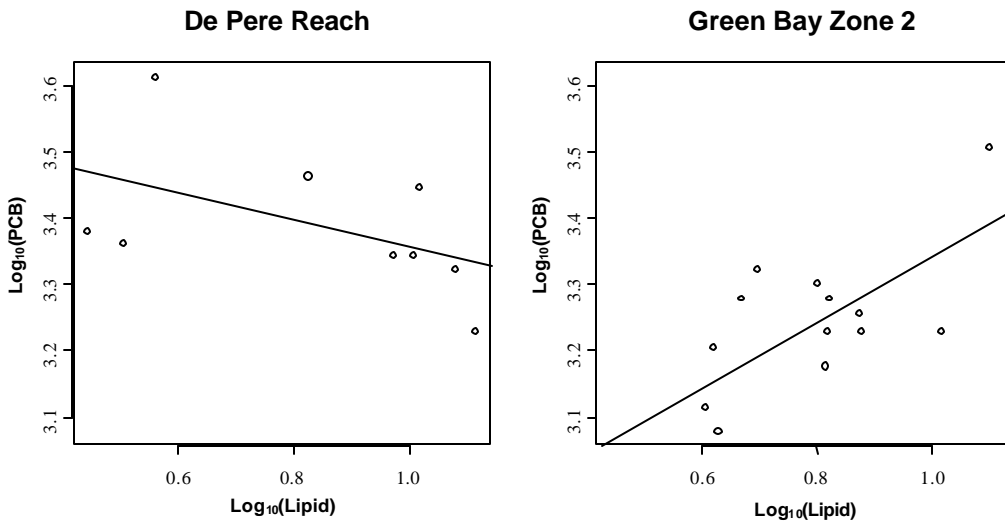


Figure 42 Log PCB Concentration versus Log Percent Lipid for De Pere Reach and Green Bay Zone 2 for Gizzard Shad, Whole Body, 1989

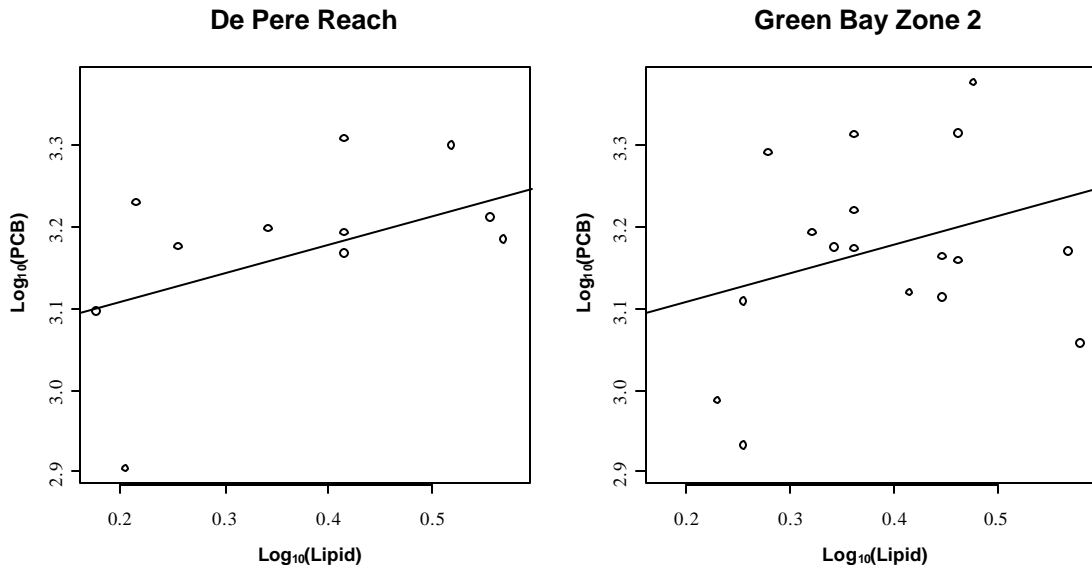


Figure 43 Log PCB Concentration versus Log Percent Lipid for De Pere Reach and Green Bay Zone 2 for Walleye, Skin-on Fillet, 1989–1991

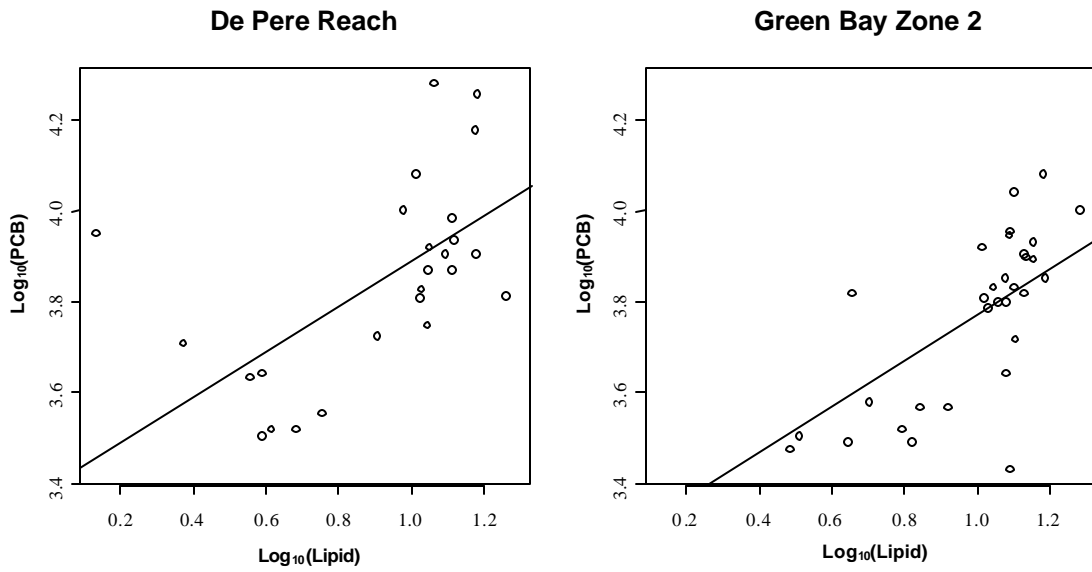


Figure 44 Log PCB Concentration versus Log Percent Lipid for De Pere Reach and Green Bay Zone 2 for Walleye, Whole Body, 1989

In the gizzard shad samples, the slope of log PCB versus log lipid in the De Pere Reach is -0.20 (Table 27, Figure 42). That slope is negative in the De Pere Reach is biologically implausible and probably randomly different from zero or a slightly positive value. The negative slope is significantly different ($p = 0.0009$) from the positive coefficient of 0.50 in Green Bay Zone 2. In Green Bay Zone 2, a doubling of percent lipid in the gizzard shad species would yield an expected 41 percent increase in PCB concentration, while in the De Pere Reach, if one takes the fitted model at face value, the PCB concentration would decrease. If zero or a small positive value is the true slope for log PCB concentration versus log percent lipid in the De Pere Reach, a doubling of lipid content in this reach would cause only a slight change in PCB concentration.

For these two species, carp and gizzard shad, the plots (Figure 41 and Figure 42) indicate that the PCB concentrations differ most at low lipid levels and tend to converge at higher lipid levels. Thus, for each of the two species, the fish samples in the two reaches will have similar PCB concentrations at higher lipid levels and dissimilar PCB concentrations at lower lipid levels.

In two of the three other snapshot analyses (alewife and walleye, both “whole body”), slopes of log PCB versus log lipid were not significantly different between the reaches ($p = 0.3$ and 0.2 , respectively), but the mean PCB concentration differed, controlling for lipid level ($p = 0.0001$ and 0.006 , respectively). The plots (Figure 40 and Figure 44) clearly convey this offset between the PCB-lipid relationship.

The difference between reaches in mean log PCB concentration for a specified lipid content is the coefficient b_2 in the snapshot equation with, in these two analyses, b_3 set equal to zero and the $L \cdot R$ term excluded from the model. The De Pere Reach minus Green Bay Zone 2 difference in expected log PCB is $2.943 - 2.668 = 0.275$ for alewife and 0.121 for walleye. These differences correspond to a geometric mean PCB concentration that is $10^{0.275} = 1.9$ times higher (90 percent higher) for alewife in De Pere Reach than in Green Bay Zone 2, and $10^{0.121} = 1.3$ times higher (30 percent higher), correspondingly, for walleye.

5.4.2 De Pere Reach versus Green Bay Zone 2: Time Trends Analysis

All three analyses comparing alewife, carp, and gizzard shad between De Pere Reach and Green Bay Zone 2 yield statistically significant differences in time trends between the reaches, as shown in Table 28. The trends are also plotted on Figure 45 through Figure 47. All results here are based on analyses of whole body samples. The slopes for alewife (log PCB versus time in years) are -0.023 for the De Pere Reach and 0.004 for Green Bay Zone 2. They imply that the PCB concentration in De Pere Reach alewife has been decreasing by 5 percent per year and increasing by 1 percent per year in Green Bay Zone 2, a difference in rates of 6 percent per year. Similar comparisons for the other species, based on the slopes in the table, yield, for carp, a 14 percent per year greater rate of decrease in

Green Bay Zone 2 than in the De Pere Reach. For gizzard shad, De Pere Reach concentrations have been decreasing 10 percent per year faster than the Green Bay Zone 2 concentrations.

Table 28 Log_{10} (PCB Concentration) versus Time in De Pere Reach and Green Bay Zone 2 for Species with Sufficient Data During 1989–1998

| Sample Type | Species | Time Trend PCB Comp: Years | De Pere Reach | | Green Bay Zone 2 | | Equal Slopes Likelihood Ratio <i>p</i> - value |
|---------------------------------|--------------|-------------------------------------|---------------|---------|------------------|----------|---|
| | | | Intercept | Slope | Intercept | Slope | |
| Whole Fish/Whole Body/Composite | alewife | 1989–1998 | 49.743 | -0.0232 | -4.336 | 0.00382 | 0.045* |
| Whole Fish/Whole Body/Composite | carp | 1989–1998 | -6.218 | 0.005 | 105.89 | -0.05131 | 0.0099** |
| Whole Fish/Whole Body/Composite | gizzard shad | 1989–1998 | 55.954 | -0.0264 | -24.037 | 0.01368 | 0.0031** |

Notes:

- * $p < 0.05$
- ** $p < 0.01$
- *** $p < 0.001$

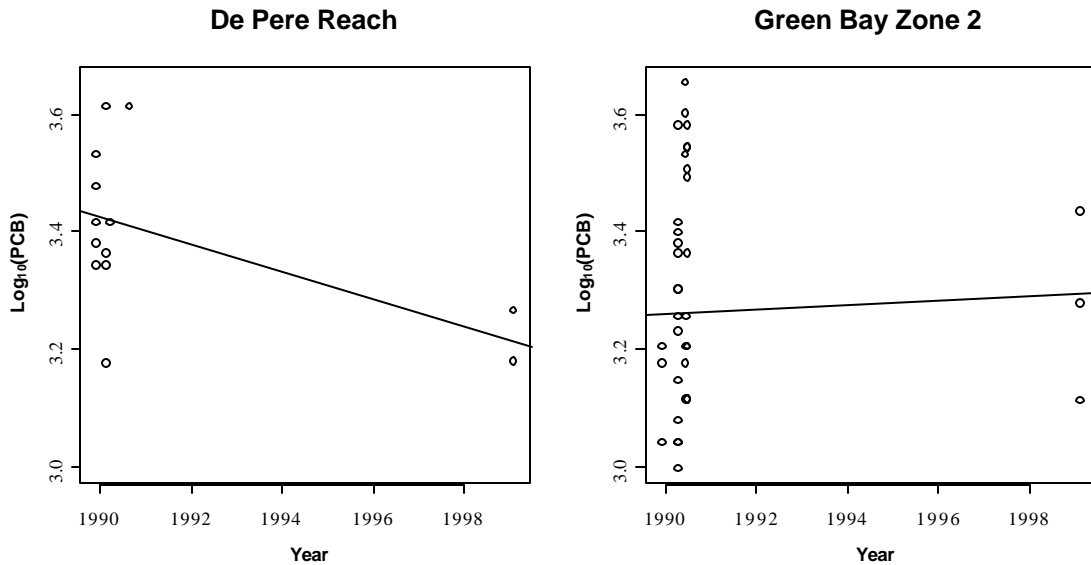


Figure 45 1989–1998 Time Profile Comparison of PCBs Between De Pere Reach and Green Bay Zone 2 for Alewife, Whole Body

Alewife whole body samples from the De Pere Reach show higher levels of PCBs around 1989–1990 than alewife in Green Bay Zone 2. By 1998, the PCB levels in the De Pere Reach appear to have dropped to levels comparable to those of Green Bay Zone 2.

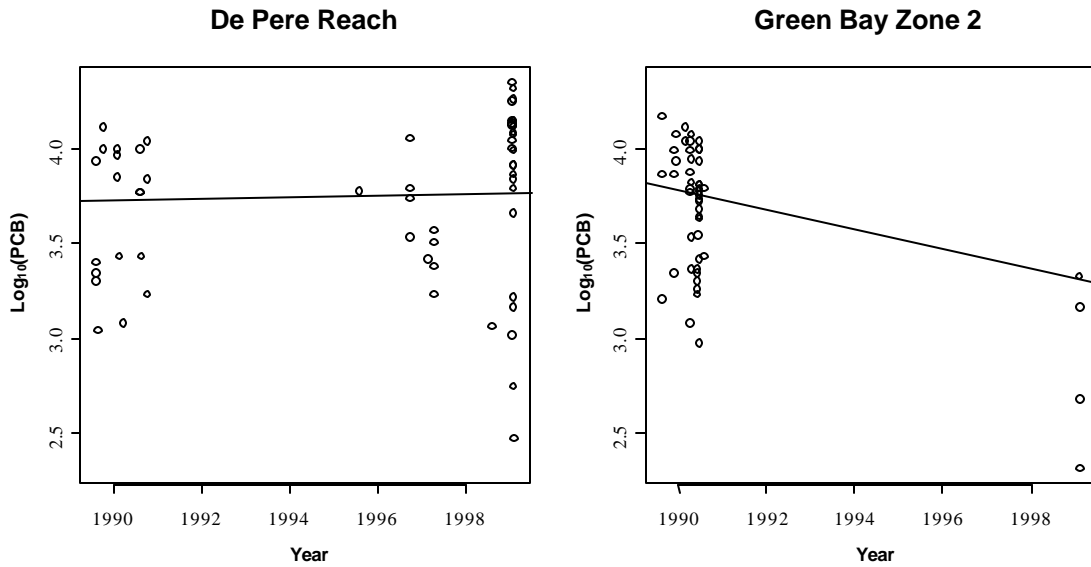


Figure 46 1989–1998 Time Profile Comparison of PCBs Between De Pere Reach and Green Bay Zone 2 for Carp, Whole Body Samples

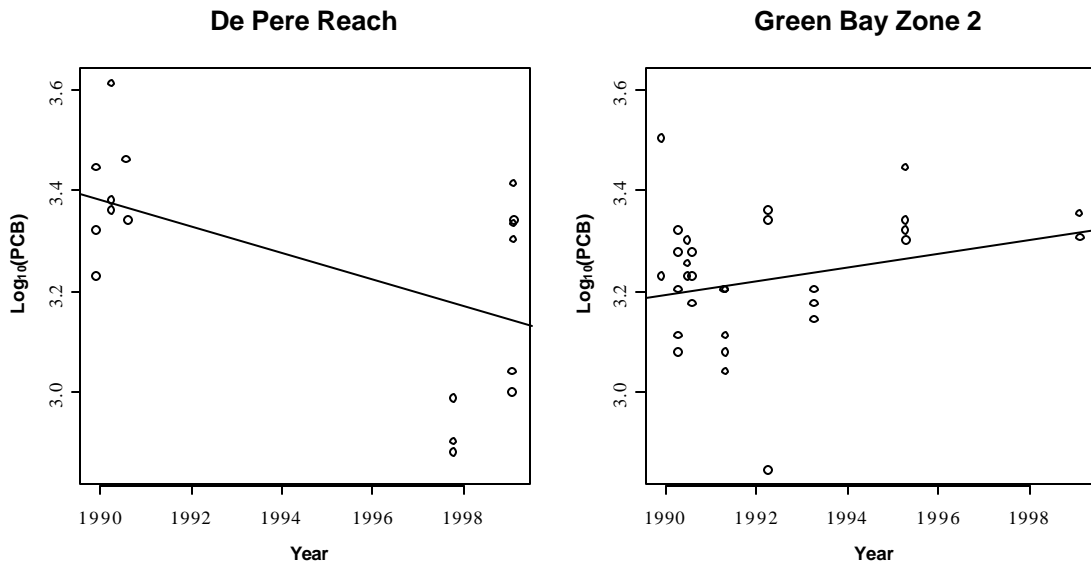


Figure 47 1989–1998 Time Profile Comparison of PCBs Between De Pere Reach and Green Bay Zone 2 for Gizzard Shad, Whole Body Samples

The majority of the analyses of species comparing De Pere Reach and Green Bay Zone 2 show statistically significant differences. Although no solid barriers

separate the two zones, the fishes sampled exhibit enough differences to suggest that the fish in the two zones are heterogeneous in either exposure to PCBs or processing of PCBs or both.

5.4.3 De Pere Reach versus Green Bay Zone 2: Without Adjustment for Lipid Concentrations

We carried out a second comparison of De Pere Reach and Green Bay Zone 2 in terms of PCB concentrations in fish. In this second analysis, the lipid weight as a percentage of tissue weight was excluded from the analysis. Analyses of PCB concentrations in fish often proceed without lipid normalization. Results presented here can then be compared to such “lipid-less” analyses. Lipid-based analyses are preferred when available, however, due to the occurrence of many highly significant associations of PCB concentration and lipid content.

The statistical model used for comparing the PCB concentration between De Pere Reach and Green Bay Zone 2 for samples collected during a short time period (snapshot analysis) is:

Equation 20

$$\log_{10}(PCB) = b_0 + b_2 R + e$$

where log PCB parameters and variables have the same definition as for the lipid-based analysis. We defined $R = 1$ for De Pere Reach and $R = 2$ for Green Bay Zone 2.

Fish sampled from De Pere Reach have an expected log concentration $b_0 + b_2$; those sampled from Green Bay Zone 2 have an expected log concentration is $b_0 + 2b_2$. Thus, if the coefficient b_2 is zero or if its difference from zero is small and not statistically significant, we would accept the hypothesis that the mean PCB concentrations in the given species and sample type are equal in the two reaches.

The model for comparing De Pere Reach and Green Bay Zone 2 when data on PCB concentrations have been collected over a longer period expands on the previous model by inserting terms involving time. The model is:

Equation 21

$$\log_{10}(PCB) = b_0 + b_2 R + b_4 t + b_5 t \cdot R + e$$

where the parameters and variables are as defined earlier (with $R = 1$ or 2). The time trend slope for De Pere Reach in this model is $(b_4 + b_5)$ and $(b_4 + 2b_5)$ for Green Bay Zone 2.

For the comparison of fish PCB concentrations in the De Pere Reach versus those in Green Bay Zone 2, we would accept the hypothesis that a given fish species and sample type has an equal mean PCB concentration in the two reaches at any

specified time if the coefficients b_2 and b_5 are small and not significantly different from zero. When these two coefficients are zero, then the rate of change (slope) of log PCB concentrations versus time in the two reaches is the same ($b_5 = 0$), and there is no difference in the expected mean log concentration ($b_2 = 0$) at any given time.

The results of the lipid-less snapshot analysis can be presented readily as a comparison of geometric means of PCB concentrations (see Table 29). For reference, we include the corresponding results for a lipid-based analysis, using 6 percent lipid content as a “plug-in” for the lipid-based snapshot equation (3 percent for walleye). We note, in general, the weaker contrast in geometric mean PCB concentration between the two reaches without the lipid variable (compare “percent increase” columns of the table). Also, only two, rather than four, of the differences are statistically significant.

The time trend lipid-less analysis is presented in Table 30. There, three out of the four analyses show statistically significant differences between the reaches with, in each case, quite striking disparities in the annual percent change in PCB concentration. (See the top row for each species/type to find the difference in rates of change in the two reaches—parameter b_5 —and the row with “+” to view the final model after all non-significant terms have been dropped.) As noted earlier, we prefer models based on lipid content, a key variable, the absence of which may mislead.

5.4.4 De Pere Reach versus Green Bay Zone 2: Summary

The De Pere and Green Bay Zone 2 reaches do not have an equivalent relationship to PCBs based on the comparisons presented here. The same species and sample types generally differ between reaches either in the slope of time trends, the relationship of PCBs to lipid content, or in the mean PCB concentration, controlling for lipid content. As can be seen from the plots associated with this analysis, the De Pere Reach generally has higher PCB concentrations than Green Bay Zone 2.

5.4.5 Lipid Normalization

The lipid content of samples strongly predicts PCB concentrations in most of our analyses, and, therefore, is an important variable to include in the time trends models. Its association with PCB concentrations is statistically significant—and often highly significant—in 17 out of the 19 analyses of individual sample types (see Table 18). Also, 7 out of the 19 analyses have coefficients of the log lipid variable that differ significantly from 1.0, the value that yields results equivalent to the traditional lipid normalization calculated as $(PCB\ concentration)/(percent\ lipid\ content)$. Only one such significant difference—rather than seven—would be expected by chance if 1.0 were the true value for all species and sample types. Thus, the traditional lipid normalization does not always control for the lipid effect.

Table 29 Comparison of Geometric Mean Concentrations of PCB Concentrations in De Pere Reach and Green Bay Zone 2, With and Without an Adjustment for Lipid Content, Samples from 1989–1991 (“snapshot” analysis)

| Species | Sample Type | Lipids | | | | No Lipids | | | | Years | Sample Size: Total (DP/Z2) |
|--------------|-----------------------------------|---------------------------------------|--------------------------------------|---|-------------------------|-----------------------------|-----------------------------|---------------------------------------|-------------------------|-----------|----------------------------|
| | | De Pere Geometric Mean Conc. (mg/kg)* | Zone 2 Geometric Mean Conc. (mg/kg)* | Zone 2 GM Percent Increase over De Pere | Reaches differ? p-value | Zone 2 Geometric Mean Conc. | Zone 2 Geometric Mean Conc. | Zone 2 Percent Increase over De Pere* | Reaches differ? p-value | | |
| Alewife | Whole Fish/ Whole Body/ Composite | 2,799 | 1,504 | 86 | 0.00006 | 2,654 | 1,963 | 35 | 0.04 | 1989 | 45 (11/34) |
| Carp | Whole Fish/ Whole Body/ Composite | 4,158 | 2,896 | 44 | 0.02 | 4,528 | 5,116 | -11 | 0.5 | 1989 | 66 (21/45) |
| Gizzard Shad | Whole Fish/ Whole Body/ Composite | 2,514 | 1,706 | 47 | 0.0009 | 2,450 | 1,717 | 43 | 0.002 | 1989 | 23 (9/14) |
| Walleye | Skin-on Fillet | 1,672 | 1,562 | 7 | 0.7 | 1,511 | 1,476 | 2 | 0.8 | 1989–1991 | 28 (11/17) |
| Walleye | Whole Fish/ Whole Body/ Composite | 6,201 | 4,242 | 46 | 0.006 | 6,995 | 5,835 | 20 | 0.1 | 1989 | 56 (25/31) |

Note:

* Based on a fitted model and a lipid percentage of 6 percent of weight except for walleye fillet with skin, where 3 percent was used.

Table 30 Models Comparing Log (PCB Concentration) versus Time in De Pere Reach and Green Bay Zone 2, Without Adjustment for Lipid Content

| Sample Type | Species | Model | Regression Model Parameter Statistics | | | | | | | | Likelihood Ratio Tests | | | Sample Size | | Total |
|---------------------------------|--------------|--------|---------------------------------------|-----------------------------|----------------------------------|-----------------------------|-----------------------------------|-----------------------------|--|-----------------------------|--|---|--|---------------|------------------|-------|
| | | | Constant Parameter (b ₀) | Std. Err. (b ₀) | Time Parameter (b ₄) | Std. Err. (b ₄) | Reach Parameter (b ₂) | Std. Err. (b ₂) | Time x Reach Interaction Parameter (b ₅) | Std. Err. (b ₅) | III vs. II Equal Slopes: Time x Reach Effect p-value | II vs. I Equal Intercepts: Reach Effect p-value | III vs. I Equal Slopes and Intercepts: Full Reach Effect p-value | De Pere Reach | Green Bay Zone 2 | |
| Whole Fish/Whole Body/Composite | alewife | III | 89.7407 | 63.163 | -0.0433 | 0.0317 | -42.95 | 37.59 | 0.0215 | 0.0189 | 0.26 | 0.068 | 0.10 | 13 | 37 | 50 |
| | | II | 20.8587 | 18.597 | -0.0087 | 0.0093 | -0.105 | 0.057 | 0 | 0 | | | | | | |
| | | I(+) | 18.2357 | 19.169 | -0.0075 | 0.0096 | 0 | 0 | 0 | 0 | | | | | | |
| Whole Fish/Whole Body/Composite | carp | III(+) | -242.08 | 64.123 | 0.1230 | 0.0322 | 219.60 | 49.47 | -0.1100 | 0.0248 | 0.00002*** | NA | 0.00005*** | 64 | 49 | 113 |
| | | II | 27.37 | 22.514 | -0.0118 | 0.0113 | -0.1161 | 0.0926 | 0 | 0 | | | | | | |
| | | I | 11.30 | 18.635 | -0.0038 | 0.0094 | 0 | 0 | 0 | 0 | | | | | | |
| Whole Fish/Whole Body/Composite | gizzard shad | III(+) | 143.151 | 37.016 | -0.0701 | 0.0186 | -85.6 | 25.41 | 0.0429 | 0.013 | 0.0014** | NA | 0.0028** | 18 | 32 | 50 |
| | | II | 25.6608 | 13.639 | -0.0112 | 0.0068 | -0.06 | 0.048 | 0 | 0 | | | | | | |
| | | I | 20.0254 | 13.061 | -0.0084 | 0.0066 | 0 | 0 | 0 | 0 | | | | | | |
| Whole Fish/Whole Body/Composite | walleye | III(+) | -24.0765 | 49.96 | 0.0141 | 0.0251 | 70.136 | 38.47 | -0.0353 | 0.019 | 0.0707 | 0.0569 | 0.0319* | 44 | 34 | 78 |
| | | II | 62.2372 | 16.529 | -0.0293 | 0.0083 | -0.119 | 0.062 | 0 | 0 | | | | | | |
| | | I | 51.1061 | 15.85 | -0.0238 | 0.008 | 0 | 0 | 0 | 0 | | | | | | |

Notes:

(+) Model indicated by likelihood ratio test. Coefficients appear in Equation 21.

* $p < 0.05$

** $p < 0.01$

*** $p < 0.001$

The statistically significant coefficients of the log percent lipid term in the models range from 0.43 up to 1.09. We noted earlier that a change in the lipid content by a multiplicative factor of F (e.g., $F = 2$, doubling the percent) leads to a change in PCB concentration by a multiplicative factor of F^{b_1} , where b_1 is the coefficient of \log_{10} lipid percentage in a regression model. The percentage change corresponding to F is $100\% * (F^{b_1} - 1)$. The observed range of significant lipid coefficients of 0.43 to 1.09 in the 19 analyses implies that a doubling of lipid percentage, for example, leads to a range of 34 to 113 percent increase in PCB concentration. The strong association between lipids and PCB concentration is illustrated by an example, Figure 48, where the positive association between log (PCB concentration) and log (percent lipid) is evident from the sparseness of points in the upper left and lower right of the plot.

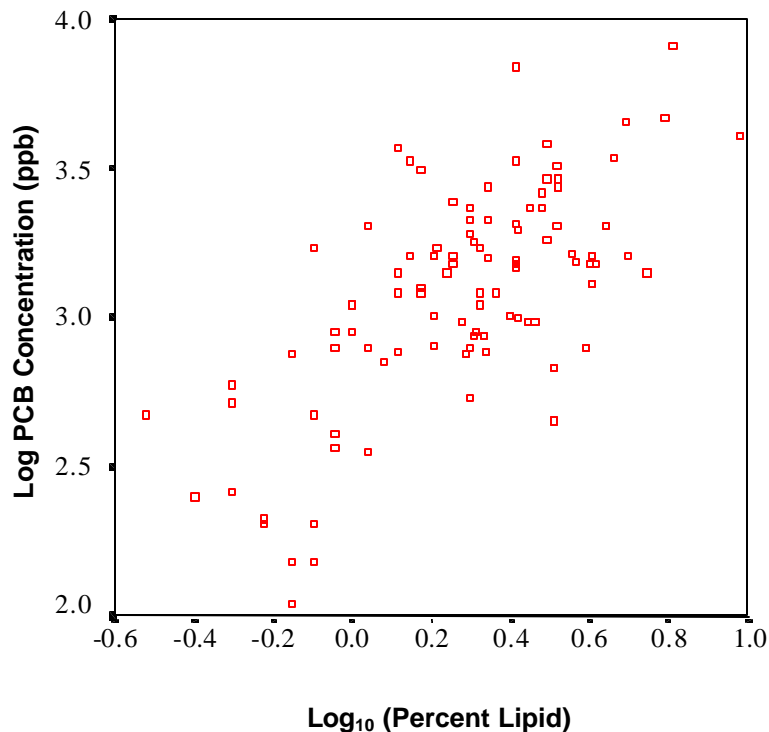


Figure 48 Log_{10} PCB Concentration (ppb) versus Log_{10} Percent Lipid Content for Walleye, Skin-on Fillet, De Pere to Green Bay Reach

The relationship between total PCBs and percent lipids (a measure of body fat) is strong. To adjust for this relationship, \log_{10} (percent lipids) must be included as an independent covariate in regression analyses.

6 Conclusions and Discussion

The analysis of trends in fish tissue and sediment over time in the Lower Fox River has led us to several significant conclusions. These conclusions, supporting statements, and discussion are included in Section 6.1. In addition, Section 6.3 identifies uncertainties associated with this trends analysis.

6.1 Conclusions

Data collected in the Lower Fox River and Green Bay show that concentrations of PCBs in fish tissue and surface sediments declined following the elimination of PCB point source discharges. However, further analysis of that data has identified statistically significant breakpoints in the decline for most of the fish species examined, suggesting that the decline has slowed down or, in some cases, that tissue concentrations of PCBs have increased. Furthermore, the analysis shows that it is not possible to project PCB concentrations into the future for fish or sediment with confidence because time trends appear to be quite changeable and confidence intervals for rates are quite wide.

Data on PCBs in sediment samples taken from surface sediments suggest that PCB concentrations have generally declined over time. Trends in concentrations of PCBs in subsurface sediments are less clear—some deposits show declining trends, while others show trends either close to zero or not significantly different from zero and yet others show increasing trends.

Specific conclusions of the time trends analysis include:

- **Fish tissue concentrations have generally declined over the period of time for which there are data in the Lower Fox River and Green Bay Zone 2.** Fish tissue PCB concentrations generally showed a slow rate of decline throughout the Lower Fox River and Green Bay Zone 2. Most time trend slopes were negative, and all statistically significant slopes were negative except one.
- **Significant “breakpoints” in the decline were identified for most of the fish species examined, suggesting that rates of decline in PCB tissue concentrations are changeable and slowing and, in some cases, tissue concentrations may be increasing.** Fish tissue concentrations have not declined at a constant rate since the 1970s. Among fish time trends analyzed, 7 out of 19 combinations of reach, species, and sample type showed a statistically significant change in slope (log scale) between earlier and later periods. In Little Lake Butte des Morts, De Pere Reach, and in Green Bay Zone 2, there were steep declines in fish tissue PCB concentrations from the 1970s, but with significant breakpoints in declines for some species beginning around

1980. A meta-analysis of time trends showed that the most recent slopes averaged across species showed a 5 to 7 percent decline per year for three of the reaches. Six species showed an increasing rate in their final slope, but only two of these rates were statistically significant (carp, whole body, in De Pere Reach and in Green Bay Zone 2). The existence of breakpoints plus a meta-analysis to detect non-constant trends suggest that rates of change are changeable and not constant.

- **PCBs in surface sediment samples have generally declined over the period of time for which there are data for the Lower Fox River.** Surface sediment PCB concentrations combined within each reach showed statistically significant decreasing trends in all reaches except Appleton to Little Rapids. There were wide confidence intervals for rates of change, both for individual deposits and combined deposits, indicating that rate estimates are not precise. This imprecision and other uncertainties associated with the data do not support accurate future projections. Surface sediments of individual deposits within the reaches included a mixture of positive and negative slopes. Among the 16 negative slopes, 6 were statistically significant; neither of the 2 positive slopes were statistically significant.
- **Time trends in PCB concentrations in sediments below the surface sediment are less clear—some indicate a decline, others indicate no change or increases, others are unchanging or even increasing.** There is a strong trend toward fewer and weaker negative slopes at increasing depths. For Little Lake Butte des Morts, subsurface trends are mixed. The only statistically significant subsurface trend shows an increase and the other trends are a mixture of positive and negative trends. In the Appleton and De Pere reaches, there is a mixture of positive and negative trends that is not clearly distinguishable from a zero overall trend. For Little Rapids to De Pere, there are consistently negative trends in the 10- to 30-cm strata, but in the lower strata, the data are consistent with either a zero trend (30 to 50 cm), or an increasing trend (50 to 100 cm and 100+ cm).
- **Projection of PCB concentrations into the future for fish or sediments is questionable because of imprecision and other uncertainties identified in the analysis.** The analyses carried out cannot assure a continued decline in PCB concentrations over time. Even though there are a number of negative time trends that suggest PCB declines, future projection is questionable. Increases in PCB concentrations in some deeper sediments and breakpoints, and other non-linear phenomena in fish PCB time trends (on the log scale) suggest that the river, its sediment, and its fish species could experience an arrest or reversal of such a decline.

- **PCB concentrations may increase or decrease in the future.** Some, perhaps all, of the changes in slope from before to after a breakpoint in the fish analysis may be genuine, due to unpredictable events, such as floods accompanied by scouring and deposition. As discussed in the Remedial Investigation, sediment bed elevations have been altered historically and may also undergo changes in the future due to scouring and redistribution of sediments. The occurrence of these breakpoints in the past suggests that the river may change again in the future. The presence of non-constant slopes (which we refer to as “curvature”) in the post-breakpoint period also suggests change. If so, such events will continue adding variability to PCB concentration over time, making predictions based on the assumption of a future decline at a constant rate questionable. The presence of curvature is consistent with the more dramatic changes represented by breakpoints and supports the notion of a dynamic process, liable to change, rather than a steady state with future constant linear rates of change.

The last two bullets are especially germane to use of the time trends analysis in other elements of the Lower Fox River Risk Assessment and Feasibility Study. The time trends were estimated only for the period of time for which data exist. These analyses are not suitable for accurately projecting trends into the future. Of particular importance, the data do not provide assurance of a continuing future decline in PCB concentrations.

The time trends analysis has dealt strictly with the testing of changes in PCB concentrations over time in the Lower Fox River, and not with the mechanisms that could control changes in sediment and tissue loads. The apparent decline of PCBs observed in surface sediments and fish from the Lower Fox River are consistent with the continued observed transport of PCBs from the river to Green Bay, as discussed in detail in the Remedial Investigation. Changes in sediment bed elevations have been documented and are discussed in Technical Memorandum 2g (WDNR, 1999a) and in the Remedial Investigation. Some of the variability observed in the data may be accounted for by changes in river profile, burial, scour by flood or ice, and propeller wash in the lower reaches of the river. As the analysis focused solely on the existing data in the Lower Fox River and Zone 2 of Green Bay, these potential mechanisms were not introduced into the analysis and thus could not be controlled. What is important to note, however, is that the trends analysis is dependent upon the existing hydraulic conditions in the Lower Fox River. Any changes in those conditions might result in exposure of underlying PCB-laden sediments or burying of sediments, and lead to new trends that may not be similar to the trends from this analysis.

The conclusions of a general historical decrease in PCB burdens in sediments and fish of the Lower Fox River and in Zone 2 of Green Bay are similar to those reported by other Great Lakes researchers. Decreases in PCB concentrations have been observed in Lake Michigan (Offenberg and Baker, 2000; DeVault *et al.*, 1996; Lamon *et al.*, 1998), Lake Ontario (DeVault *et al.*, 1996; Gobas *et al.*,

1995), Lake Superior (Smith, 2000; DeVault *et al.*, 1996) and lakes Huron and Erie (DeVault *et al.*, 1996). The yearly rate of decline for PCBs in biota and sediment of Lake Superior has been estimated at 3 to 8 percent per year and is expected to continue at 5 to 10 percent per year (Smith, 2000), which is generally consistent with the trends observed in the Lower Fox River. However, several other researchers have also noted breakpoints, or constant levels of PCBs beginning in the mid- to late 1980s. PCB concentrations in lake trout and smelt are reported to have been relatively constant in Lake Ontario since 1985 (Gobas *et al.*, 1995) while concentrations in other fish and in sediments show a decline during the period of observed data (to about 1990) and a projected continuing decline (see Gobas *et al.*, 1995, Figures 2 and 3). PCB body burdens in Lake Erie walleye were shown to be declining during the period of 1977 through 1982, but after that period remained constant through 1990 (DeVault *et al.*, 1996). Time trends analysis for salmonids and trout in Lake Michigan showed generally decreasing tissue concentrations (Lamon *et al.*, 1998). The uncertainty in rates is often large, and some trends are not significantly different from a zero rate or have confidence intervals that include positive rates of increase (e.g., lake trout, see DeVault *et al.*, 1996, Figure 3). These findings are consistent with the time trends analysis for the Lower Fox River and suggest that there may continue to be slow, gradual declines, or a steady state in PCB concentrations in fish and sediment in the future. The possibility of some increases cannot be ruled out.

6.2 Time Trends Discussion

6.2.1 General Issues

The time trends analysis has shown that PCB concentrations in surface sediments (0 to 10 cm) and fish are generally decreasing over time. In both sediment and fish analyses, the magnitude and level of statistical significance of time trends varies widely. All except one statistically significant fish time trend indicated decreasing concentrations. The time trends in subsurface (10+ cm) sediments contain a mixture of positive and negative rates of change, and it is difficult to reach a firm conclusion about the subsurface PCB time trends. The time trends in sediment generally exceed in magnitude (positive or negative) those in fish. Most significant and non-significant sediment trends were negative, but there were some statistically significant positive trends for deeper strata. More is known about the trends in surface deposits because a larger fraction of the surface deposit groups than subsurface deposits were analyzed.

Sediment samples taken from the surface sediments have more negative than positive slopes. However, there was a trend toward fewer negative and more positive slopes as depth increased. In sediments sampled from the surface, 89 percent of slopes were negative. Below 50 cm, 71 percent of slopes were positive. The time trend analysis has shown that rates of change of PCB concentrations in fish are themselves liable to change, calling into question the value of projecting concentrations under an assumed but unverifiable steady-state model. By

implication, sediment—particularly surface sediment—as the primary source of PCBs in fish, is also likely to be changeable in its time trends of PCB concentrations.

The meta-analysis (pooled results from all surface sediment deposits) for trends in surface sediments showed an average rate of decrease in PCB concentrations of 18 percent per year in Little Lake Butte des Morts, 0.6 percent per year increase in the Appleton Reach, 10 percent per year decrease in the Little Rapids Reach, and 15 percent per year decrease in the De Pere Reach. These meta-analysis trends were statistically significant except for the small trend in the Appleton Reach. Thus, surface sediments show decreasing PCB concentrations over time, and at a fairly rapid average rate during the period covered by the data, except for the Appleton Reach.

It is important to emphasize that it is the average rate of change over a period of time that is strikingly negative in three out of the four reaches, and not necessarily the individual deposit rates or even the rates at each point in time covered by the data. Given the findings of fish time trends that seem to vary over time, as evidenced by both breakpoints and curvature, it is likely that the sediment time trends may also be volatile over time, perhaps due to scouring and deposition, which are described in a companion document (WDNR, 1999a). There are simply too few distinct time points of measurement of sediment concentrations to support a breakpoint and curvature analysis such as that carried out for fish. Since the ultimate source of PCBs in fish is sediment, however, it is difficult to imagine that fish have volatile time trends with sediment volatility.

The fish meta-analyses within the three out of four reaches with more than one fish category available for analysis show that PCB concentration was most recently declining at a rate of 5 to 7 percent per year (Little Lake Butte des Morts, De Pere, Green Bay Zone 2). The single fish category that could be analyzed for the Appleton Reach also shows a decline of 10 percent per year.

However, the fish time trends are changeable. Little Lake Butte des Morts had a steeper decline in PCB concentrations in earlier years. All analyses with a breakpoint in this reach show a steeper decline before the breakpoint. In the other reaches, the data for each fish category considered individually are consistent with a constant rate of decline over the whole time period, except for 2 out of 12 combinations of species and sample type. Nevertheless, the collective evidence demonstrates that slopes (on the log scale) tend to be non-constant. Based on a meta-analysis, the hypothesis of constant final slopes for all species was rejected and we must accept the concept of non-constant time trends for the post-breakpoint period for at least some species. In this regard, we note that it is possible to not detect curvature for analysis of individual species and yet to detect the presence of curvature from a global meta-analysis (and accept changing slopes for some individual species), because the meta-analysis has more power.

A practical dilemma in estimating future concentrations of PCBs is the choice of a statistical model to use in projecting concentrations forward in time, both for sediment and fish. For sediment, there are insufficient data to test for “curvature” (a non-constant slope over time), though the fish analysis implies curvature and changeability of slopes. Using the fitted time trends as presented in this report for projection and ignoring the possibility of non-constant sediment time trend slopes assumes a steady state in the river and, consequently, could lead to erroneous future projections. Such error in the projection is likely to be smaller, when one aggregates the results of projections of individual deposits into larger geographic units, such as a reach or the entire river. There is disagreement between fish and sediment time trends. The average rates of decrease of PCB concentration in the meta-analysis of surface sediments generally exceed those observed in the meta-analysis of fish PCB trends. Biologically, fish rates should have to be linked with and similar to those for sediment. One possible explanation for the mismatch is that the sediment rate of decrease may have slowed down recently. There are too few time points with sediment data, per deposit group and depth, to detect such a slowing, and the calculated rate of change for sediment PCB concentration may be an average of a faster earlier rate and a slower recent rate.

6.2.2 Fish Lipids

Lipid content of samples distinctly assisted in reducing unexplained variance for most analyses of fish PCB time trends. Since it is so helpful, efforts should be taken in the future to explore ways to more powerfully incorporate lipids into the analysis. The time trend analysis used lipid content as a linear independent variable. We prefer this approach to the alternative of dividing PCB concentration by the percent lipid content, which is equivalent to using lipids as an independent variable but forcing its coefficient to be unity.

Only two analyses of time trends in gizzard shad (for two different reaches) showed no significant relationship between lipids and PCBs, suggesting that some species may handle PCBs in a different fashion. The variety of coefficients relating PCBs to lipid content among the various species and sample types suggests that species are not identical in their PCB-lipid relationships.

6.2.3 Strengths of the Study

There are a number of strengths of the study. The maximum likelihood method used to handle data below a detection limit allowed these values to contribute to the analysis without having to impute a proxy value. The methods used to detect and handle spatial correlation of sediment samples have allowed us to avoid overstating statistical significance of time trends. In fact, statements of statistical significance should be quite conservative. Our approaches to quantifying and testing for non-constant rates of change in fish time trends (breakpoints and curvature) have allowed us to assess the changeability of time trends. Our use of regression analysis of lipid content as a factor in PCB concentrations makes good use of the lipid data and does not impose a pre-specified coefficient relating PCBs

to lipid content. The use of meta-analysis of rates has increased precision and power in time trend estimates. The remarkable agreement of the data with the lognormal distribution and the need to address only two outliers in over 2,000 observations, support the overall validity of the PCB concentrations used in the time trends analysis. The inherent very great variability of the PCB concentrations has been thoroughly described quantitatively, through confidence intervals of slopes, and graphically, by scatter plots of concentrations versus time. Finally, clear statements about confidence in and statistical significance of the various quantitative trends have been provided to guide the reader in the use of the trends.

6.3 Sources of Uncertainty in the Time Trends Analysis

The conclusions and discussion presented above are based upon the statistical analyses of the data as received by us. However, there are areas of uncertainty that may have played a role in this analysis. By “uncertainty,” we mean either random variation (such as fish-to-fish variation in PCB concentration) or systematic variation due to unmeasured factors, such as age and gender of fish or changes in the absolute elevation of the sediment-water interface. While statisticians use terms such as “variation” and “sources of unexplained variation” for these two effects, we will use the term “uncertainty,” a term more familiar to readers, to specifically designate the combination of these two effects. While there is no uncertainty about the methodology, the results should be considered as possibly influenced by unmeasured factors, hence uncertain to that extent.

In addition to the uncertainty arising from sheer randomness, there are sources of uncertainty associated with laboratory and analytical variation and other factors that could not be included in the analysis. The various sources of uncertainty are discussed below.

6.3.1 Statistical Uncertainty — Statistical Significance and Confidence Intervals

The data used for both sediment and fish time trends analyses are inherently quite variable. A wide scatter of points typically surrounds the regression lines for fitted models. This variability has led to some wide confidence intervals around estimated values. The lack of statistical significance of a time trend does not imply the absence of a real trend, even a strong one. Some attention to confidence intervals shows the possibility of strong trends that may not have been detected due to the large random component in the data.

We suggest that the reader take note of the statistically significant trends and use the confidence intervals for these and other trends as statements ruling out (with high confidence) certain slopes outside the confidence intervals. Slopes within the confidence intervals (usually quite wide) are all quite plausible and consistent with the data. These confidence intervals are usually quite wide. Because the

confidence intervals are generally wide, they cannot usually be used to state that a trend is close to zero. Within the intervals, there are differing rates of change.

By examining the standard errors of slope estimates of \log_{10} PCB concentration versus time, a quantitative notion of the statistical uncertainty in the time trend estimates can be expressed. A standard error (SE) of 0.0054 for a slope estimate on the \log_{10} scale would indicate “excellent” precision because, for example, a slope of zero (zero percent change per year) with an SE of 0.0054 would lead to a 95 percent confidence interval (CI) for the rate of change of --2.5 to +2.5 percent, a tight range of 5 percentage points. None of the 46 sediment trends and only 3 out of the 19 fish trends have this precision (Table 9 and Table 18).

“Good” or “fair” precision would be an SE of 0.01 or less, which, for a zero slope, would have a 95 percent CI of ± 5 percent, a range of 10 percentage points. Two sediment and nine fish time trends have this precision. Among the meta-analyses, all of the fish combined time trend slopes have good-to-excellent precision (Table 19), but none of the combined surface sediment time trends has this precision (Table 10). Even “good” or “fair” precision of ± 5 percent provides room for very different future scenarios. A rate of 5 percent decrease per year for 10 years leads to a 40 percent loss in PCB concentration, while a 5 percent increase per year for 10 years leads to a 63 percent increase in PCB concentration. The range -40 to +63 percent is a wide zone of uncertainty.

Indeed, one of the firm conclusions of this study must be that, in some cases, a firm conclusion cannot be reached. An increasing or decreasing time trend that is statistically significant, or a trend that is not significantly different from zero but with a tight confidence interval around zero, provides a clear outcome. Non-significant trends with wide confidence intervals impart little information and do not provide a clear outcome. Thus, Table 31 and Table 32 show which calculated time trends provide a “clear outcome” and which trends have “good” or “fair” precision.

Table 31 Sediment Time Trends: Analyses with Clear Outcomes and Good Precision

| Type of Analysis | | Clear Outcome: Significant Increase or Decrease, or Confidently Close to Zero? ⁺ | Is Precision Good or Fair? ⁺⁺ |
|------------------------------------|----------|---|---|
| Sediment | | | |
| <i>Little Lake Butte des Morts</i> | | | |
| Deposit Group AB | 0-10 cm | Decrease | — |
| | 10-30 cm | — | — |
| | 30-50 cm | — | — |
| C | 0-10 cm | — | — |
| | 10-30 cm | — | — |
| POG | 0-10 cm | — | — |

Table 31 Sediment Time Trends: Analyses with Clear Outcomes and Good Precision

| Type of Analysis | | Clear Outcome: Significant Increase or Decrease, or Confidently Close to Zero?* | Is Precision Good or Fair? ** |
|--|---|---|----------------------------------|
| D | 0-10 cm | Decrease | — |
| | 10-30 cm | Increase | — |
| F | 0-10 cm | Decrease | — |
| | 10-30 cm | — | — |
| GH | 0-10 cm | Decrease | — |
| Little Lake Butte des Morts Surface Meta-analysis | | Decrease | — |
| <i>Appleton Reach</i> | | | |
| Deposit Group IMOR | 0-10 cm | — | — |
| | N Pre-dredge 0-10 cm | Decrease | Yes |
| | 10-30 cm | — | — |
| VCC | 30-50 cm | — | — |
| | 0-10 cm | — | — |
| | 10-30 cm | Decrease | — |
| | 30-50 cm | — | — |
| | Appleton Reach Surface Meta-analysis | | — |
| <i>Little Rapids Reach</i> | | | |
| Deposit Group Upper EE | 0-10 cm | — | — |
| | 10-30 cm | — | — |
| | 30-50 cm | — | — |
| Lower EE | 0-10 cm | Decrease | — |
| | 10-30 cm | — | — |
| | 30-50 cm | Increase | — |
| FF | 0-10 cm | — | — |
| | 10-30 cm | Decrease | — |
| GGHH | 0-10 cm | — | — |
| | 10-30 cm | — | — |
| | 30-50 cm | — | — |
| | 50-100 cm | — | — |
| | 100+ cm | — | — |
| Little Rapids Reach Surface Meta-analysis | | Decrease | — |

Table 31 Sediment Time Trends: Analyses with Clear Outcomes and Good Precision

| Type of Analysis | | Clear Outcome: Significant Increase or Decrease, or Confidently Close to Zero? ⁺ | Is Precision Good or Fair? ⁺⁺ |
|--|-----------|---|---|
| <i>De Pere Reach</i> | | | |
| SMU Group 2025 | 0–10 cm | — | — |
| | 10–30 cm | — | — |
| | 30–50 cm | — | — |
| | 50–100 cm | — | — |
| 2649 | 0–10 cm | Decrease | Yes |
| | 10–30 cm | — | — |
| | 50–100 cm | — | — |
| | 100+ cm | — | — |
| 5067 | 0–10 cm | Decrease | — |
| | 10–30 cm | — | — |
| | 50–100 cm | Increase | — |
| | 100+ cm | — | — |
| 6891 | 0–10 cm | — | — |
| | 10–30 cm | — | — |
| 92115 | 0–10 cm | — | — |
| De Pere Reach Surface Meta-analysis | | Decrease | — |

Notes:

- 1. “Yes” indicates increase or decrease is statistically significant compared to zero rate of change ($p < 0.05$), **or** 95 percent confidence interval for percent change is within ± 5 percent of zero.
- 2. Uncertain outcome noted by “—” (not “Yes” to above).

⁺⁺ Standard error of slope ≤ 0.1 .

Of the 46 deposit group analyses in Table 31 and 4 surface sediment analyses, only 16 cases can offer us a reasonably firm conclusion on time trends. Two indicate increasing, and 14 indicate decreasing, trends. The remaining 34 analyses have uncertain trends. All cases noted with a dash (“—”) in the “Clear Outcome” column may have trends that deviate more than ± 5 percent per year from a constant, 0 percent rate of change, and the rate may plausibly be either positive or negative. In these cases, a zero rate is just one among a wide range of possible rates bracketing zero. As noted in the “Precision” column of Table 31, only two analyses provide good or fair precision for their time trends.

The fish analyses provide a firmer set of conclusions (Table 32). Among the 19 primary analyses and 3 meta-analyses, 17 clearly demonstrate an “increase” or “decrease.” The other five analyses do not support a solid “no change,” zero-slope conclusion, but instead leave us with a fairly wide range of plausible increasing or decreasing slopes. As far as precision goes, 14 out of the 22 analyses provide “good” or “fair” precision for fish trend estimates.

Table 32 Fish Time Trends: Analyses with Clear Outcomes and Good Precision

| Type of Analysis | Clear Outcome: Significant Increase or Decrease, or Confidently Close to Zero? ⁺ | Is Precision Good or Fair? ⁺⁺ |
|--|---|---|
| Fish | | |
| <i>Little Lake Butte des Morts</i> | | |
| Carp, skin-on fillet | Decrease | Yes |
| Carp, whole body | — | — |
| Northern Pike, skin-on fillet | Decrease | Yes |
| Walleye, skin-on fillet | — | — |
| Walleye, whole body | — | — |
| Yellow Perch, skin-on fillet | — | — |
| Little Lake Butte des Morts Meta-analysis | Decrease | Yes |
| <i>Appleton Reach</i> | | |
| Walleye, skin-on fillet | Decrease | — |
| <i>De Pere Reach</i> | | |
| Carp, whole body | Increase | — |
| Gizzard Shad, whole body | Decrease | Yes |
| Northern Pike, skin-on fillet | Decrease | Yes |
| Walleye, skin-on fillet | Decrease | Yes |
| Walleye, whole body | Decrease | Yes |
| White Bass, skin-on fillet | Decrease | Yes |
| White Sucker, skin-on fillet | Decrease | Yes |
| De Pere Reach Meta-analysis | Decrease | Yes |
| <i>Green Bay Zone 2</i> | | |
| Alewife, whole body | Decrease | Yes |
| Carp, skin-on fillet | — | — |
| Carp, whole body | Decrease | Yes |
| Gizzard Shad, whole body | Increase | Yes |
| Yellow Perch, skin-on fillet | Decrease | Yes |
| Green Bay Zone 2 Meta-analysis | Decrease | Yes |

Notes:

- 1. “Yes” indicates increase or decrease is statistically significant compared to zero rate of change ($p < 0.05$), or 95 percent confidence interval for percent change is within ± 5 percent of zero.
- 2. Uncertain outcome noted by “—” (not “Yes” to above).

++ Standard error of slope ≤ 0.1 .

6.3.2 Physical Sources of Uncertainty

Depth of Sediments

The time trend analysis has shown that shallower sediment layers tend to have greater rates of decrease than deeper layers, where PCB concentrations may even be increasing. In Little Lake Butte des Morts, for example, Deposit Group D bears a strong and statistically significant decreasing trend at 0 to 10 cm and a

strong and highly significant increasing trend at 10 to 30 cm. Deposits with these trend patterns may be experiencing either burying of more contaminated surface sediments over time into deeper strata, or some mechanism whereby PCBs migrate downward.

Depth of sediment is closely related to PCB concentration. We used depth defined as the distance to the sediment-water interface. The Fox River database (the source of our data) does not include the absolute depth of deposits (in relation to fixed points and elevations on land). Such data would undoubtedly help in the analysis. The data available now do not allow us to track a given parcel of sediment over time. The interface may change over time due to scouring or deposition. Some of the time trends noted here may be due to a change in the depth from the sediment-water interface, where that boundary has shifted up or down due to deposition or scouring over time, so that different parcels of sediment are identified with the same depth label at different times. Time trends based on an absolute definition of depth would more accurately track what happens to PCBs in a specific volume of sediment over time.

Hydraulic Conditions

As noted above, there was no way to control in the time trends analysis for changes that may have occurred in sediment or fish tissue concentrations that could be attributed to flooding, ice scouring, propeller wash, or other mechanisms that would have caused changes to the hydraulic conditions in the river. Changes in bed elevations have been previously documented (WDNR, 1999a). While in one sense, the analysis of trends over time is concerned only with change, and not necessarily the underlying mechanism(s), an understanding of episodic events that may have influenced observed upward or downward trends would have facilitated the overall understanding of those results.

The trends reported here pertain to hydraulic conditions in the river at the time the data were collected. The system of locks and dams on the Lower Fox River currently control to a large degree where deposition and scouring occur. In the future, should those conditions change, any comparison of rates of change of PCB concentrations to the rates presented in this report, for the purpose of determining slowing or quickening of rates over time, would have to be done very cautiously.

6.3.3 Sources of Biological Uncertainty

Age and Gender of Fish

Age of fish may relate to PCB concentrations, due to different feeding habits and locations during the lifecycle. Incorporating age proxy variables (either length or mass) might reduce unexplained variance and increase power to detect trends. The relation of age to PCB concentration could be explored as either linear, curvilinear, or some type of step function (e.g., representing juveniles versus

adults). (Length data have recently become available for some samples as this analysis was completed.) Similarly, the gender of the fish and whether or not it recently spawned may be factors in PCB uptake and retention, and these factors can easily be incorporated into the analysis when data become available.

Spatial Dependence

The time trend analysis was not adjusted for and cannot, with present data, adjust for potential spatial dependence of data from fish samples. While individual fish do not have specific geographic coordinates, fish caught at about the same time and location may exhibit some dependence due to similar feeding sources.

6.3.4 Uncertainty Due to Laboratory and Analytical Factors

Our time trends analysis did not incorporate potential laboratory variation into the study. Multiple laboratories engaged in the analysis of sediments and fish tissues for the Lower Fox River and Green Bay, which is not uncommon for large environmental projects. Analytical variability amongst those laboratories is discussed in the Data Management Report (EcoChem, 2000). A “laboratory effect,” whereby different laboratories would produce a different mean PCB concentration on split samples, is possible. In addition, analytical techniques may have changed over the 1989-through-1998 period of sediment sample collection. Similarly, the 1976-through-1998 period of the fish samples included in the analysis may well have seen changes and refinements in laboratory equipment and techniques. Both the “laboratory effect” and changes in technique may have influenced the time trends.

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Appendix

Additional Data and Plots

APPENDIX B-1
TIME TRENDS IN PCB CONCENTRATIONS IN SEDIMENT
AND FISH

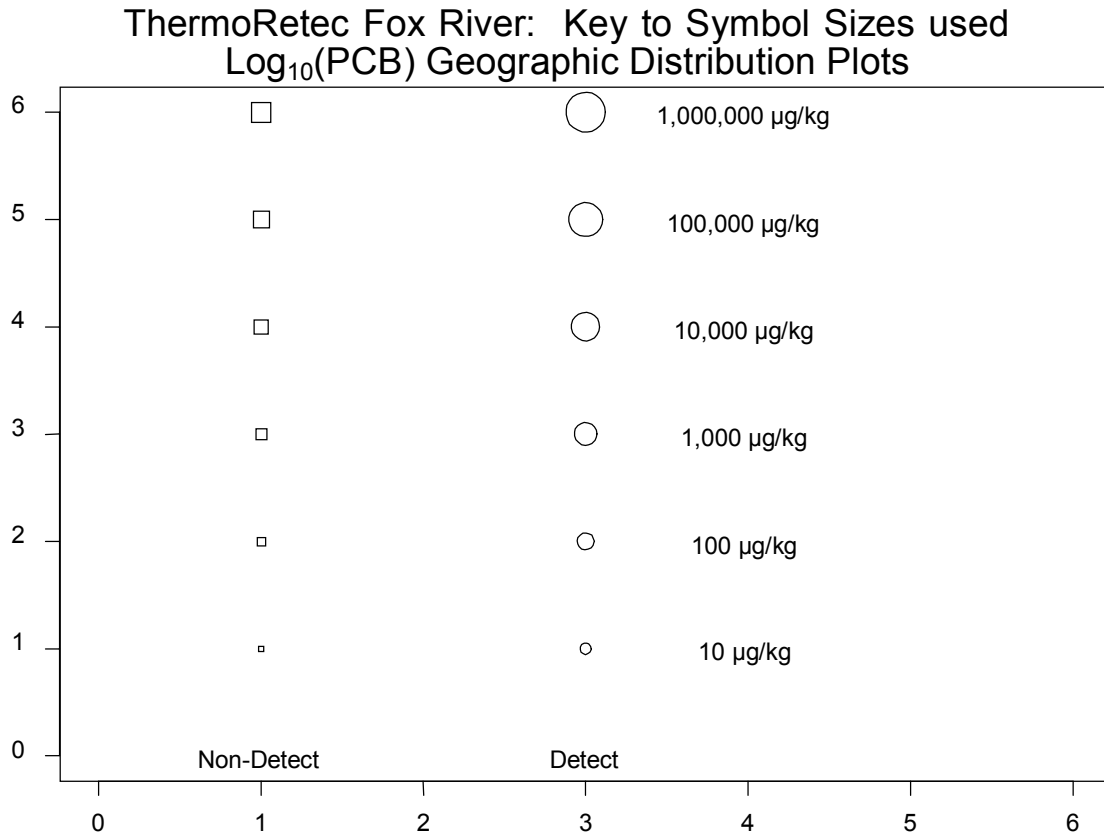


Figure A-1 Northing/Easting Plot Key

Scale plot showing the size of circles (for samples with detected PCBs) and squares (for samples with PCBs below detection limit, the square conveys the level at which the PCBs would have been detected as reported by the various testing agencies) used to convey total PCB level in the northing/easting plots of sample locations.

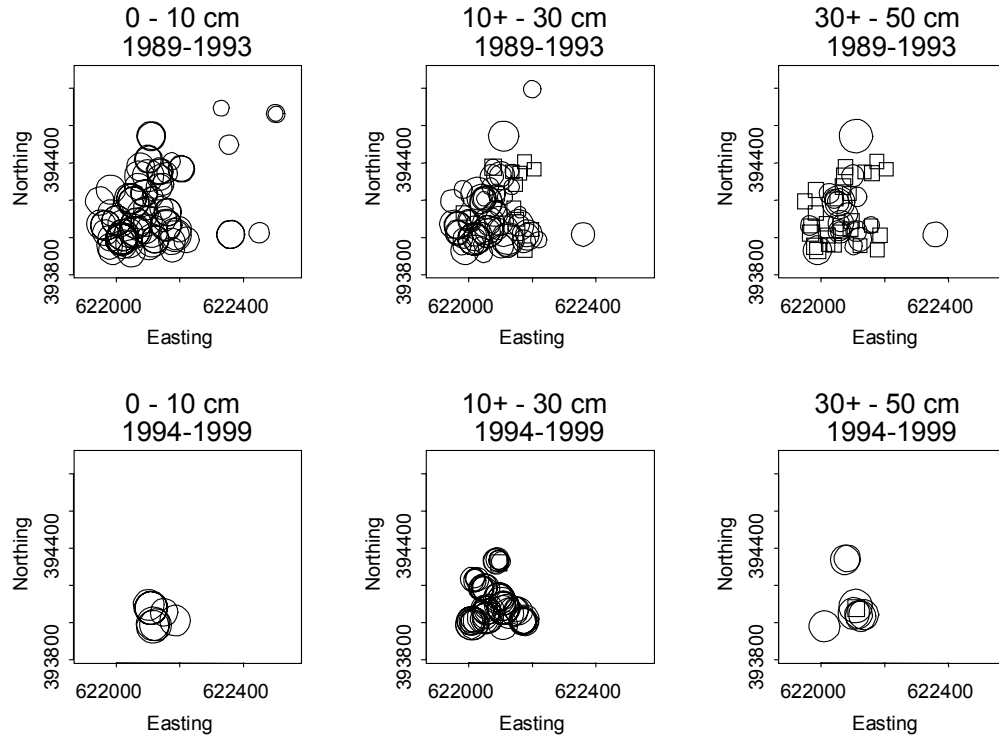


Figure A-2 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group AB (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

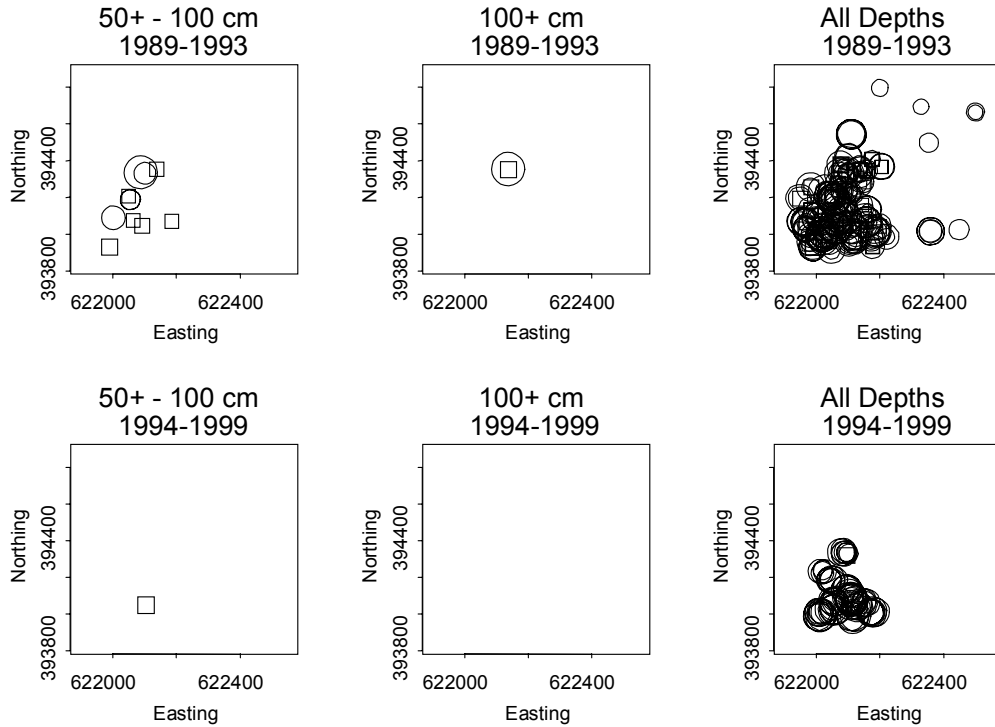


Figure A-3 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group AB (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

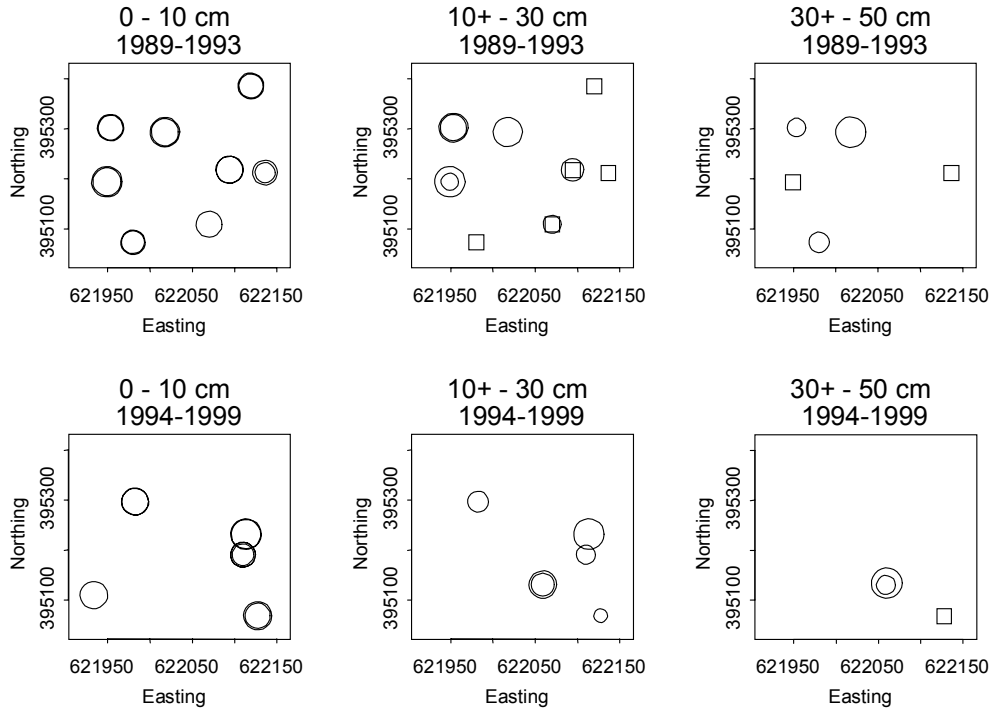


Figure A-4 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group C (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

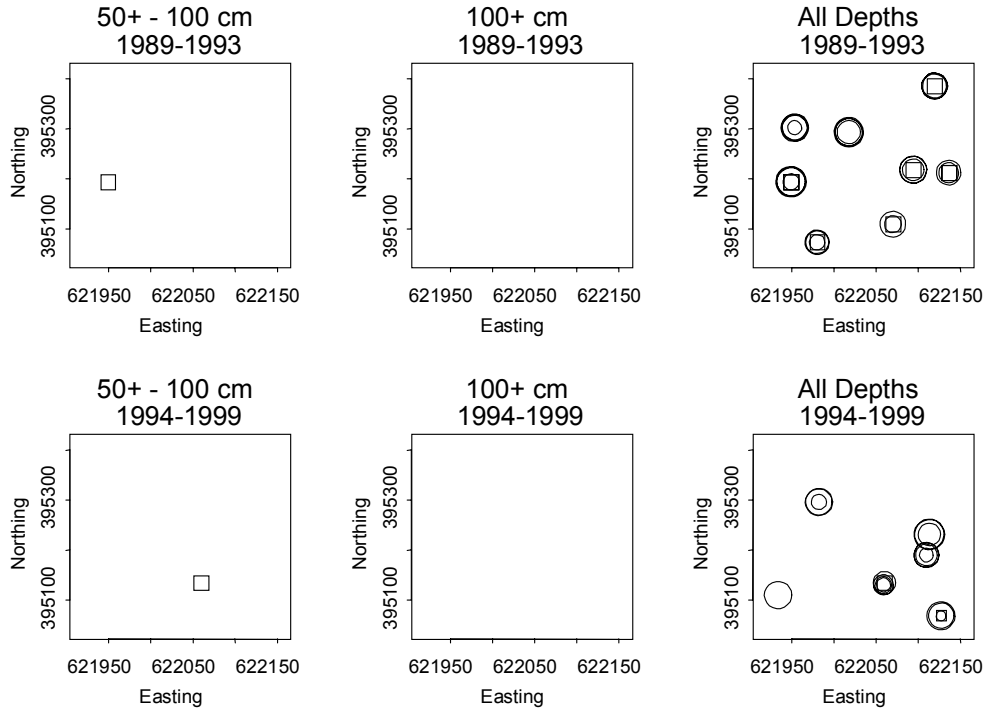


Figure A-5 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group C (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

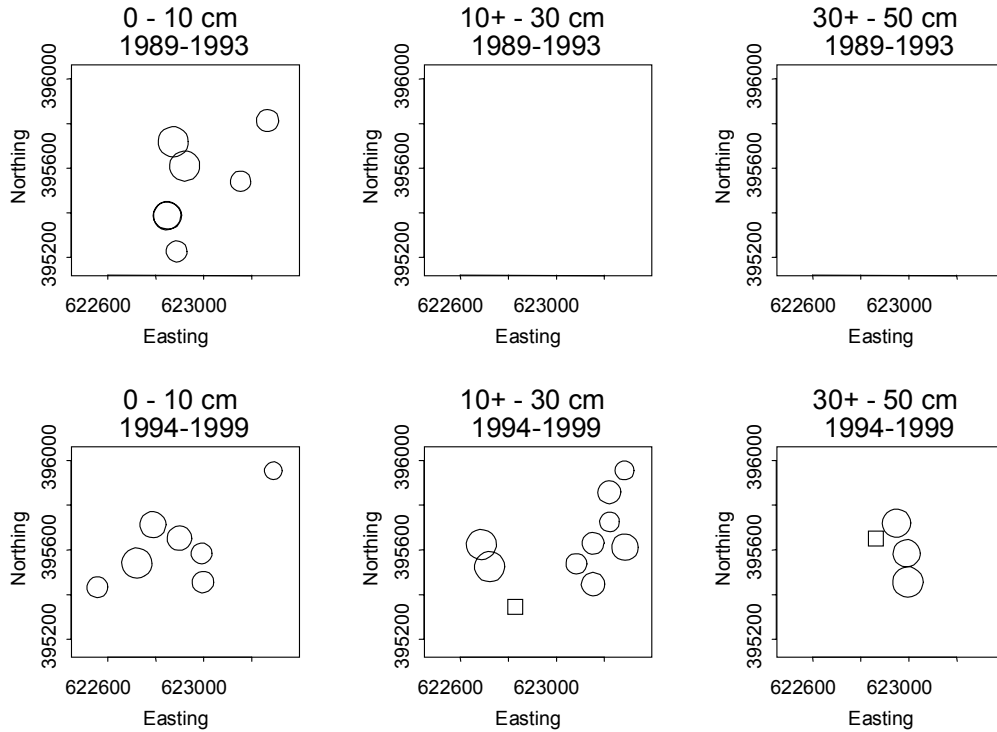


Figure A-6 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group POG (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

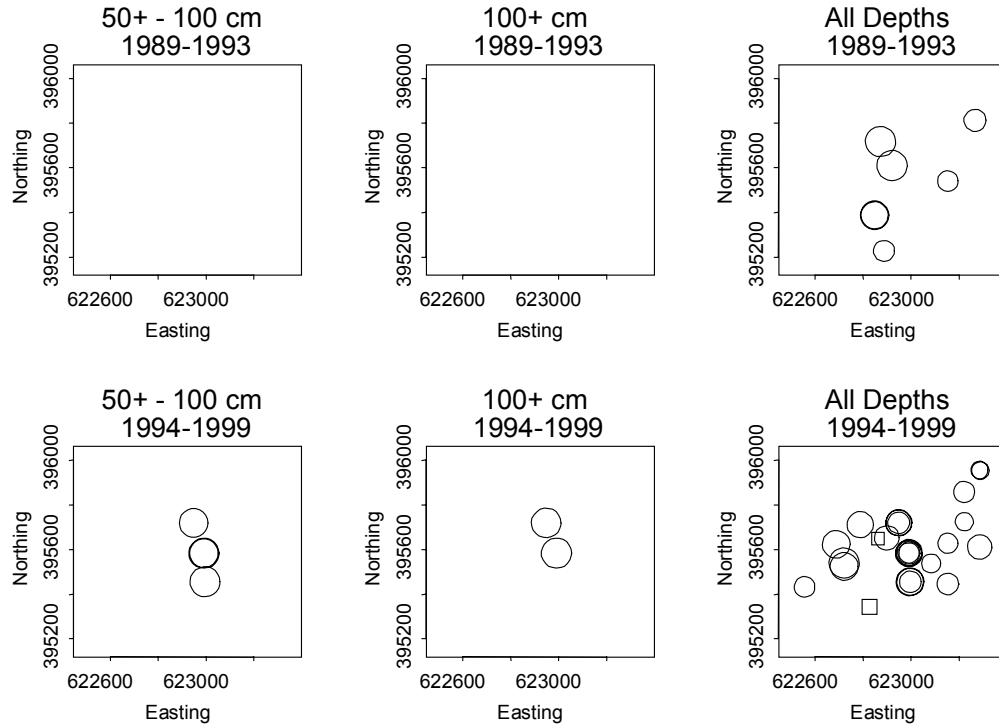


Figure A-7 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group POG (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

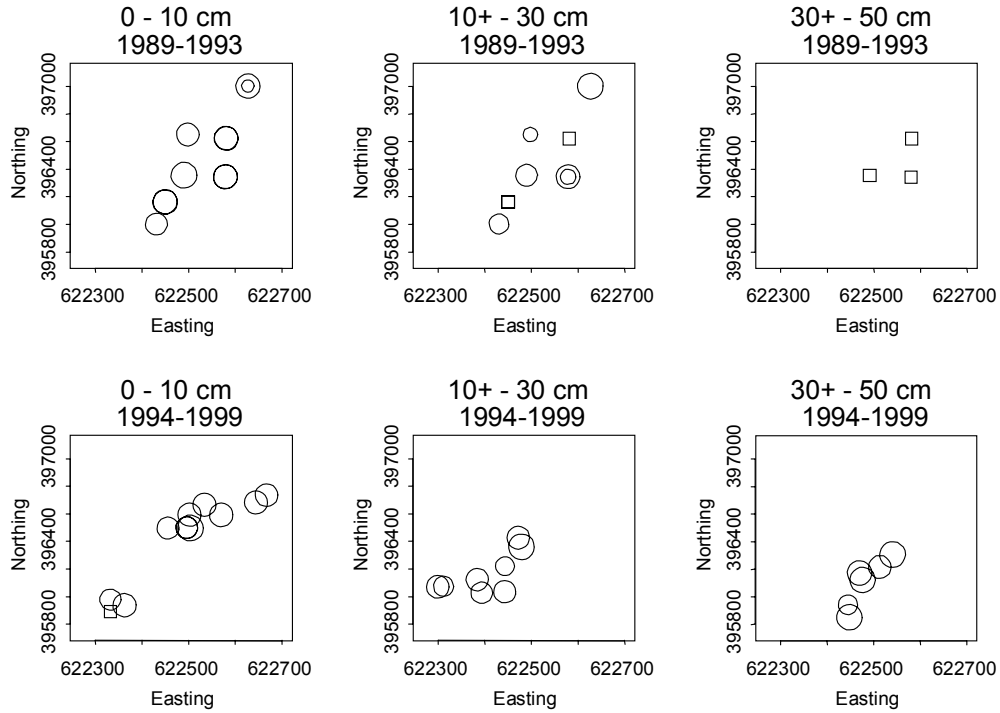


Figure A-8 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group D (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

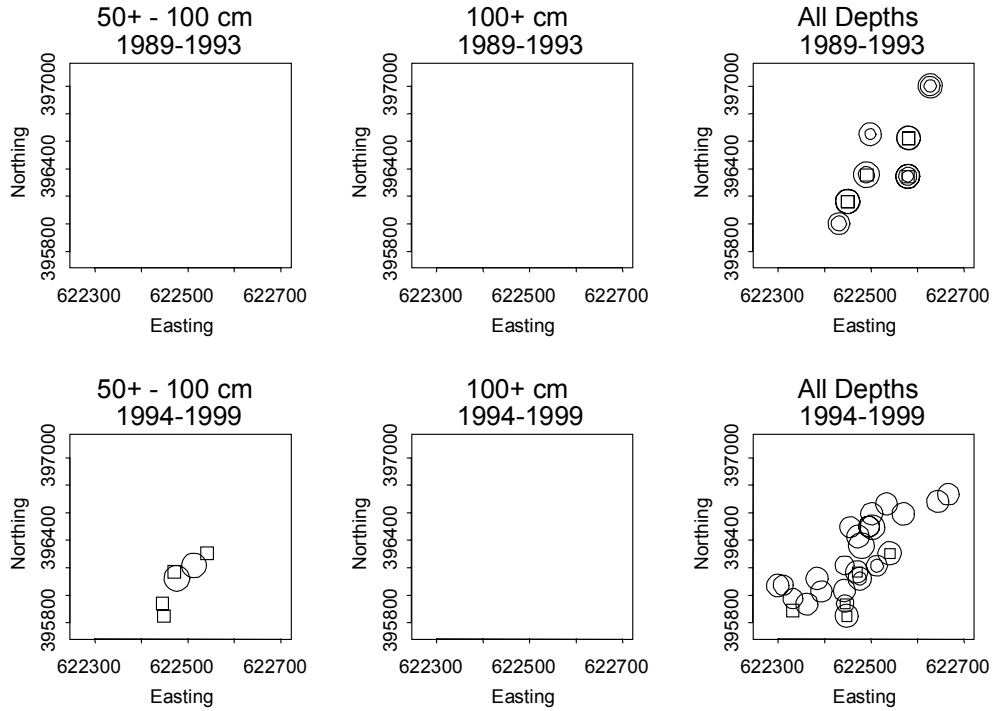


Figure A-9 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group D (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

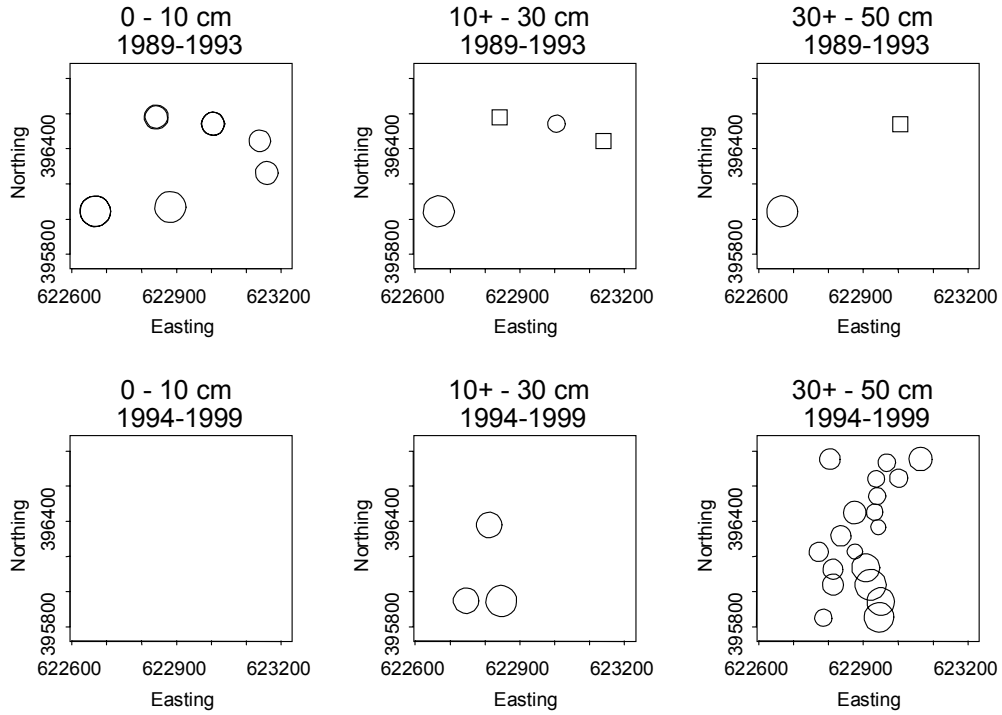


Figure A-10 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group E (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

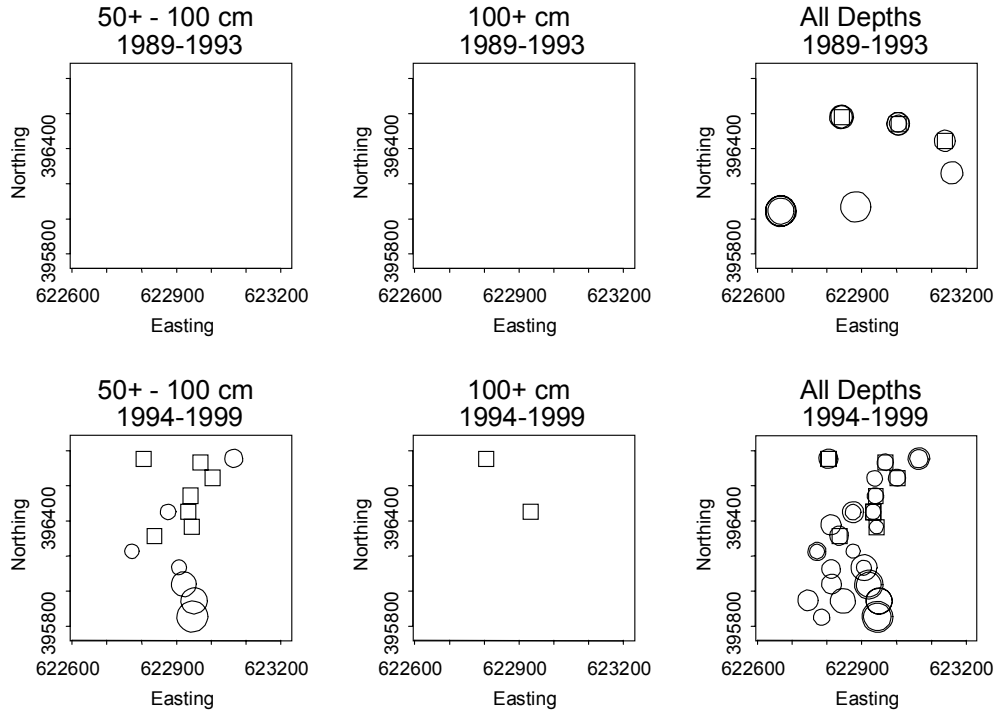


Figure A-11 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group E (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

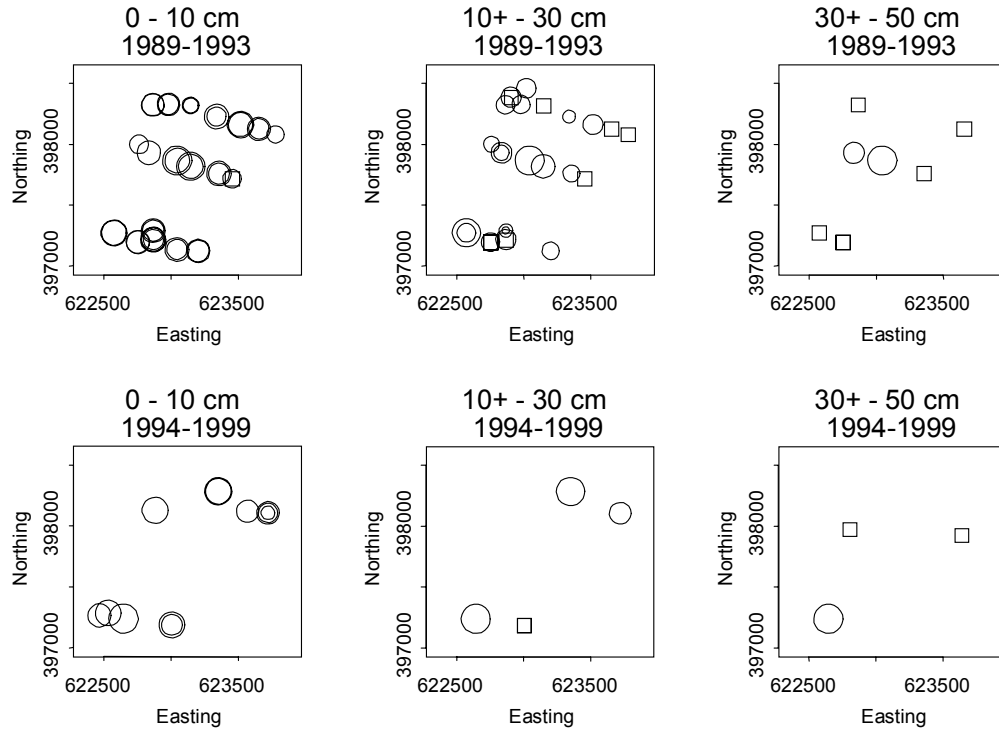


Figure A-12 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group F (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

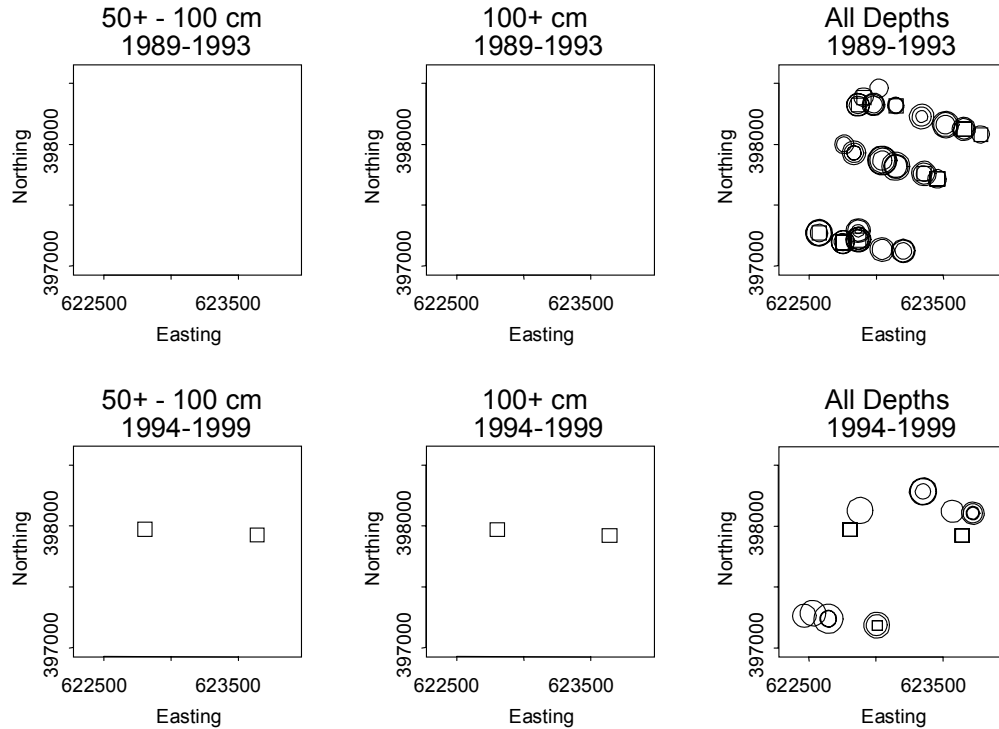


Figure A-13 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group F (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

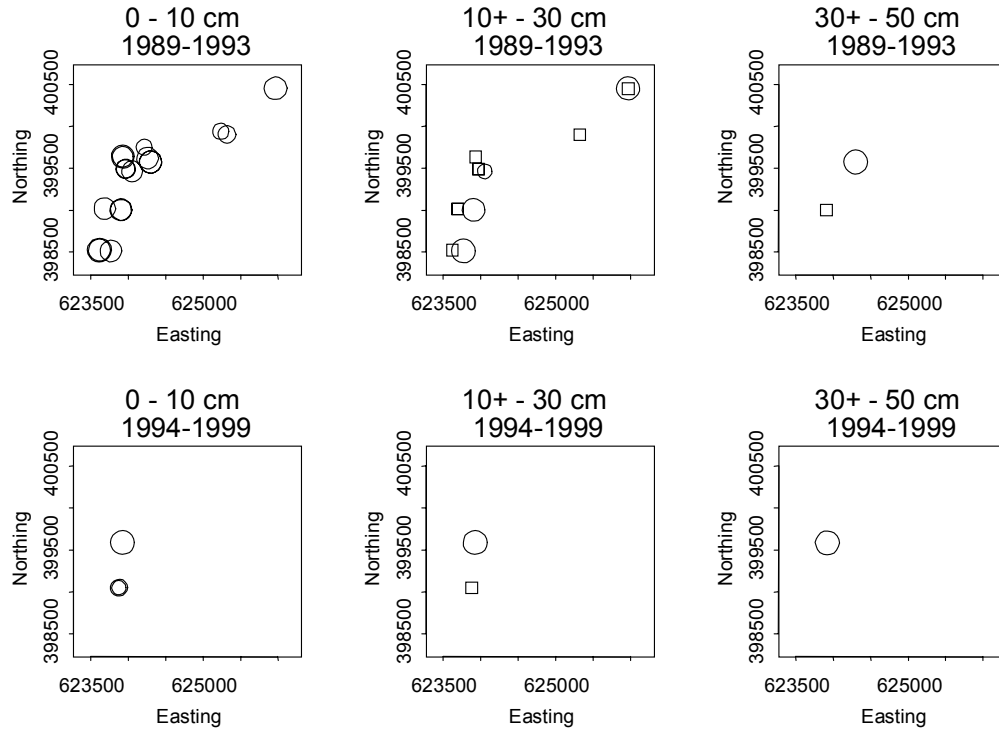


Figure A-14 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit Group GH (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

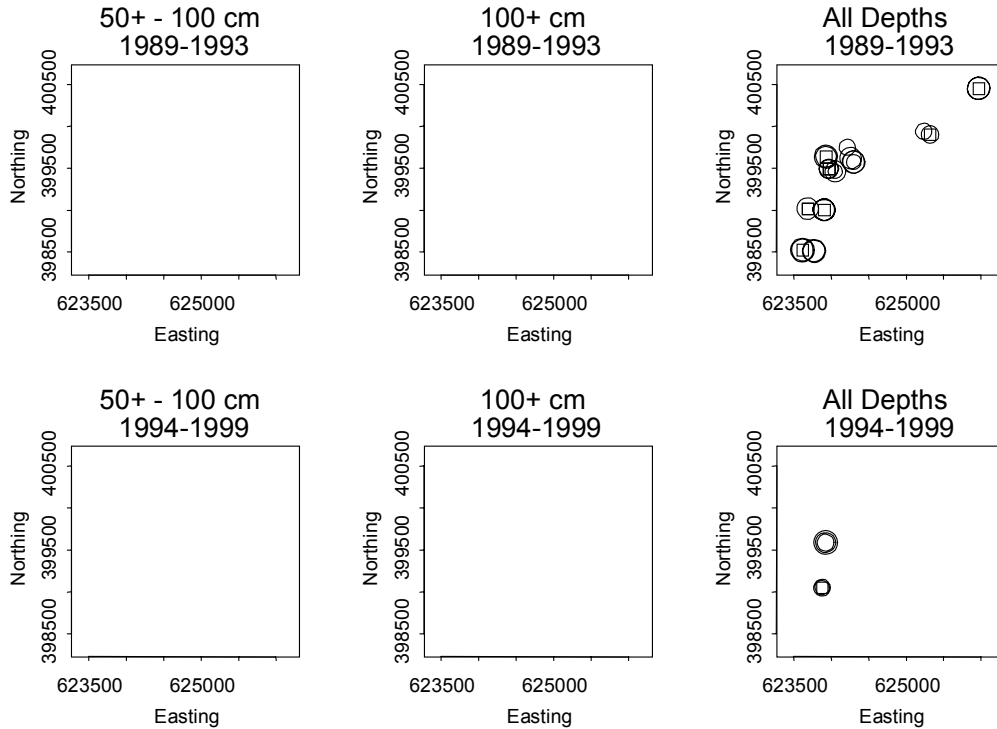


Figure A-15 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Lake Butte des Morts Deposit GH (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

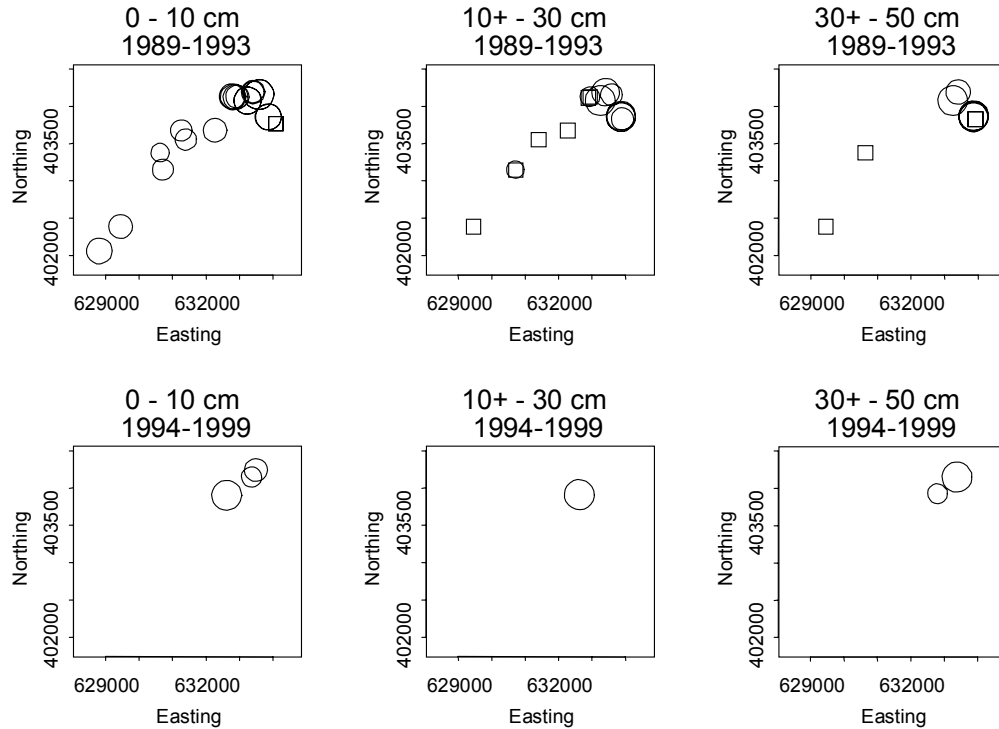


Figure A-16 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group IMOR (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

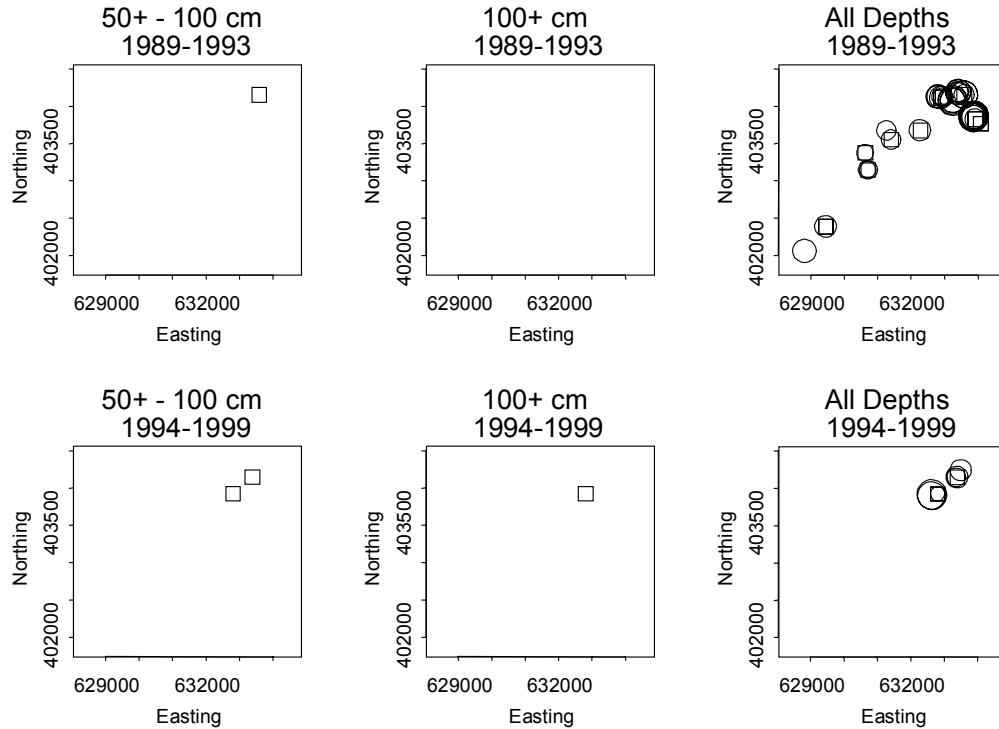


Figure A-17 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group IMOR (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

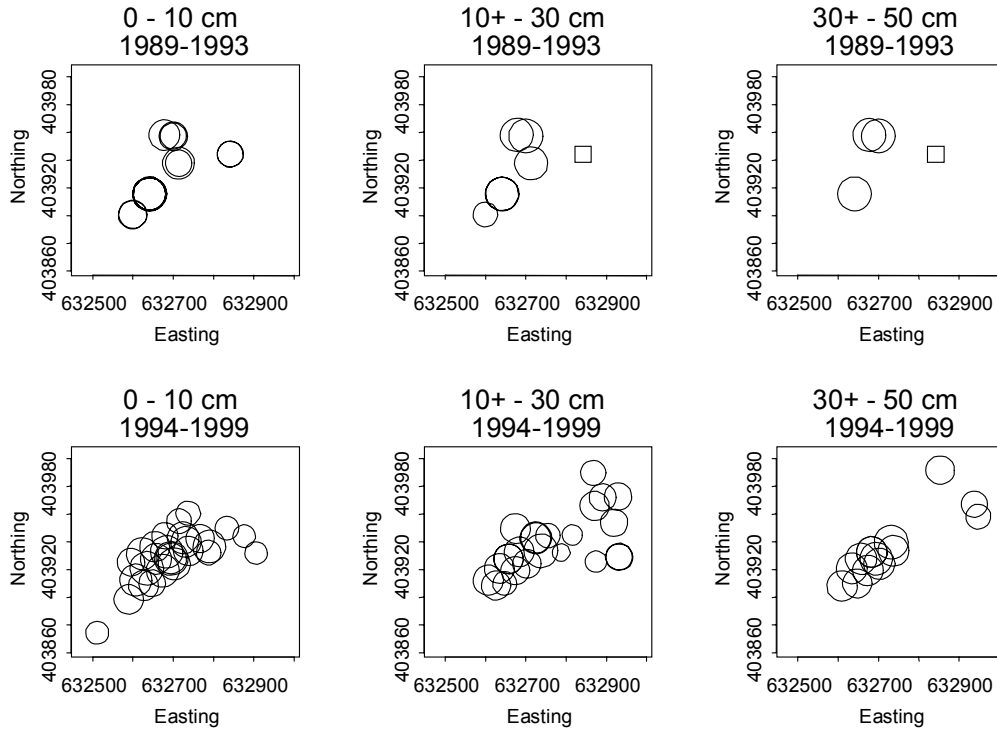


Figure A-18 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group N Before Demonstration Project (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

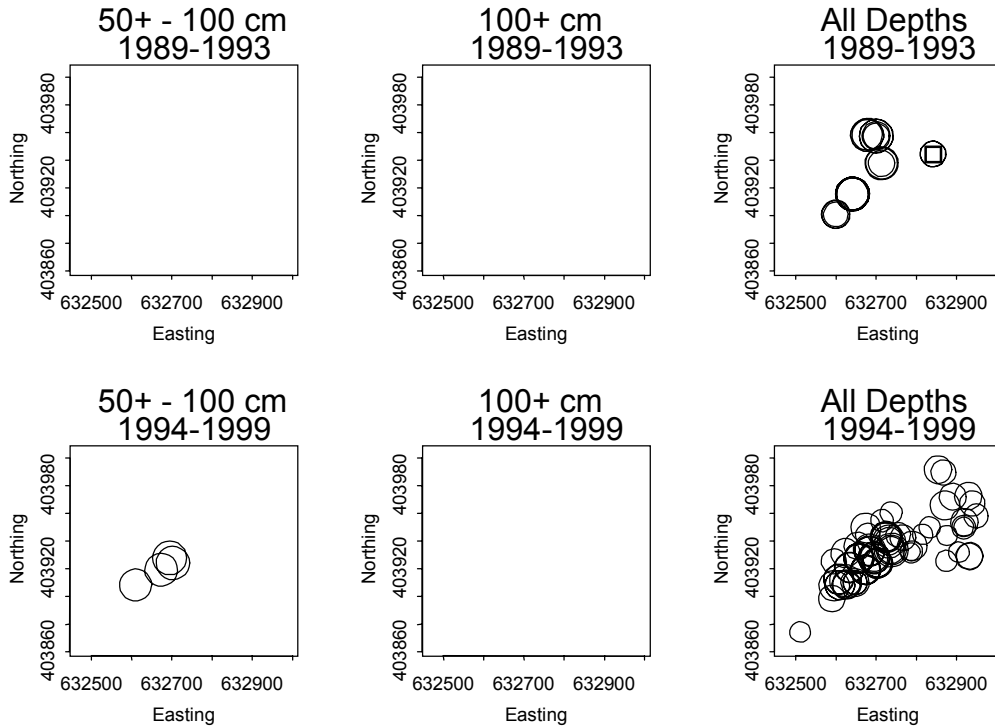


Figure A-19 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group N Before Demonstration Project (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

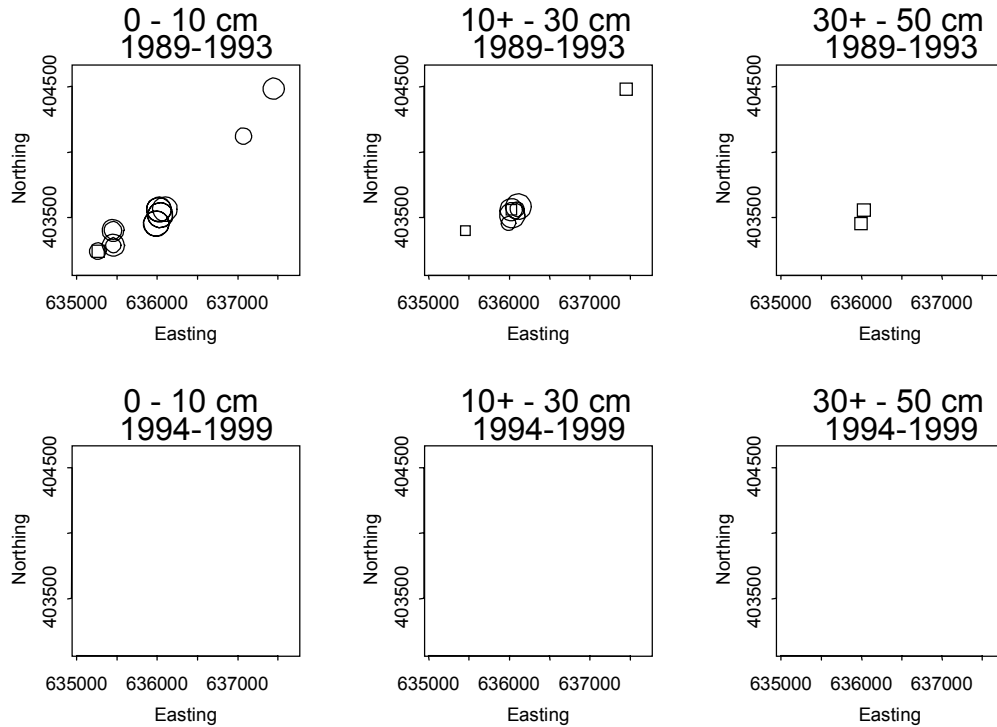


Figure A-20 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group SU (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

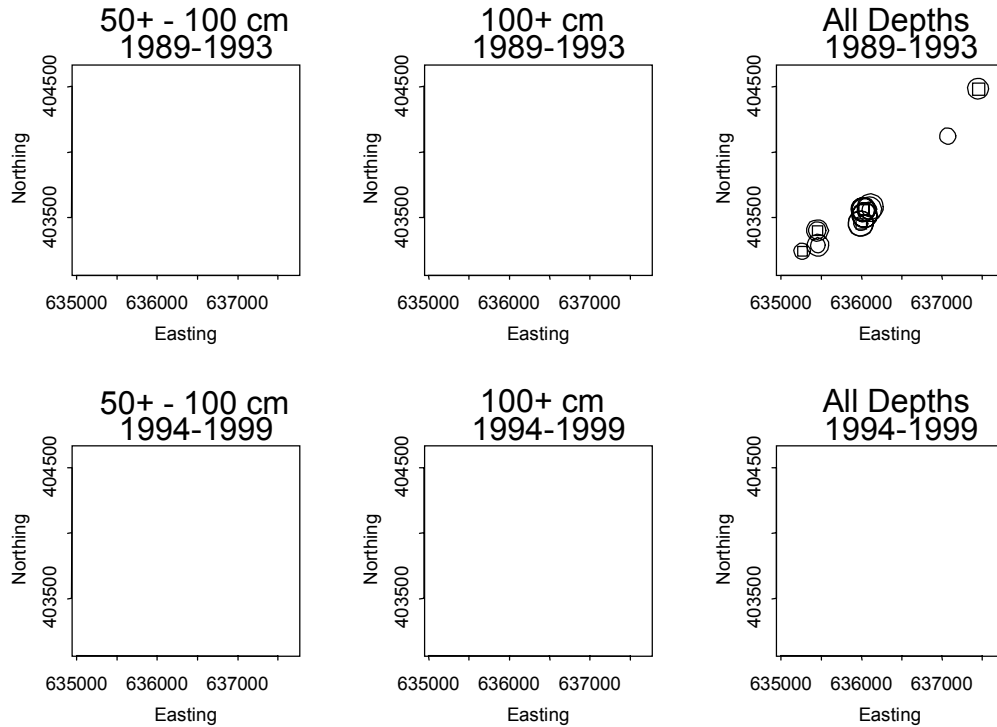


Figure A-21 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group SU (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

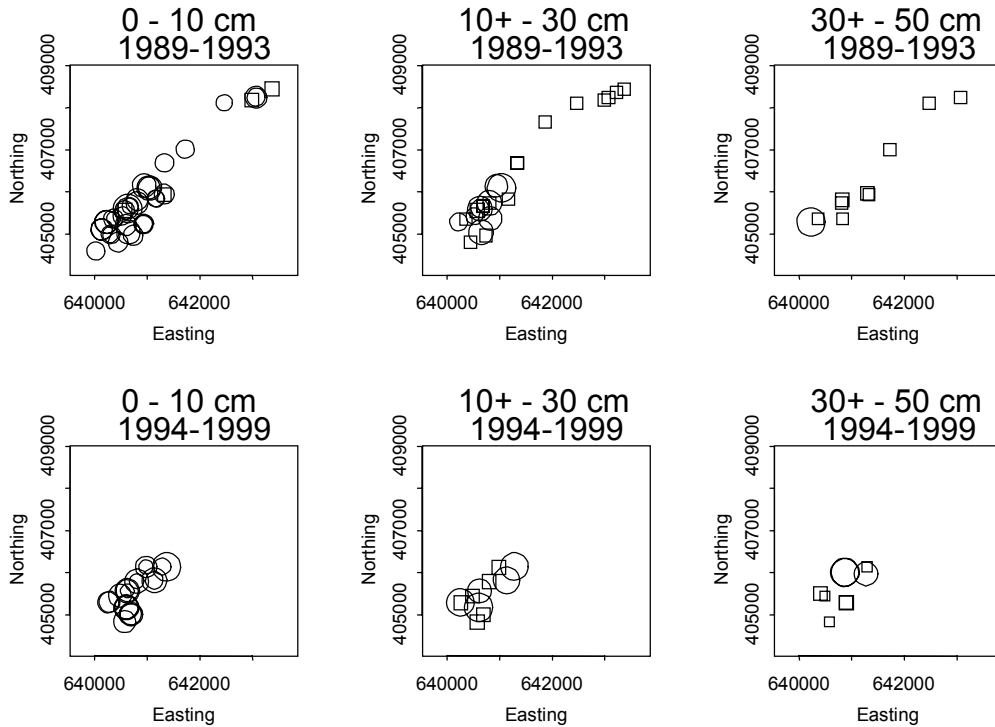


Figure A-22 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group VCC (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

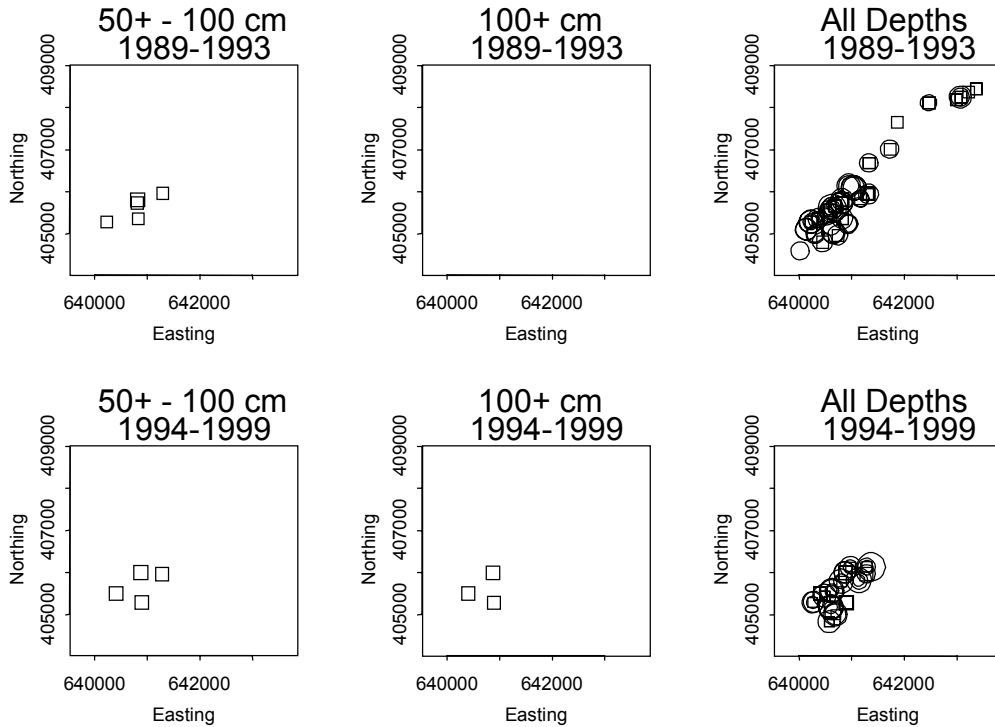


Figure A-23 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group VCC (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

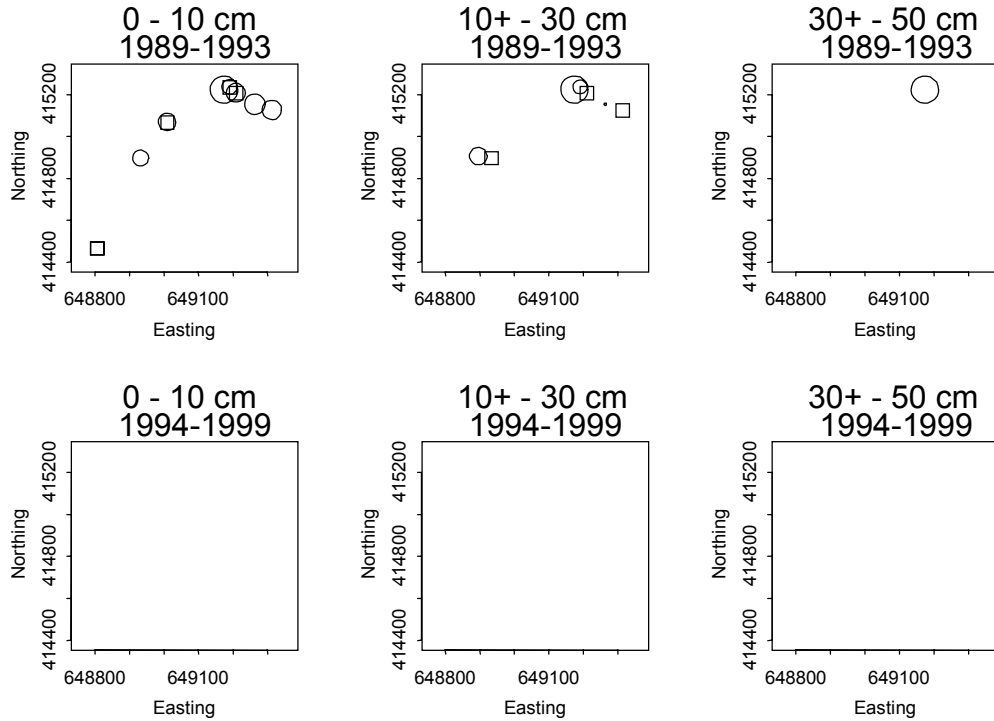


Figure A-24 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group DD (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

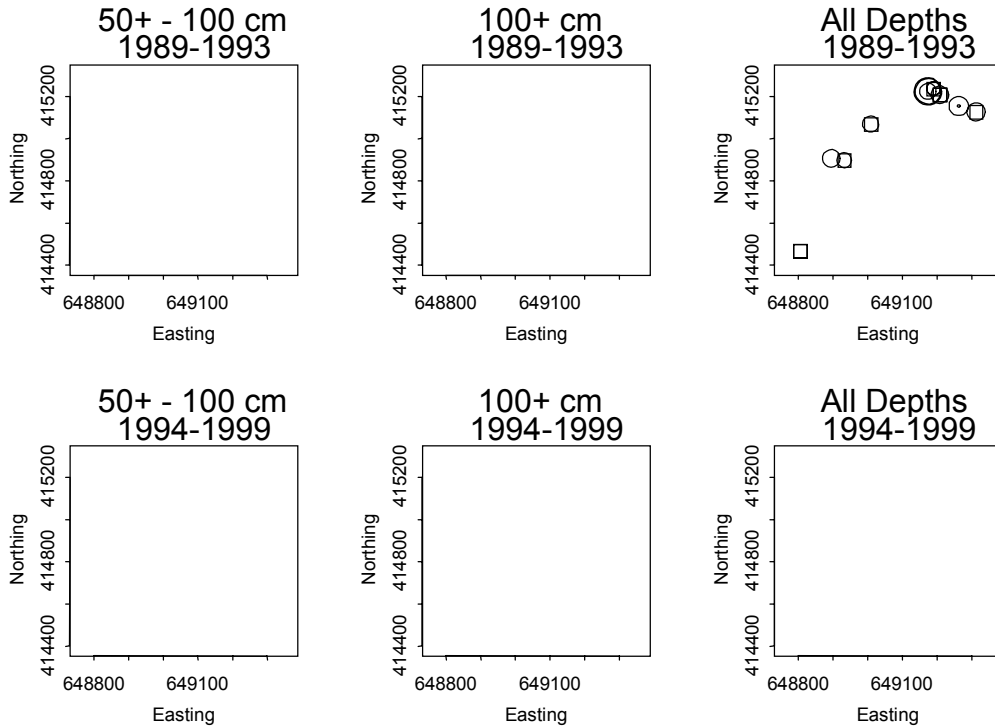


Figure A-25 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Appleton Deposit Group DD (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

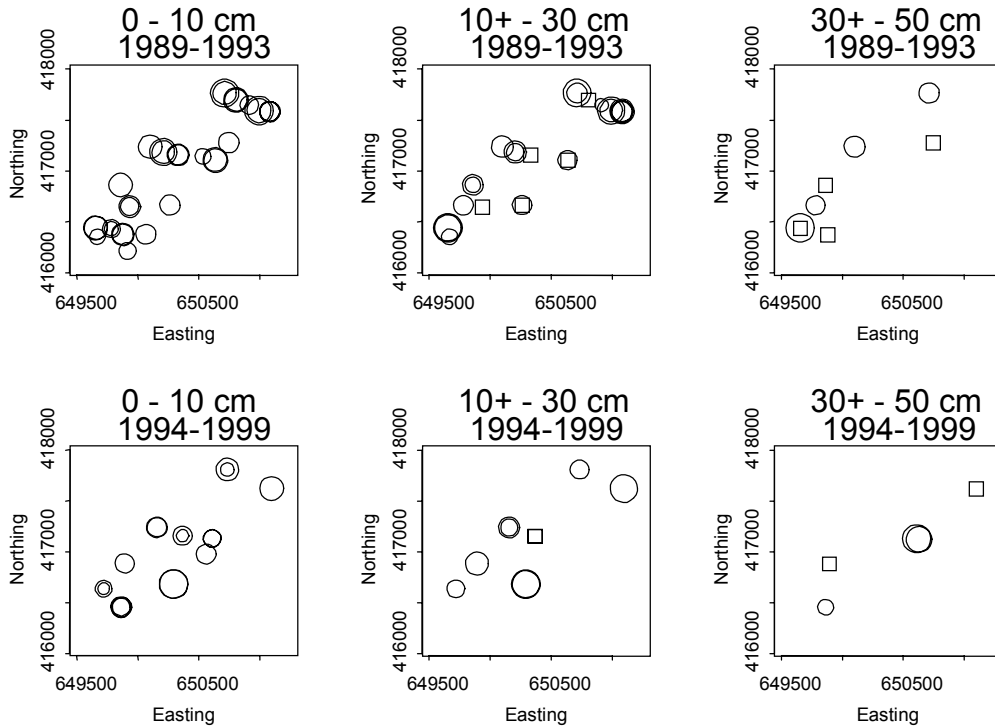


Figure A-26 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Rapids Deposit Group Upper EE (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

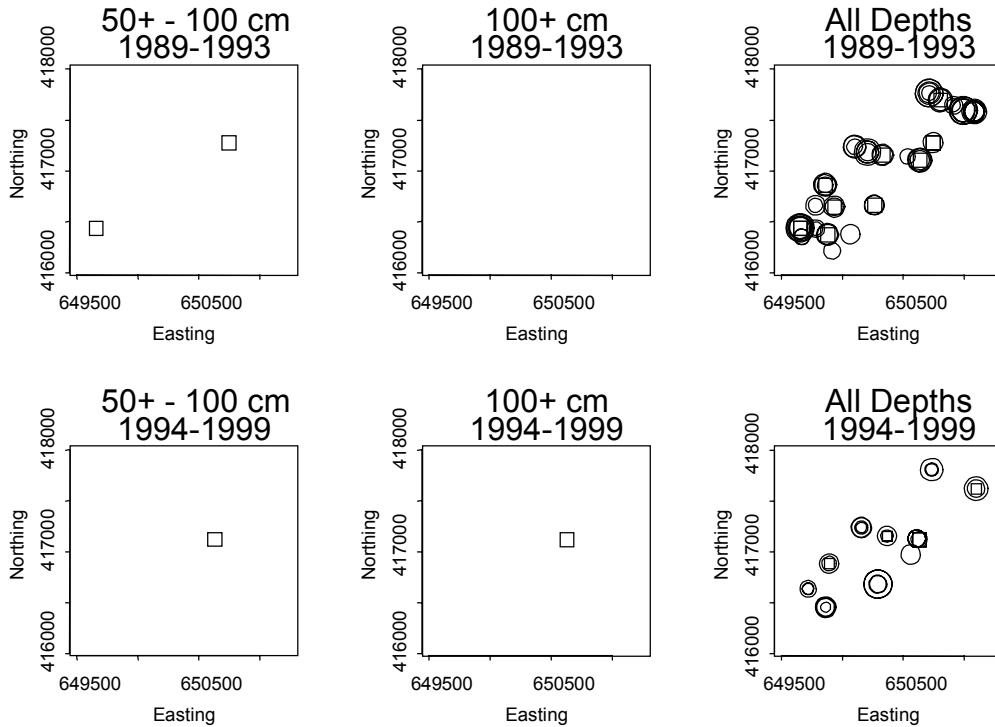


Figure A-27 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Rapids Deposit Group Upper EE (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

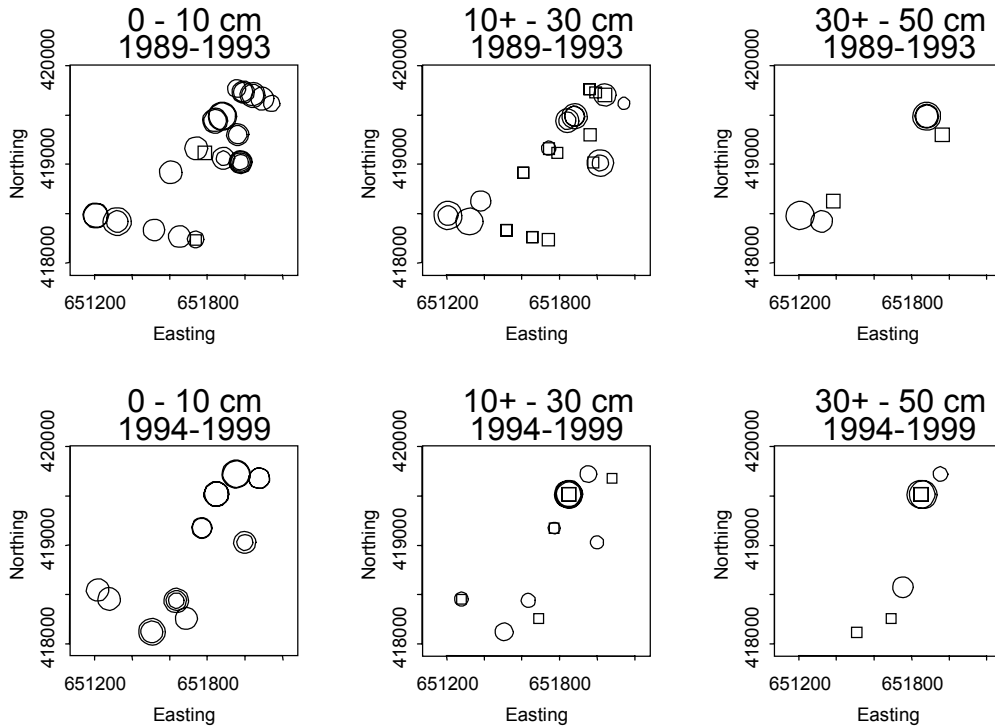


Figure A-28 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Rapids Deposit Group Lower EE (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

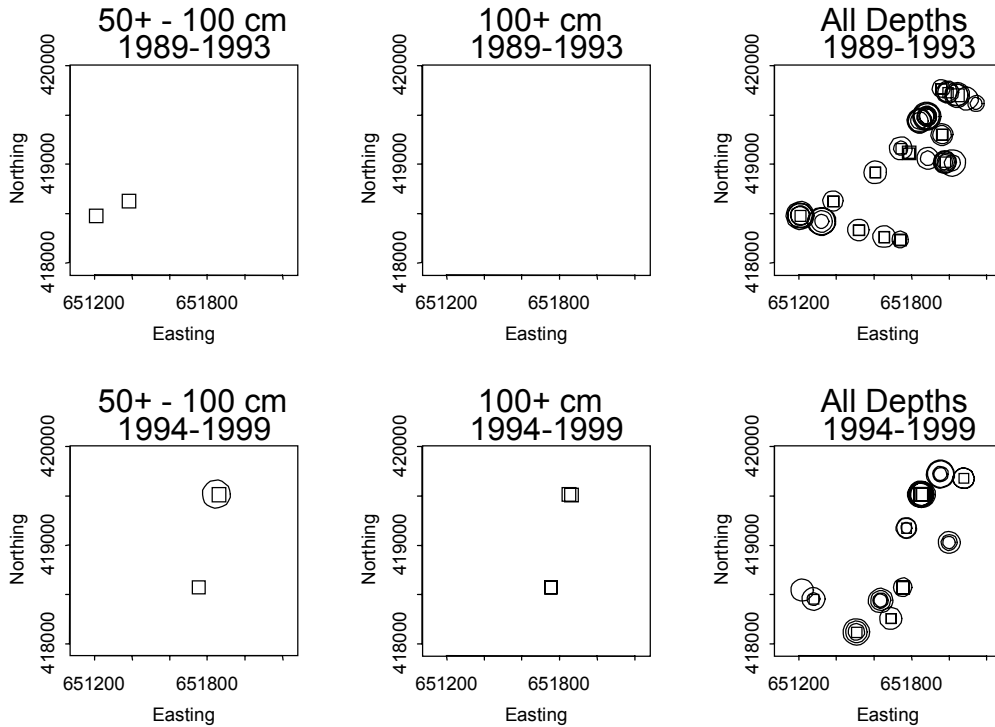


Figure A-29 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Rapids Deposit Group Lower EE (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

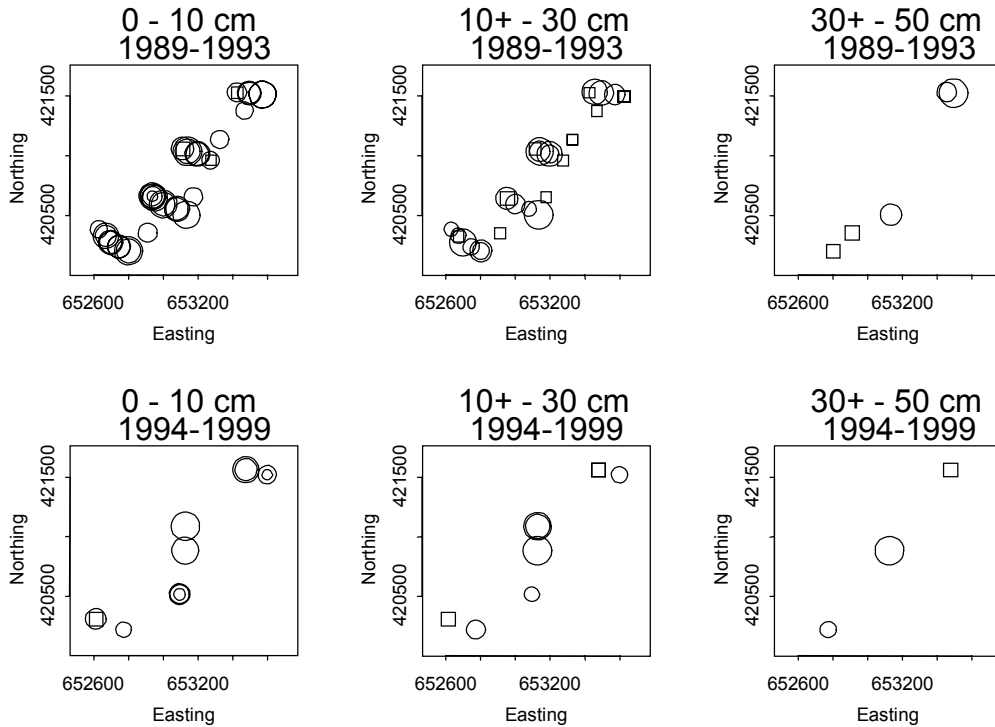


Figure A-30 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Rapids Deposit Group FF (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

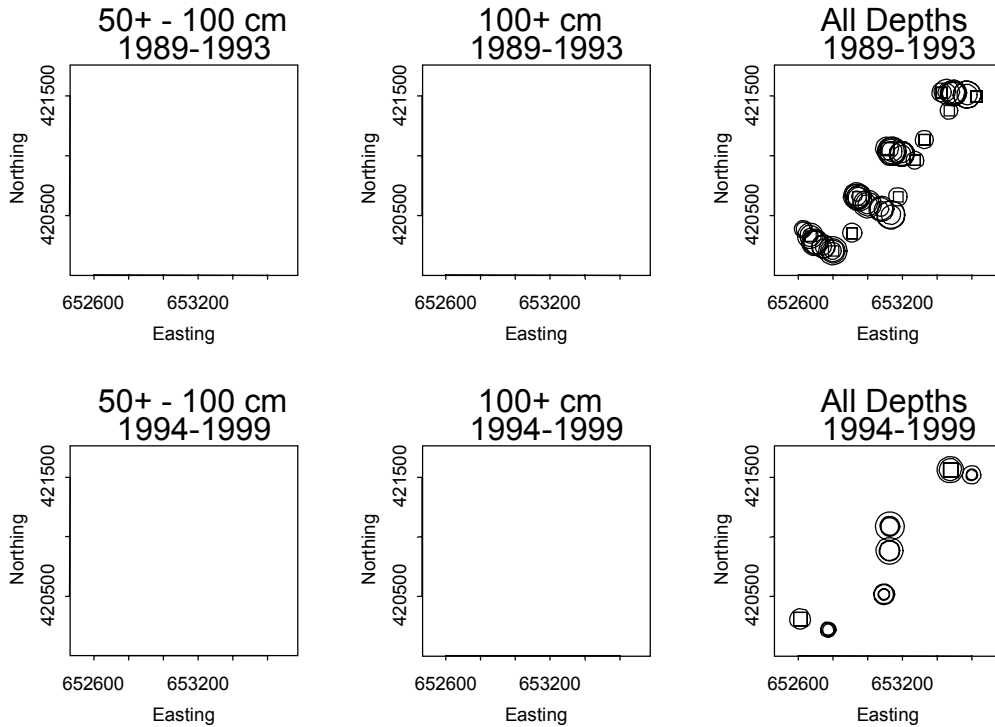


Figure A-31 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Rapids Deposit Group FF (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

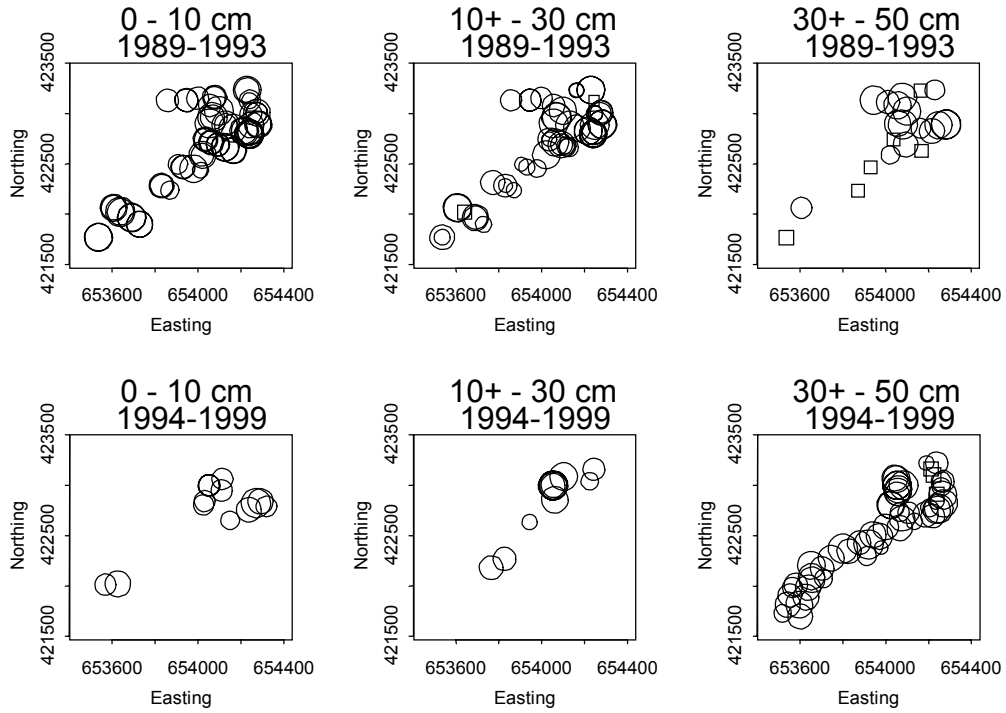


Figure A-32 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Rapids Deposit Group GGHH (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

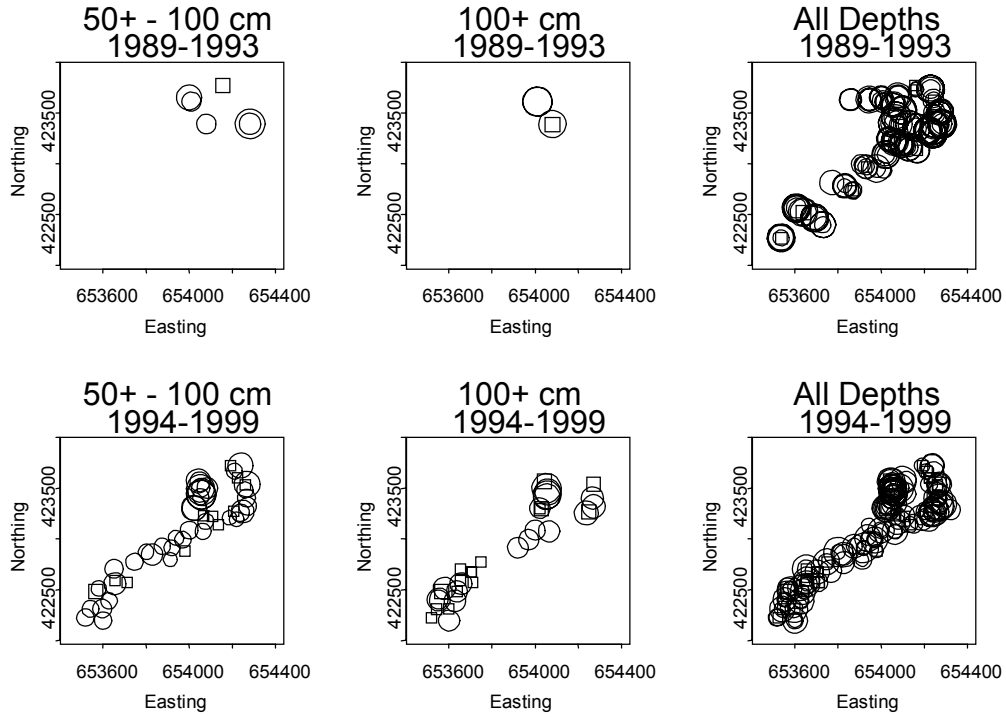


Figure A-33 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of Little Rapids Deposit Group GGHH (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

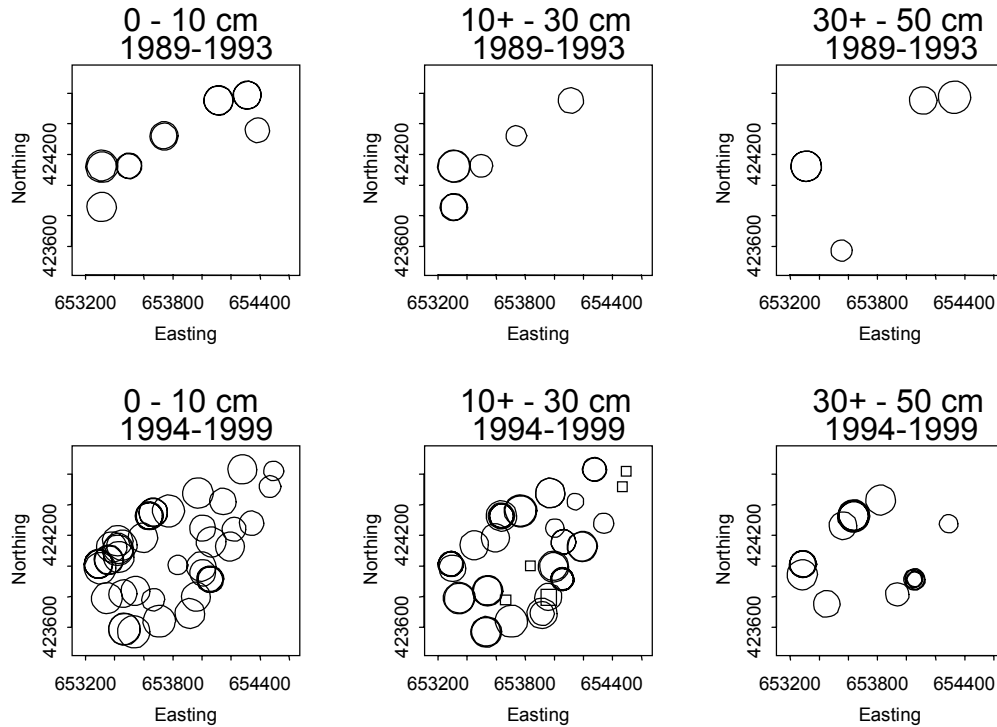


Figure A-34 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 2025 (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

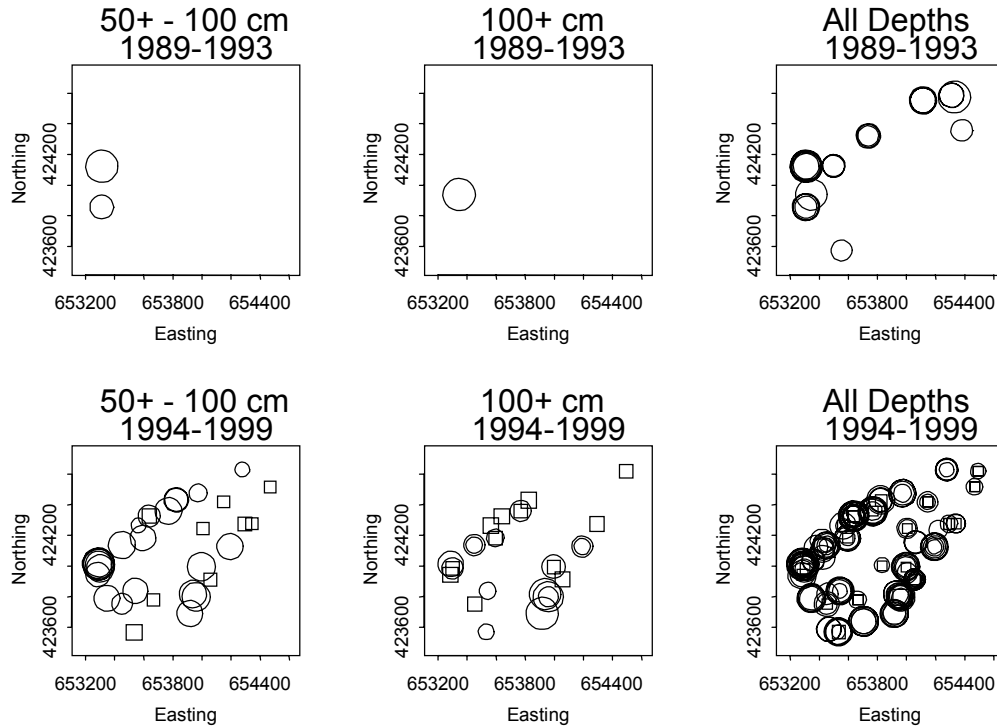


Figure A-35 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 2025 (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

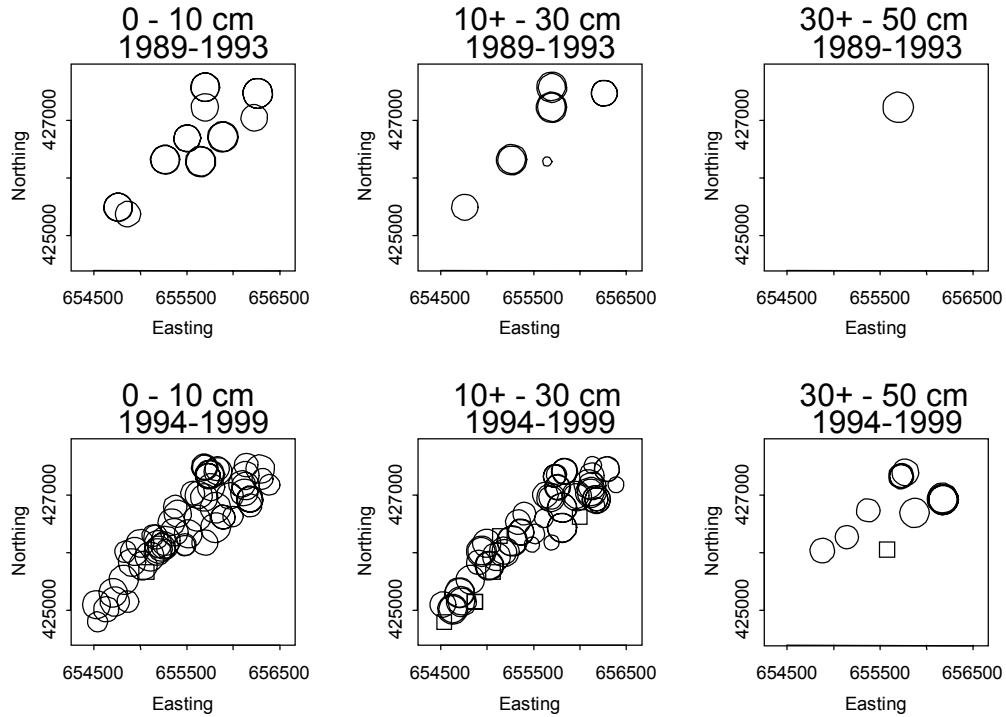


Figure A-36 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 2649 (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

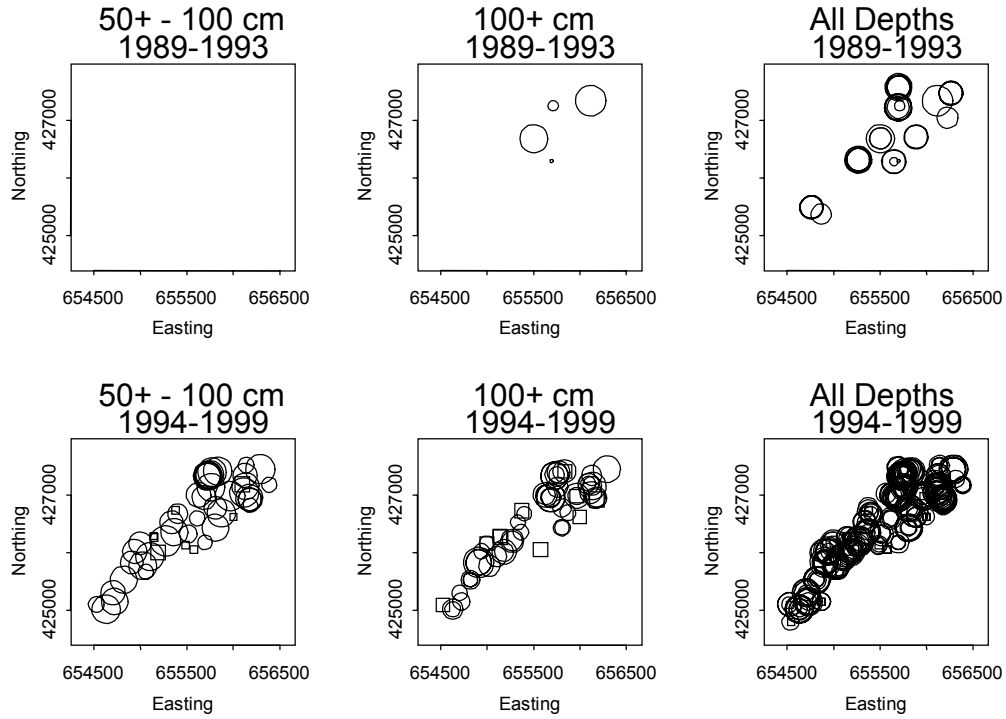


Figure A-37 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 2649 (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

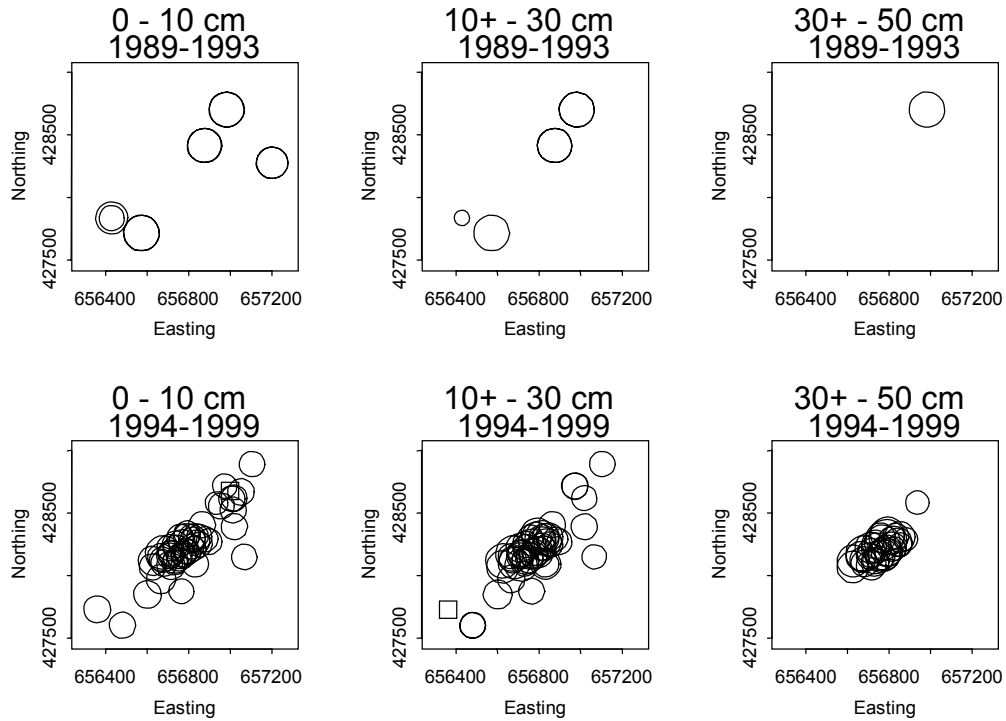


Figure A-38 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 5067 (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

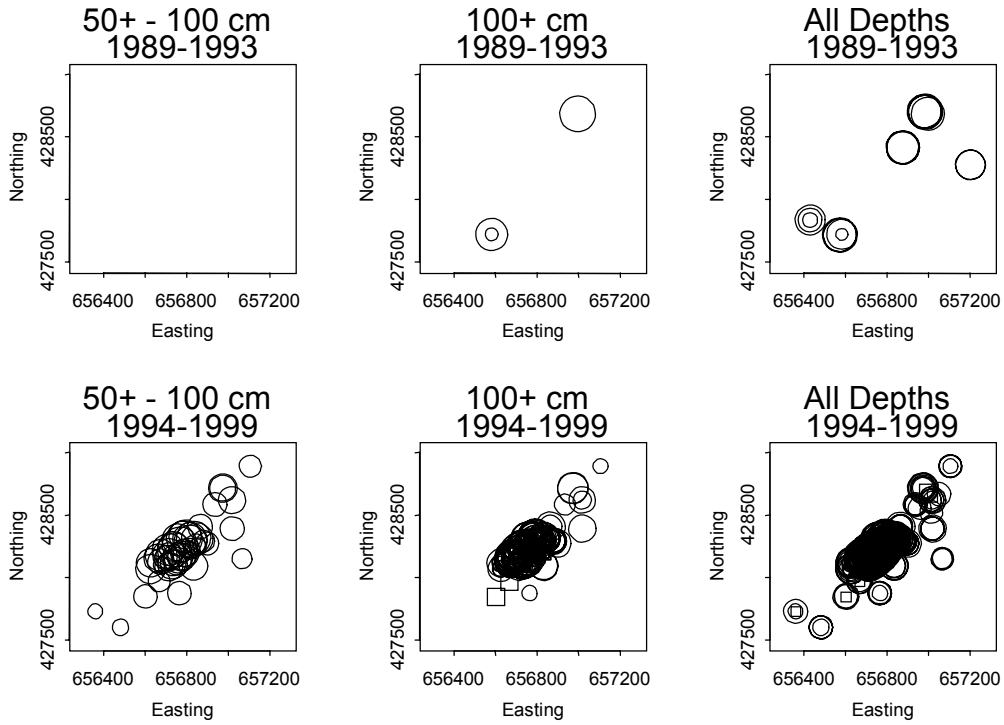


Figure A-39 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 5067 (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

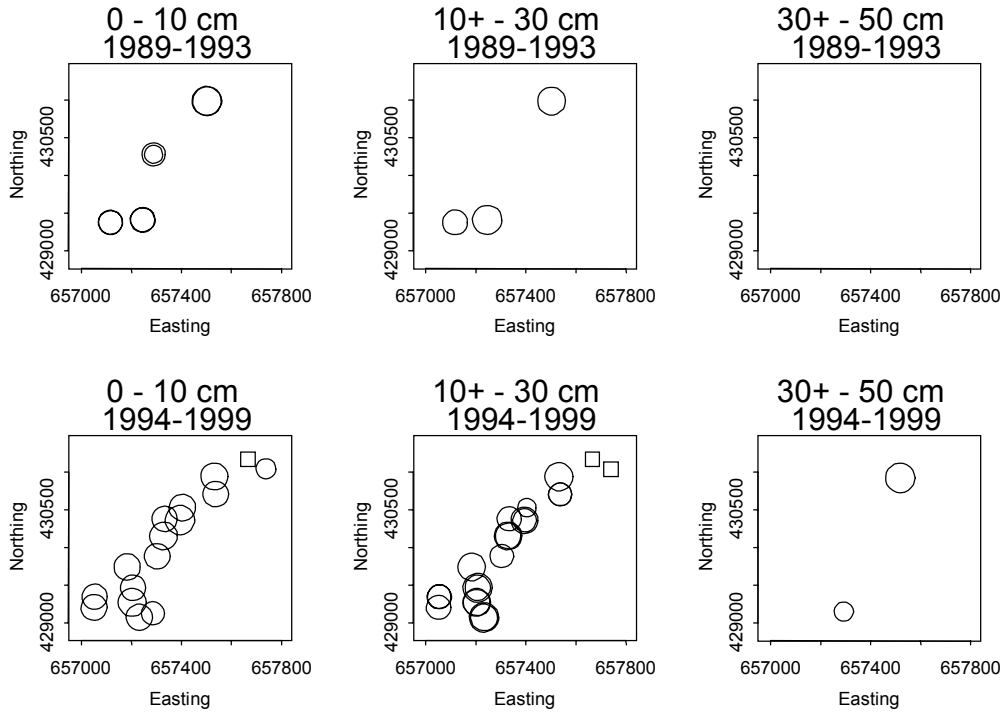


Figure A-40 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 6891 (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

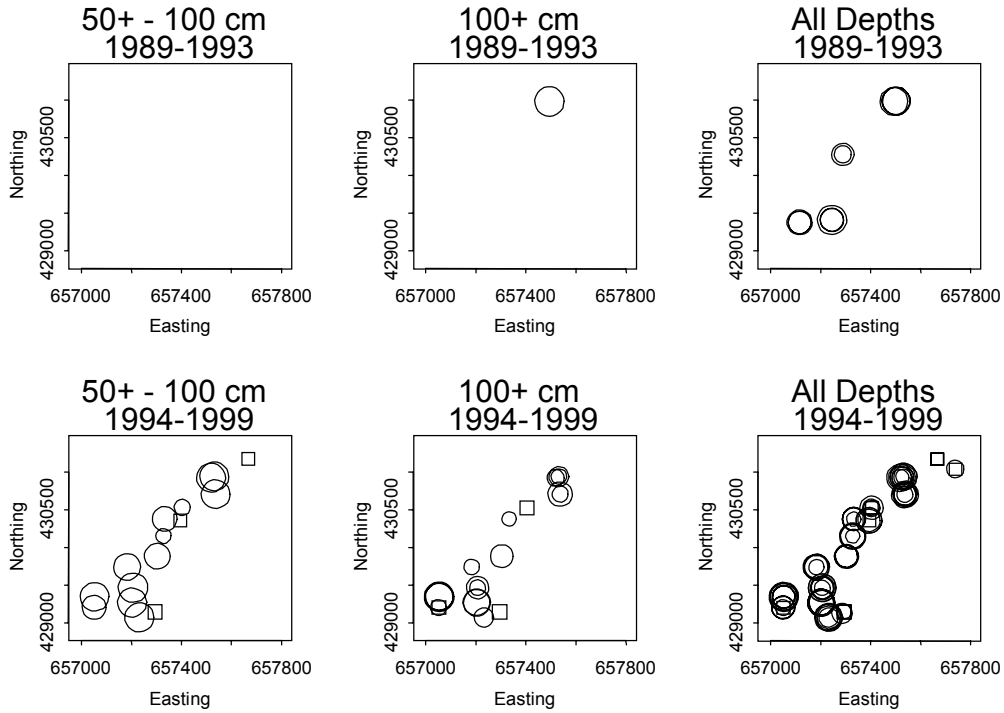


Figure A-41 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 6891 (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

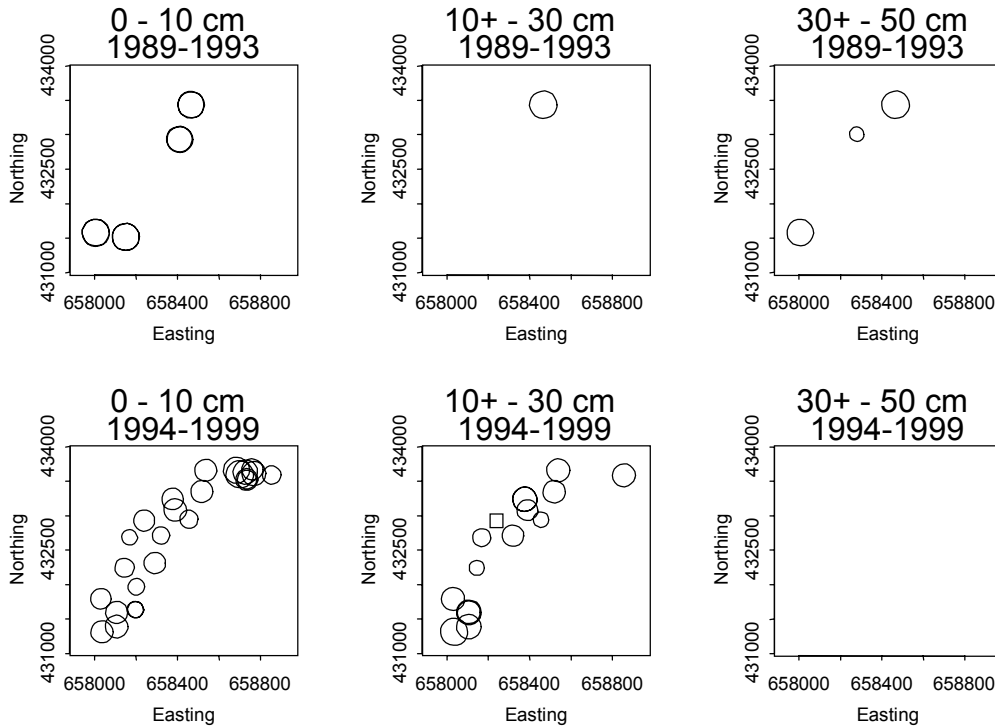


Figure A-42 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 92115 (0 to 50 cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

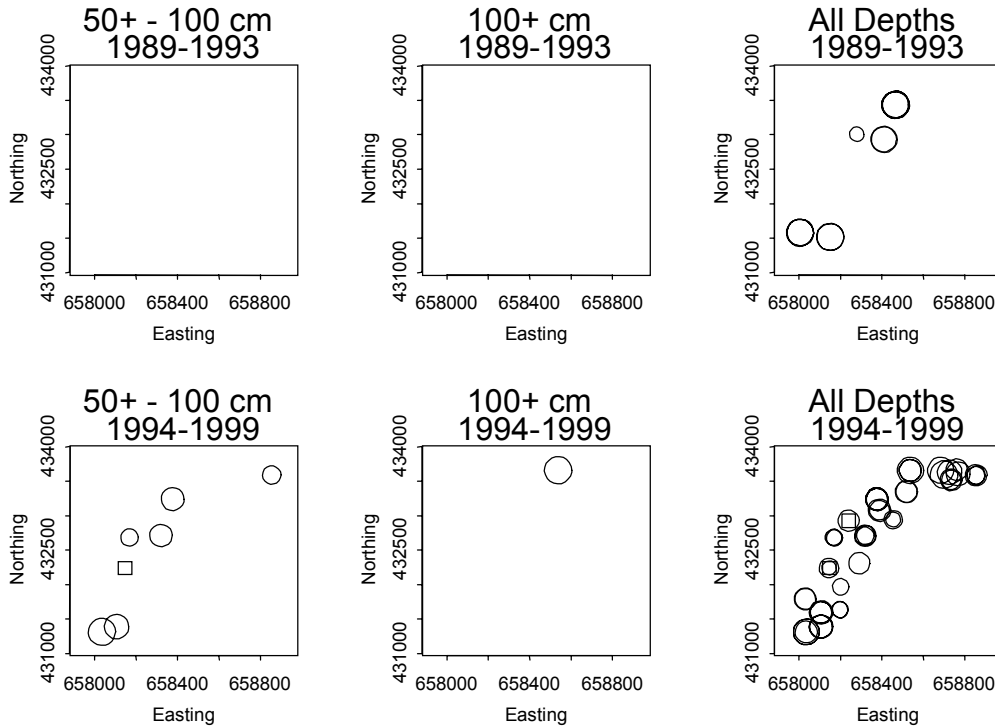


Figure A-43 Sample Locations by Northing and Easting Coordinates During 1989–1993 and 1994–1999, Depth Strata of De Pere SMU Group 92115 (50 to 100+ cm)

Larger symbols indicate higher concentrations. Circles (○) indicate measured concentrations and squares (□) indicate the detection limit of concentrations below the detection limit. Coordinates are in meters.

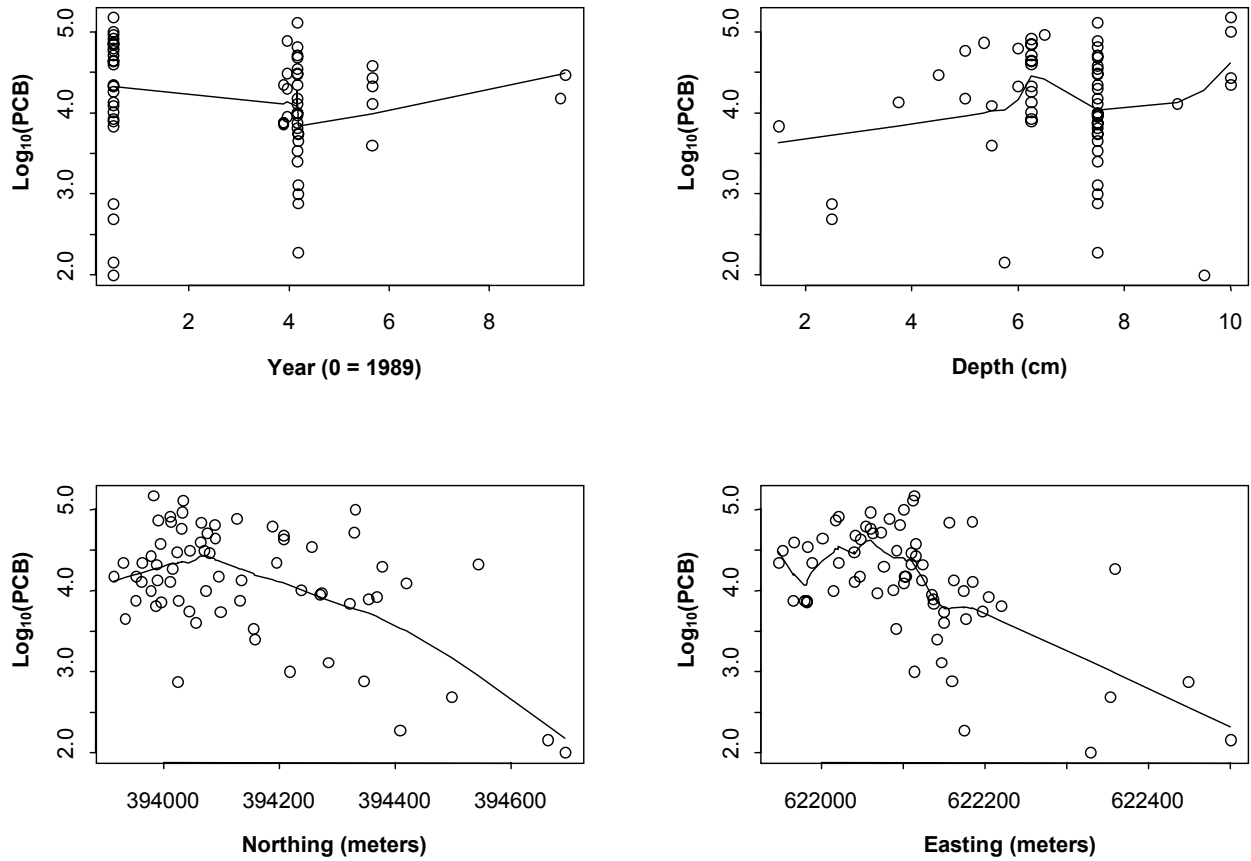


Figure A-44 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group AB (0 to 10 cm) Including Fitted Smoothed Line

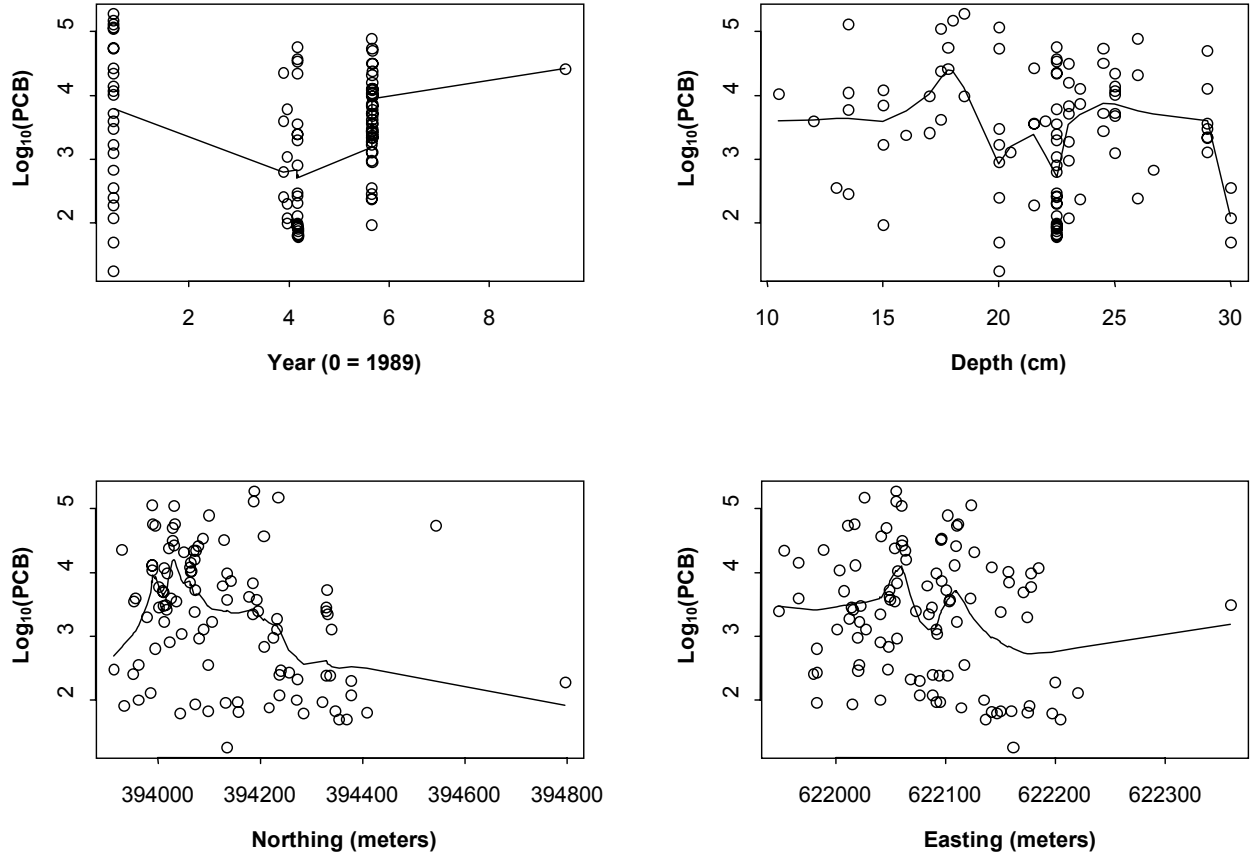


Figure A-45 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group AB (10 to 30 cm) Including Fitted Smoothed Line

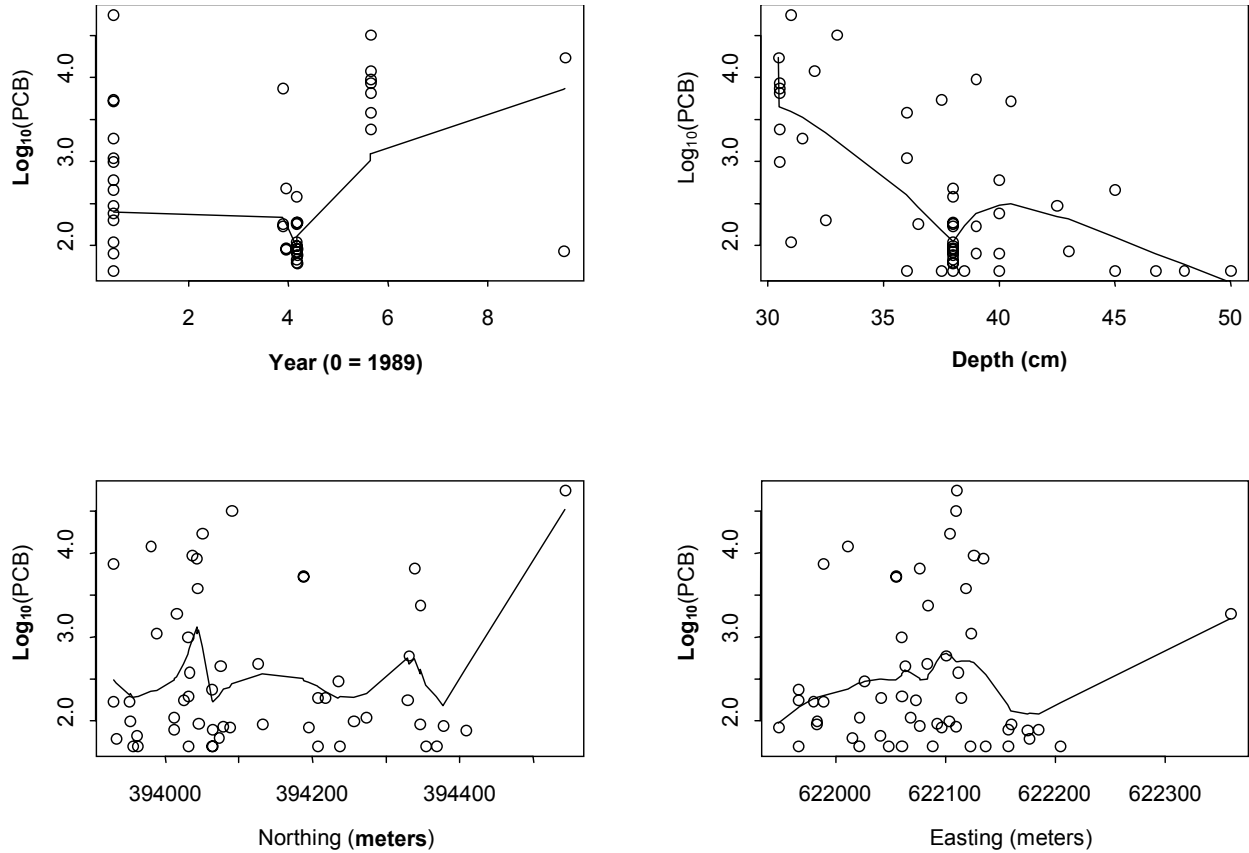


Figure A-46 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group AB (30 to 50 cm) Including Fitted Smoothed Line

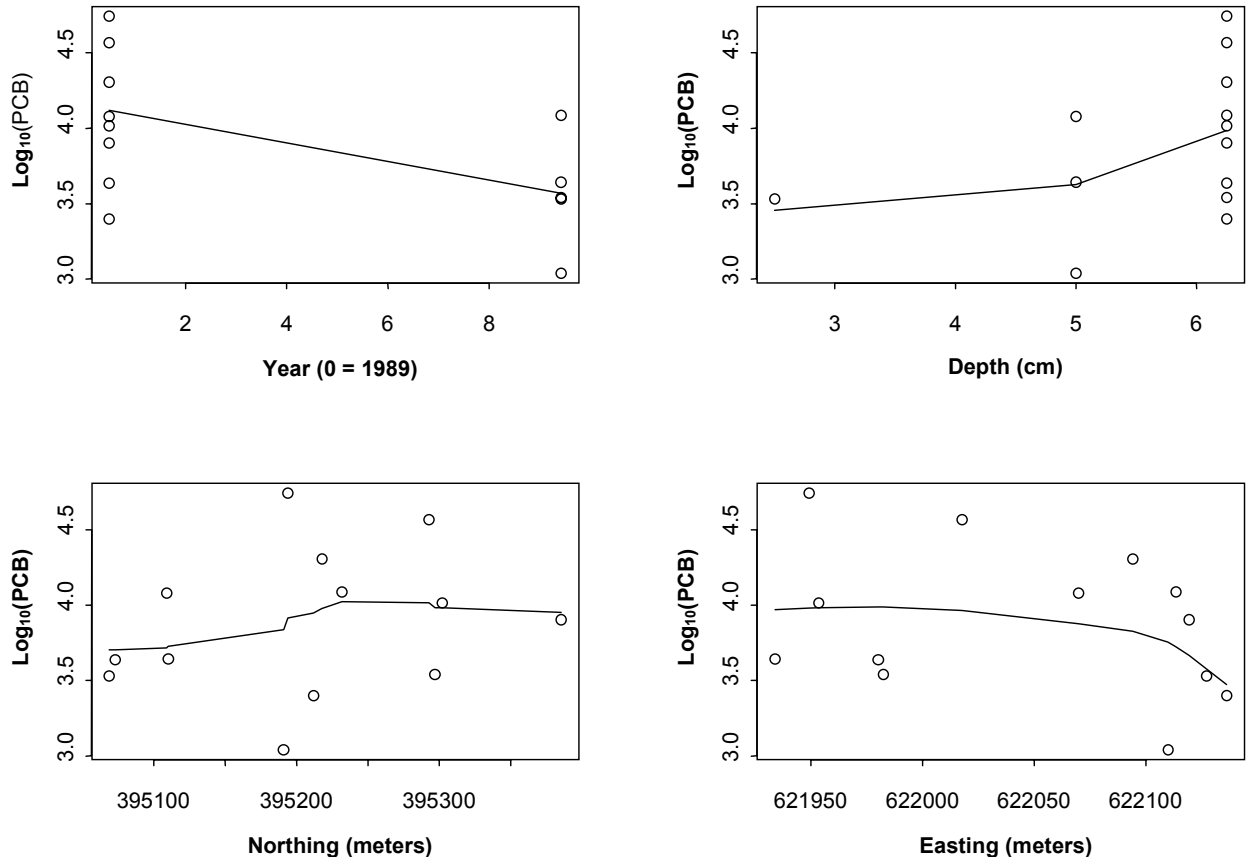


Figure A-47 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group C (0 to 10 cm) Including Fitted Smoothed Line

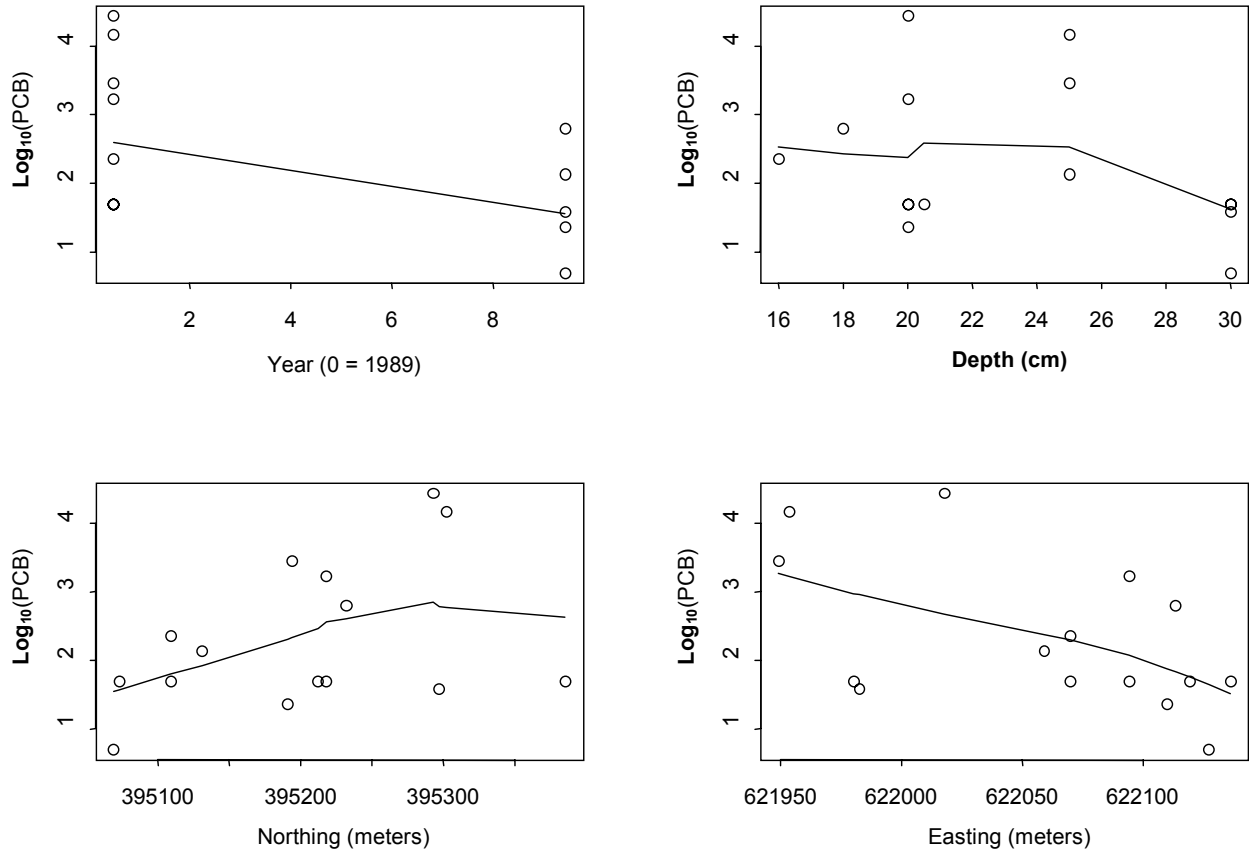


Figure A-48 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group C (10 to 30 cm) Including Fitted Smoothed Line

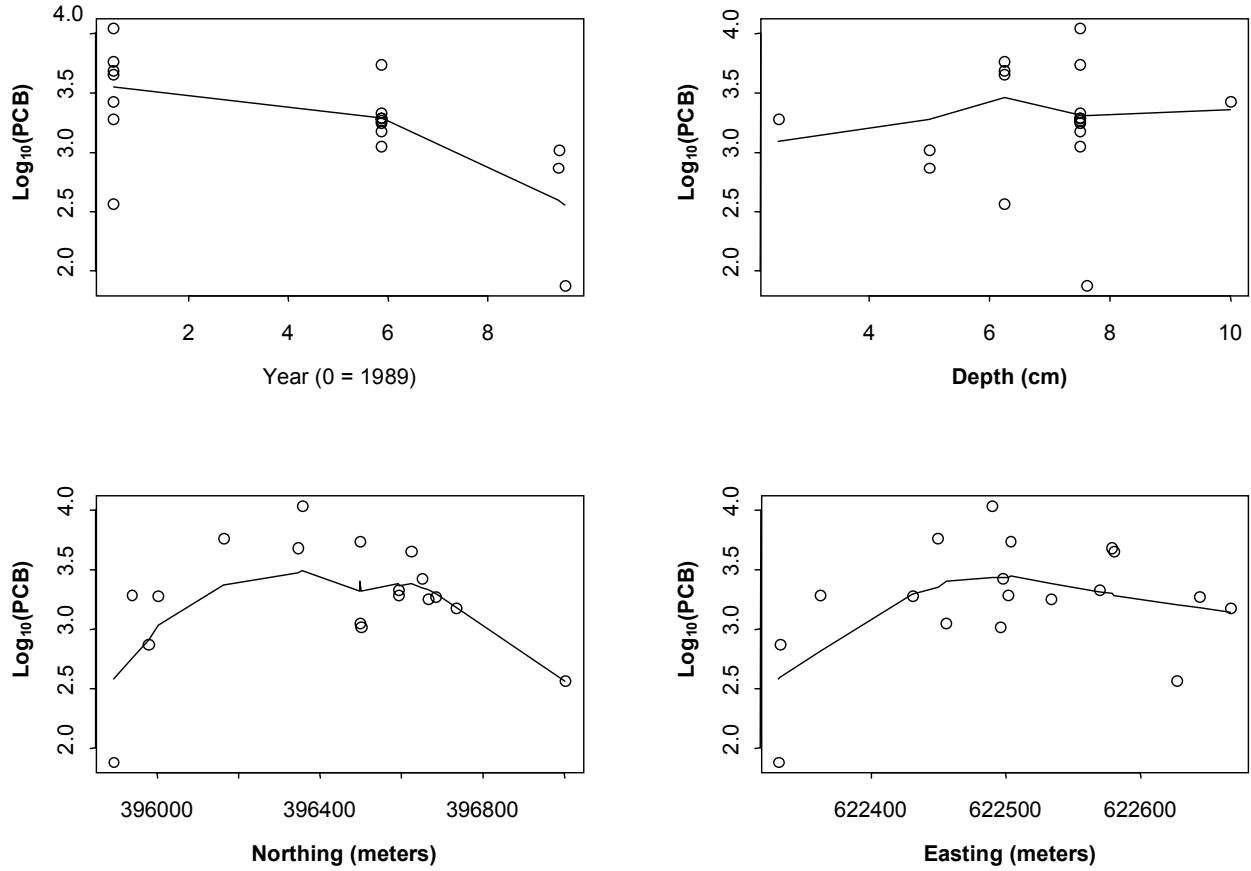


Figure A-49 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group D (0 to 10 cm) Including Fitted Smoothed Line

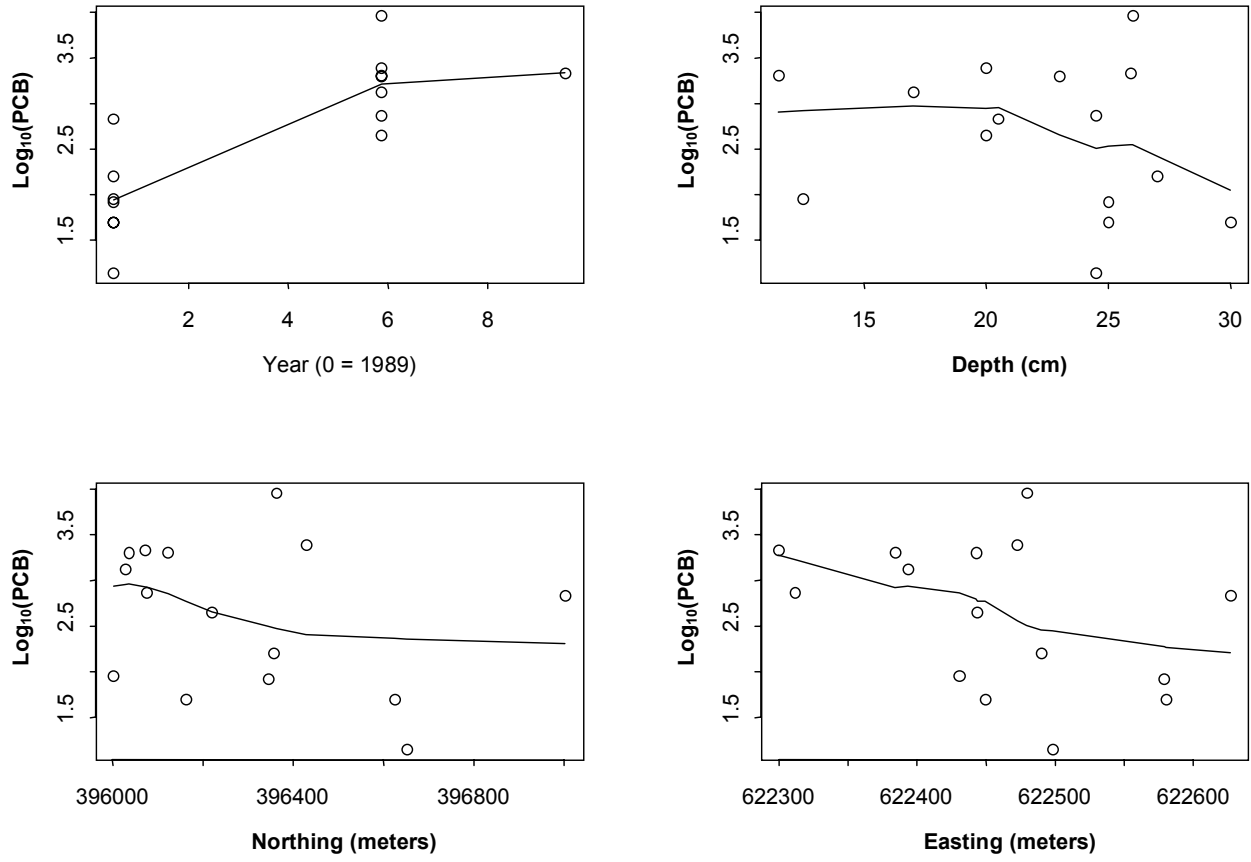


Figure A-50 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group D (10 to 30 cm) Including Fitted Smoothed Line

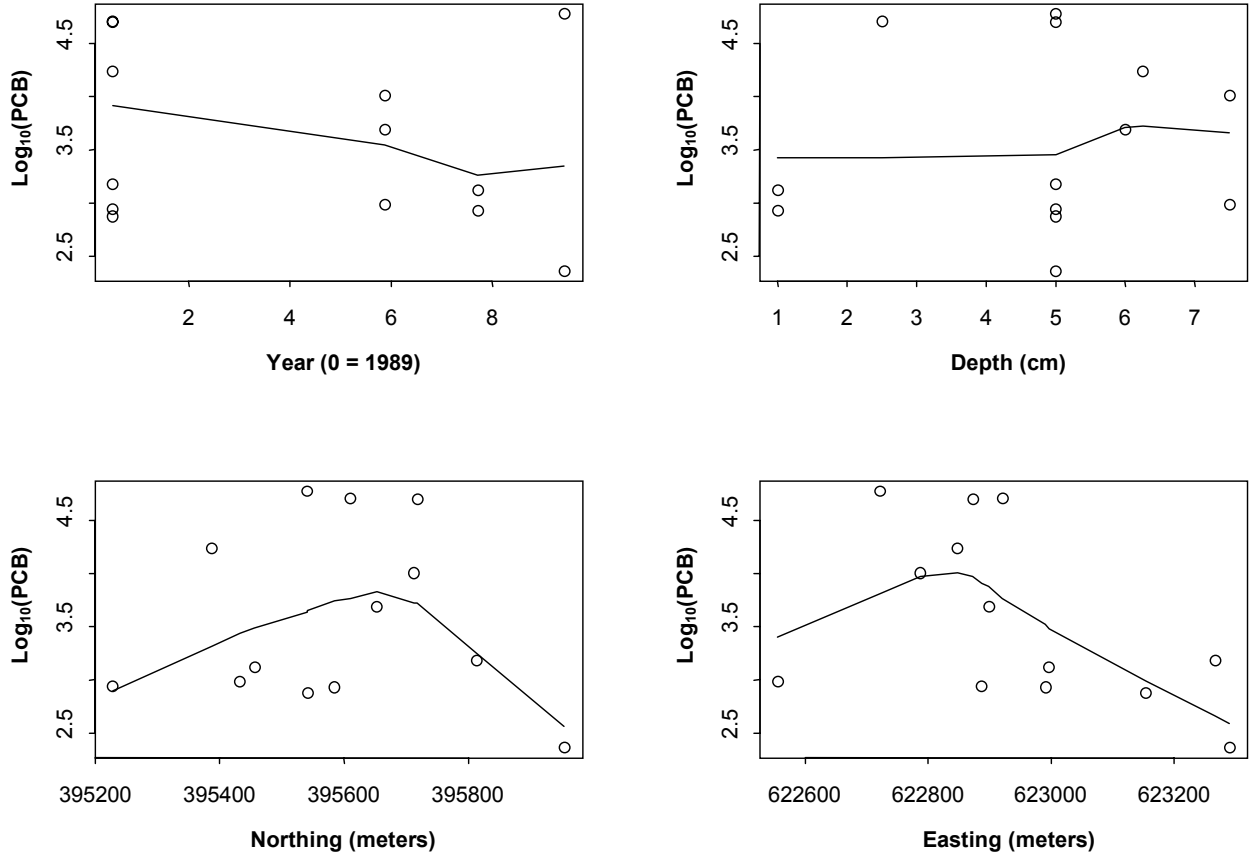


Figure A-51 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group POG (0 to 10 cm) Including Fitted Smoothed Line

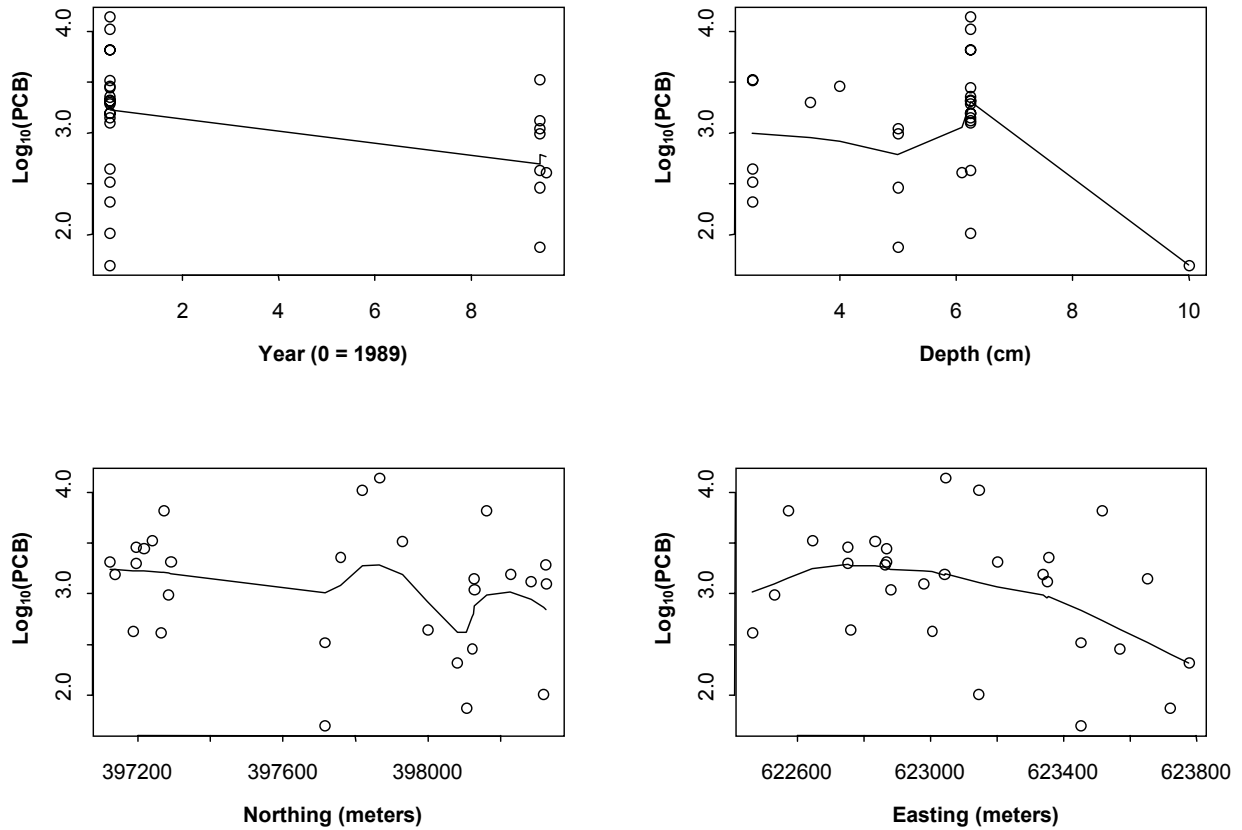


Figure A-52 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group F (0 to 10 cm) Including Fitted Smoothed Line

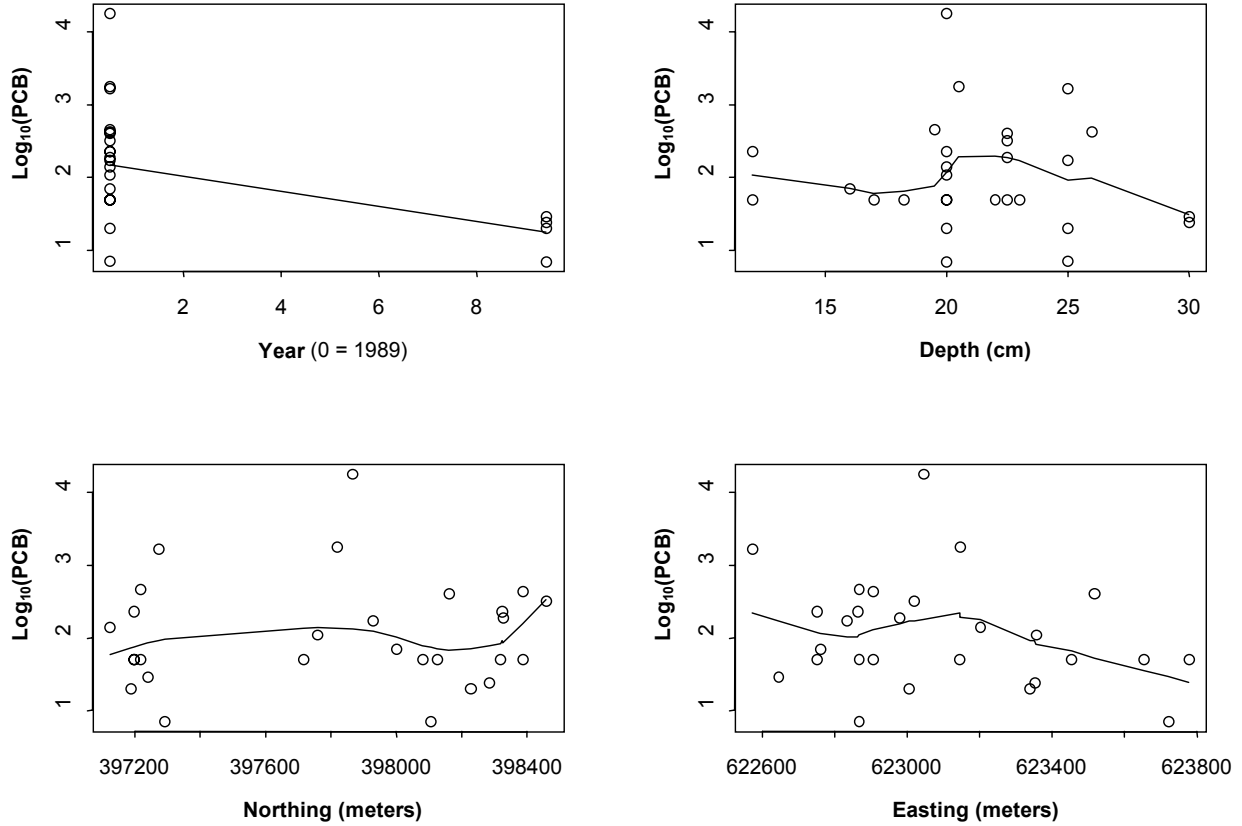


Figure A-53 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group F (10 to 30 cm) Including Fitted Smoothed Line

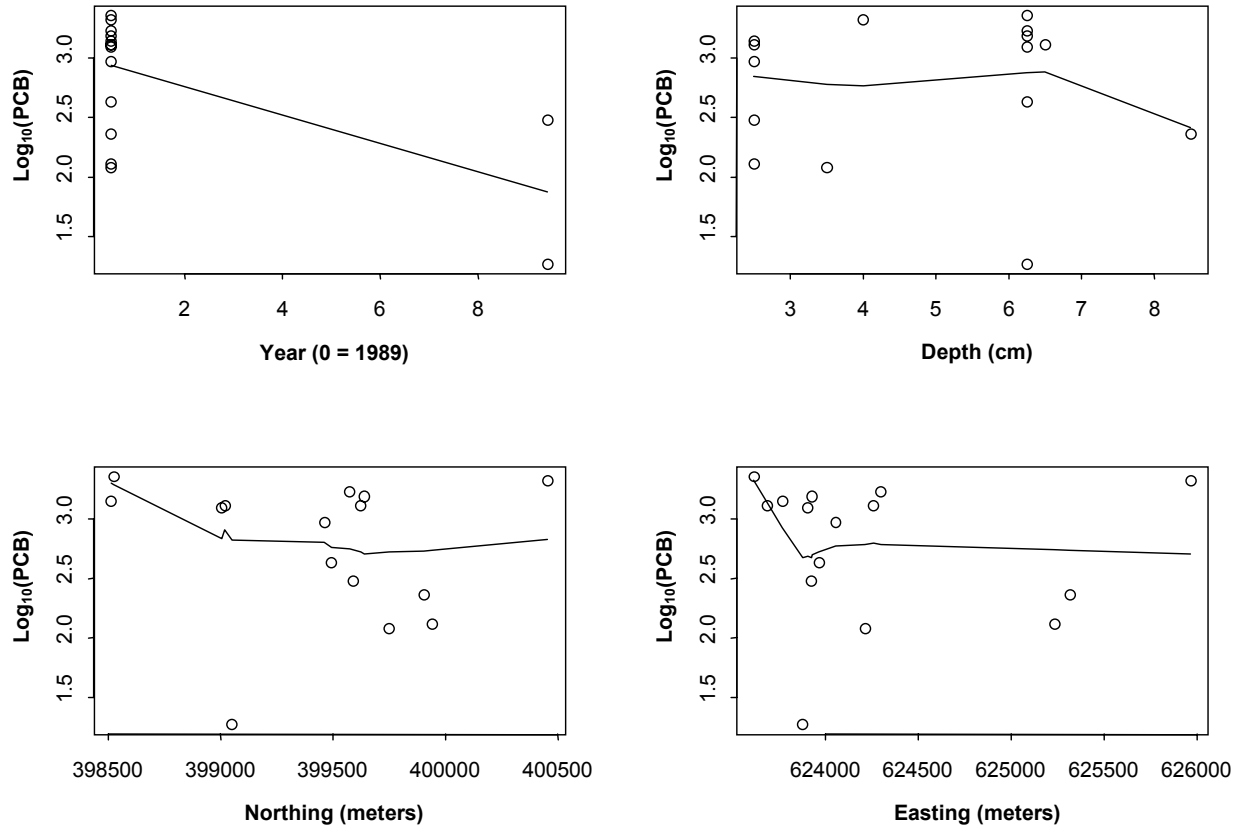


Figure A-54 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Lake Butte des Morts Deposit Group GH (0 to 10 cm) Including Fitted Smoothed Line

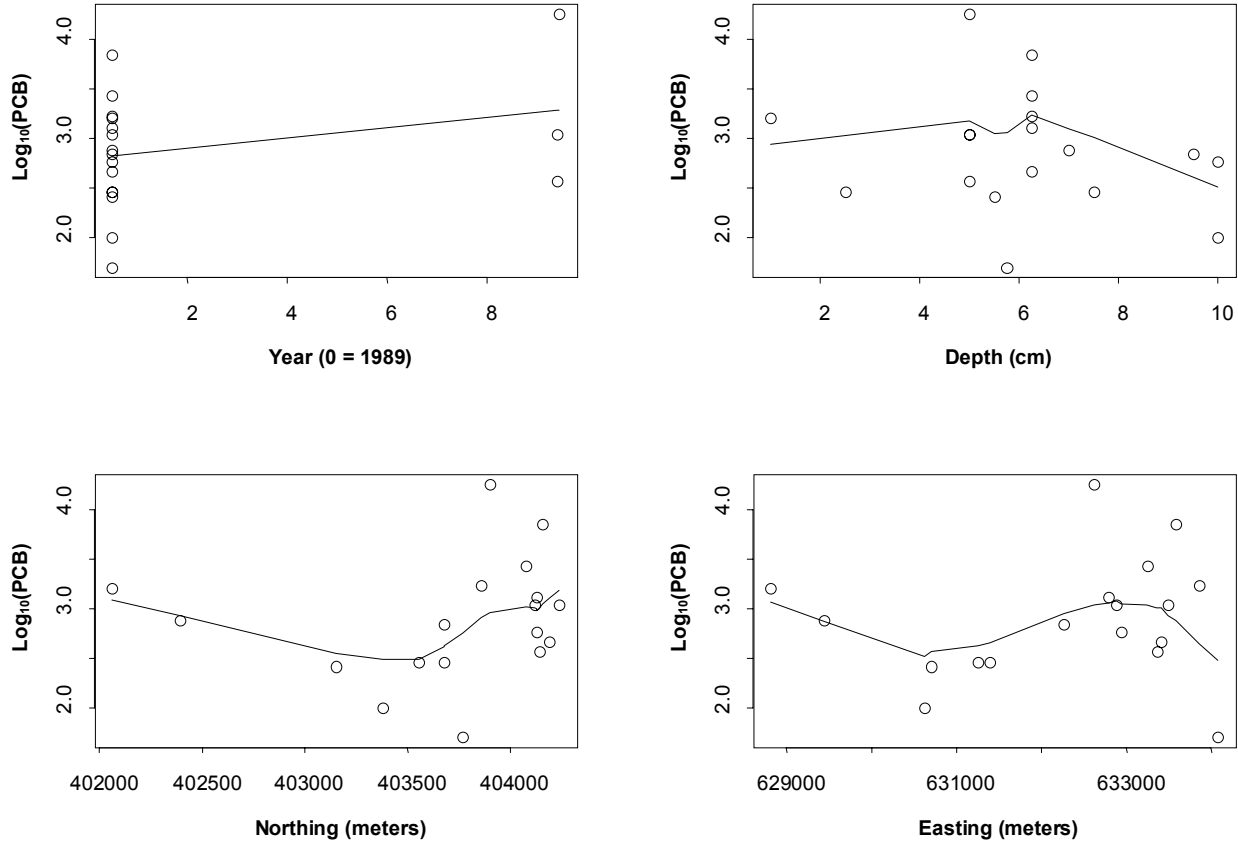


Figure A-55 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Appleton Deposit Group IMOR (0 to 10 cm) Including Fitted Smoothed Line

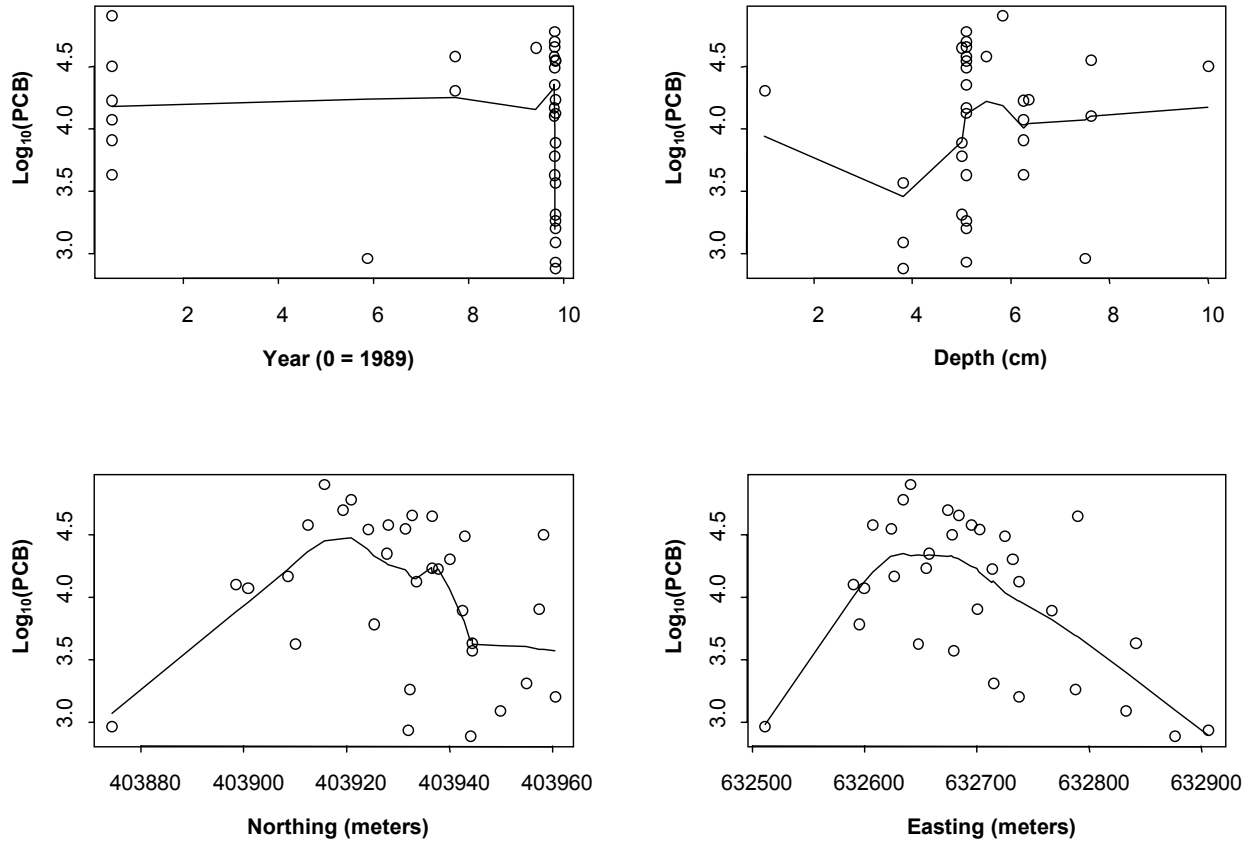


Figure A-56 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Appleton Deposit Group N Before Demonstration Project (0 to 10 cm) Including Fitted Smoothed Line

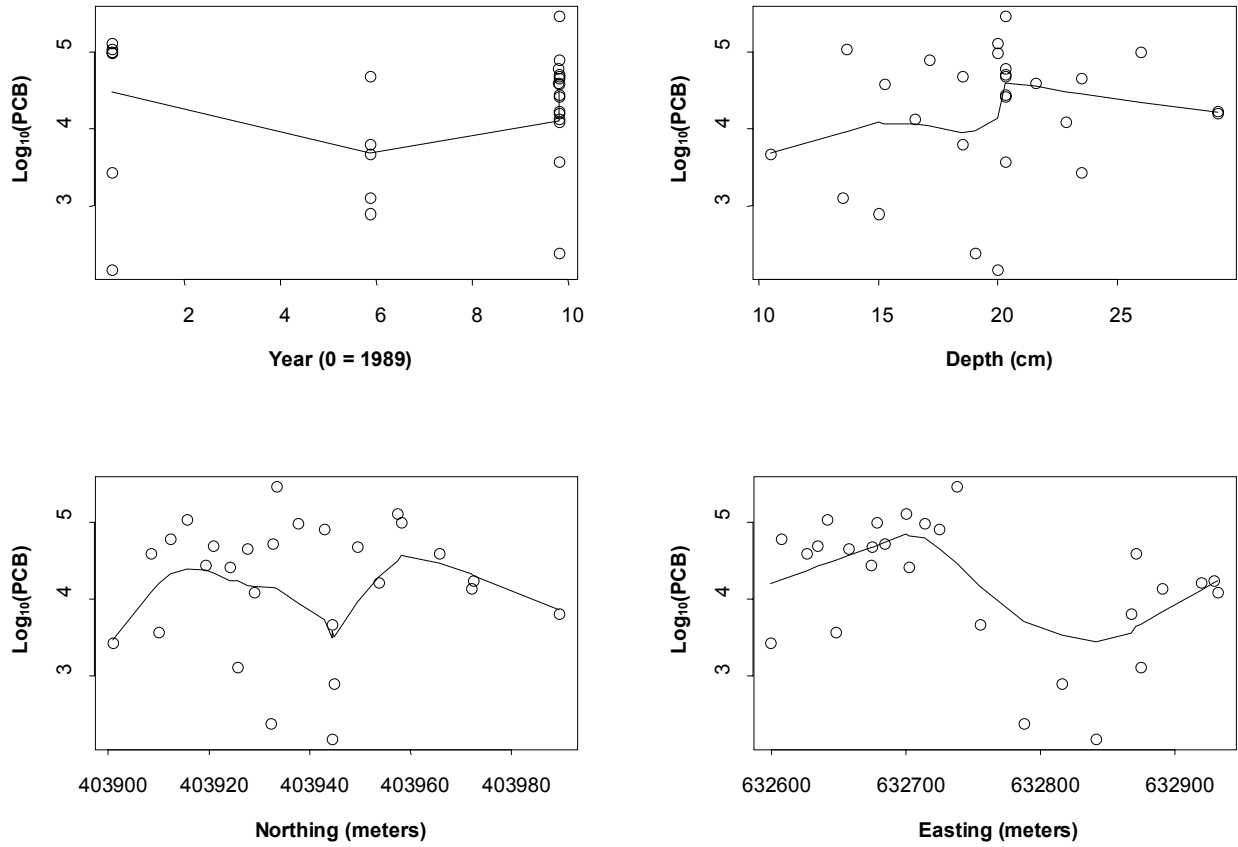


Figure A-57 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Appleton Deposit Group N Before Demonstration Project (10 to 30 cm) Including Fitted Smoothed Line

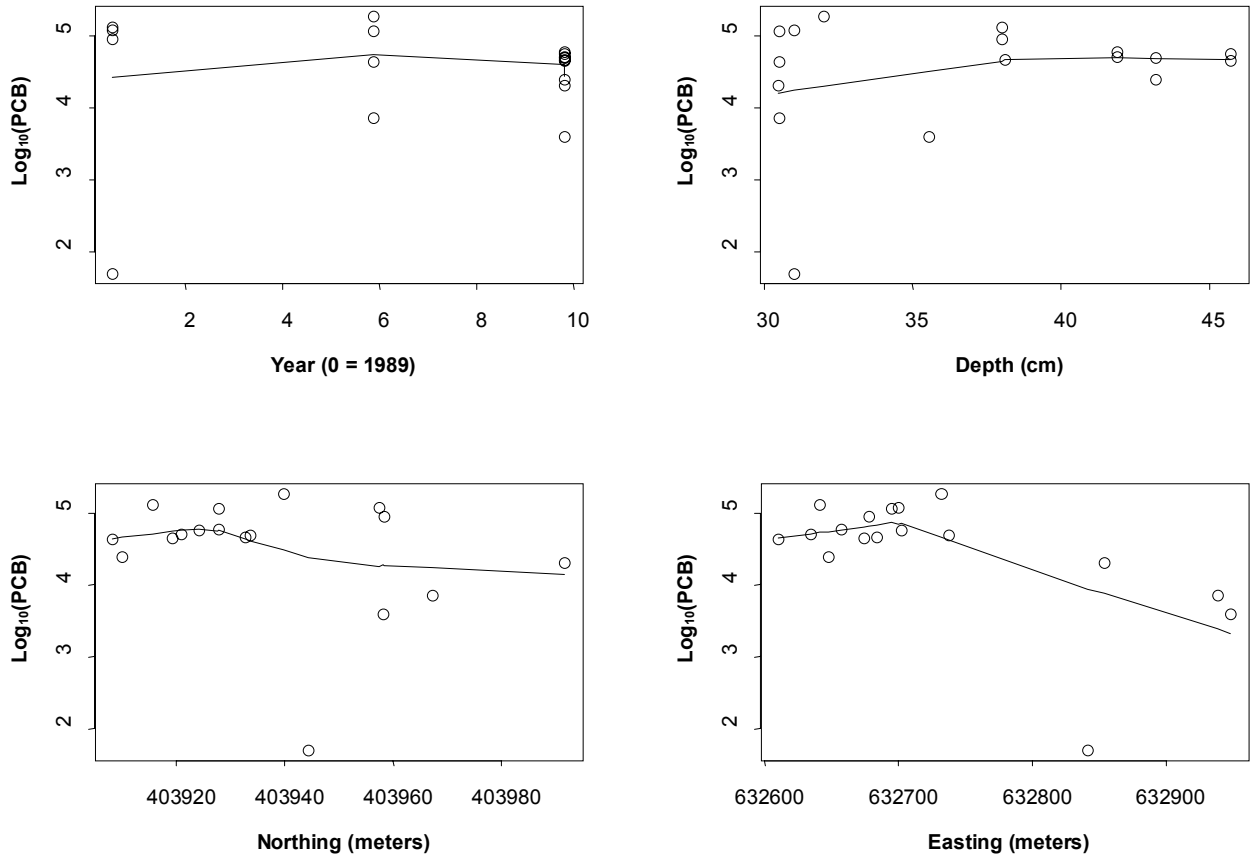


Figure A-58 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Appleton Deposit Group N Before Demonstration Project (30 to 50 cm) Including Fitted Smoothed Line

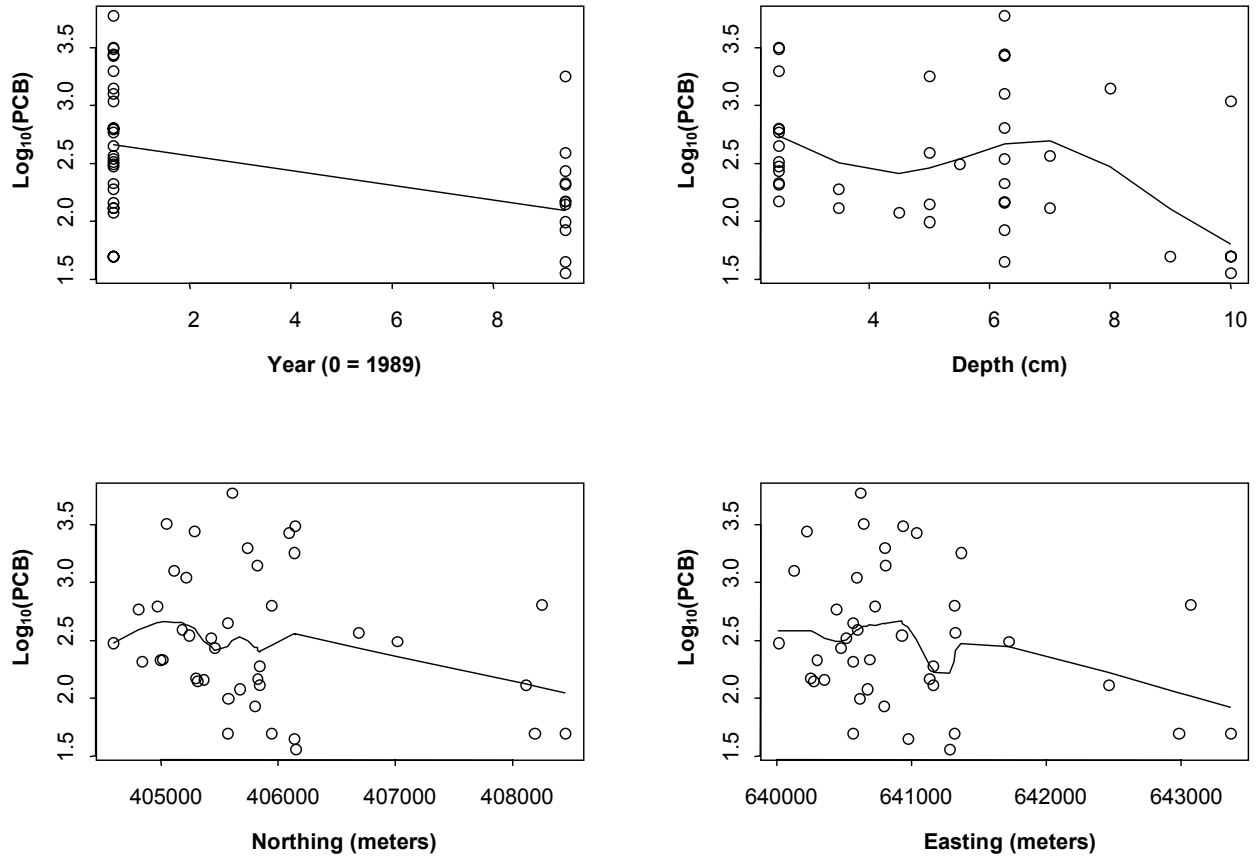


Figure A-59 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Appleton Deposit Group VCC (0 to 10 cm) Including Fitted Smoothed Line

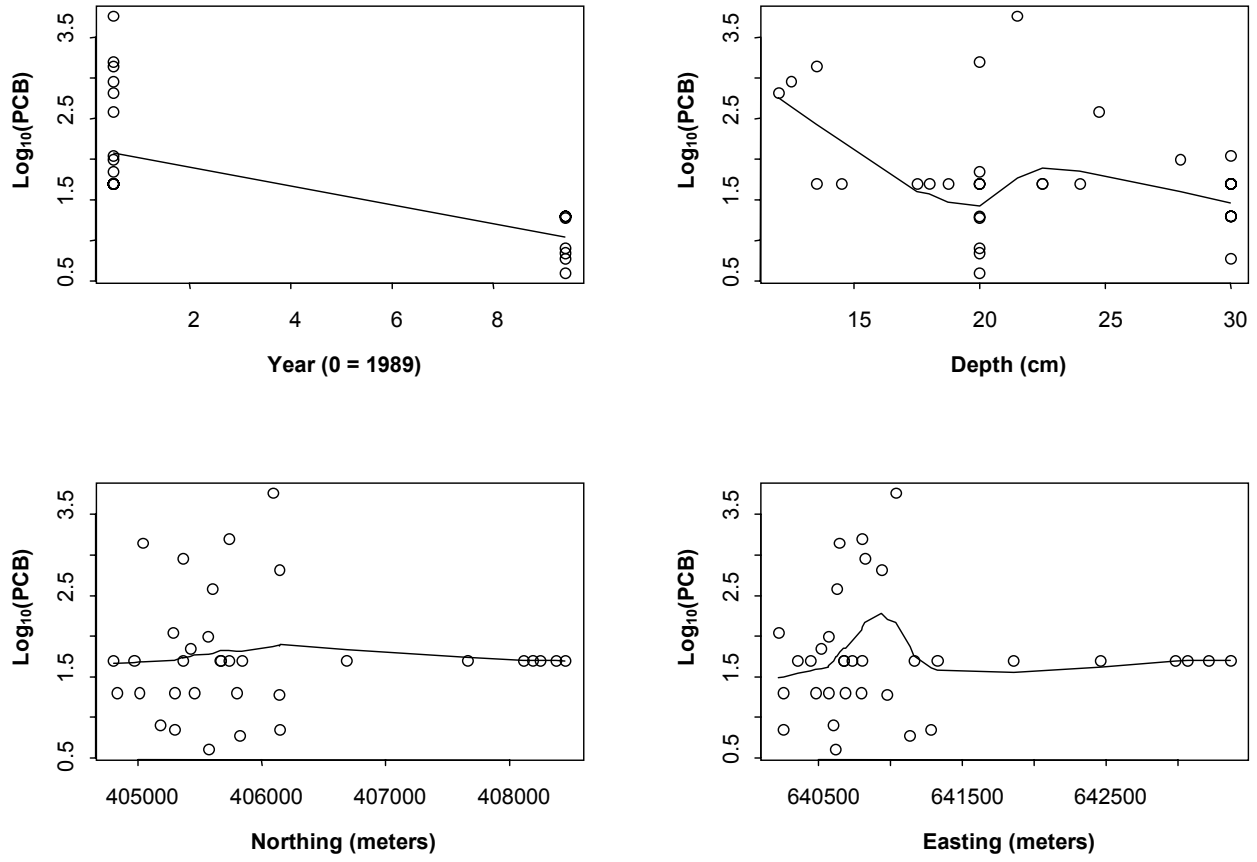


Figure A-60 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Appleton Deposit Group VCC (10 to 30 cm) Including Fitted Smoothed Line

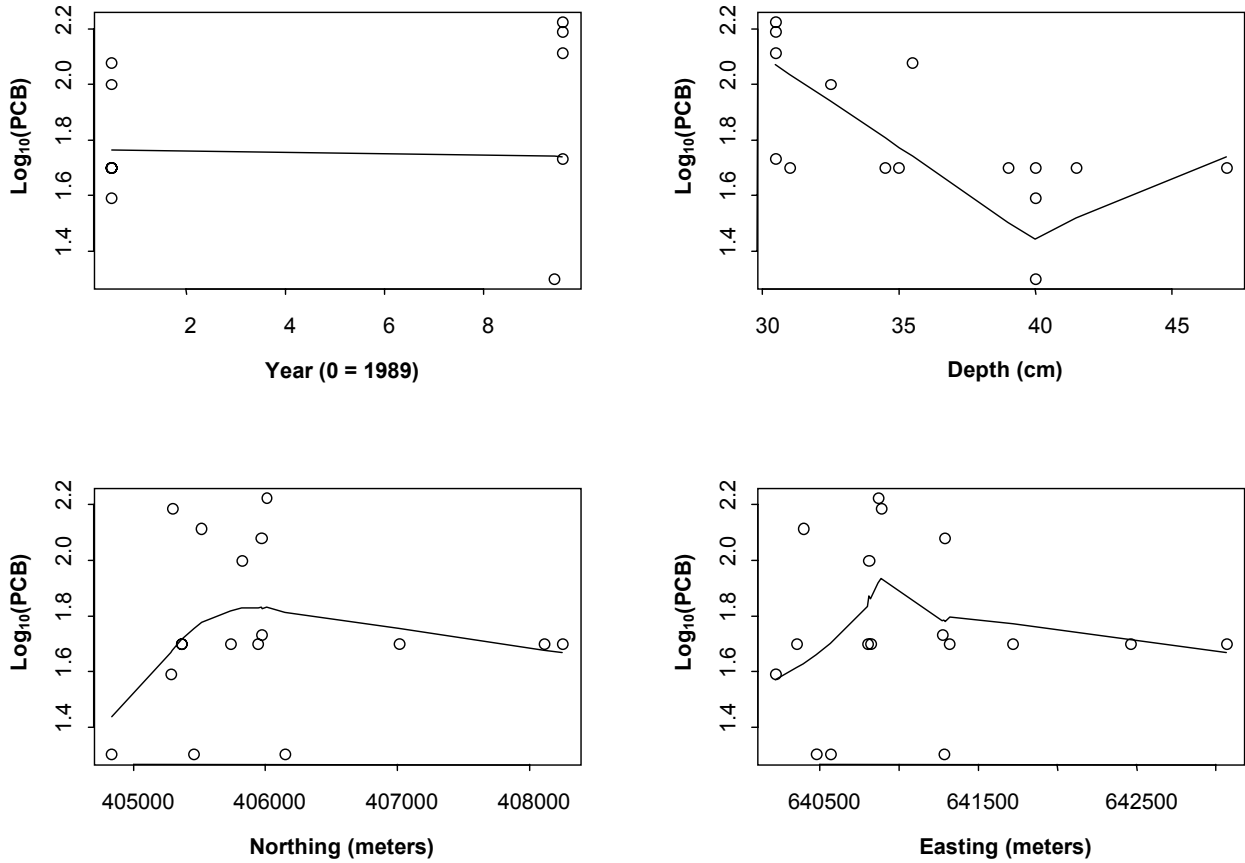


Figure A-61 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Appleton Deposit Group VCC (30 to 50 cm) Including Fitted Smoothed Line

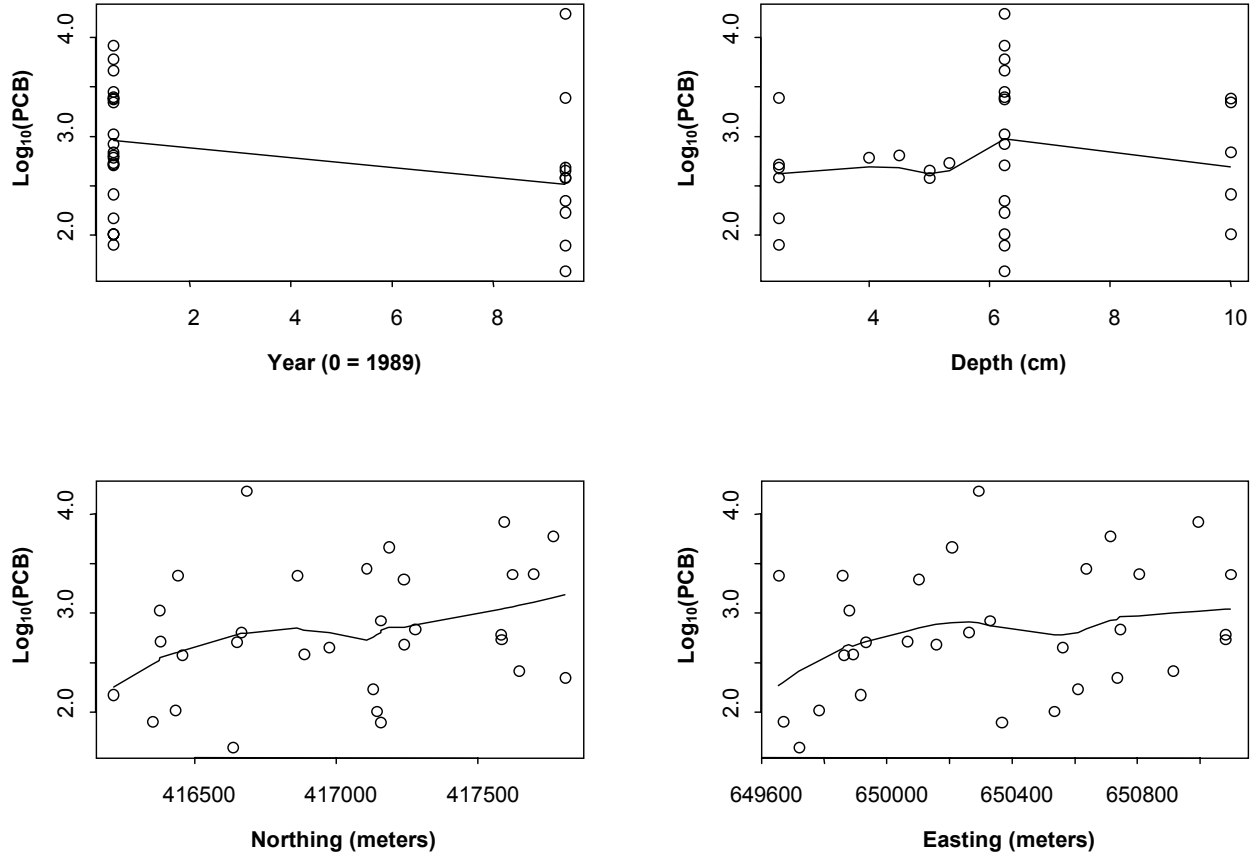


Figure A-62 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group Upper EE (0 to 10 cm) Including Fitted Smoothed Line

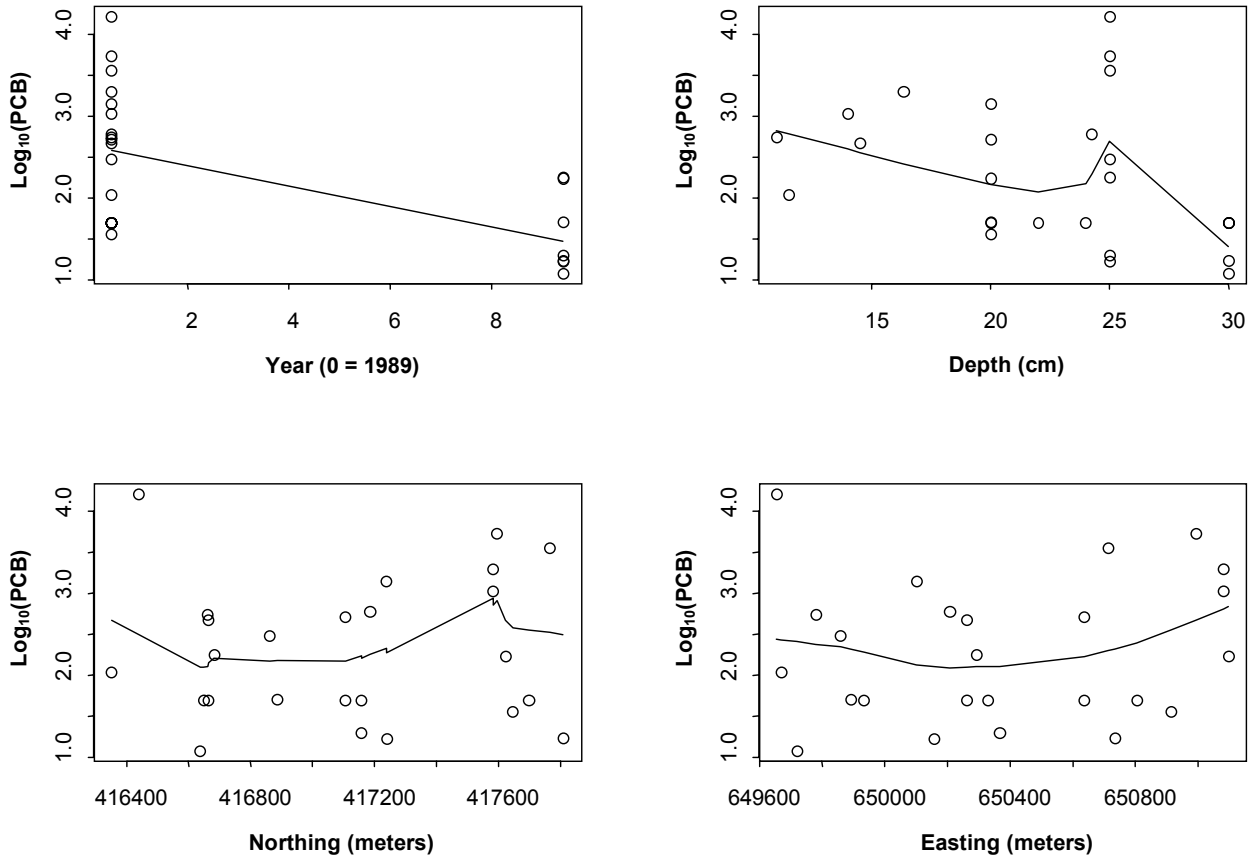


Figure A-63 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group Upper EE (10 to 30 cm) Including Fitted Smoothed Line

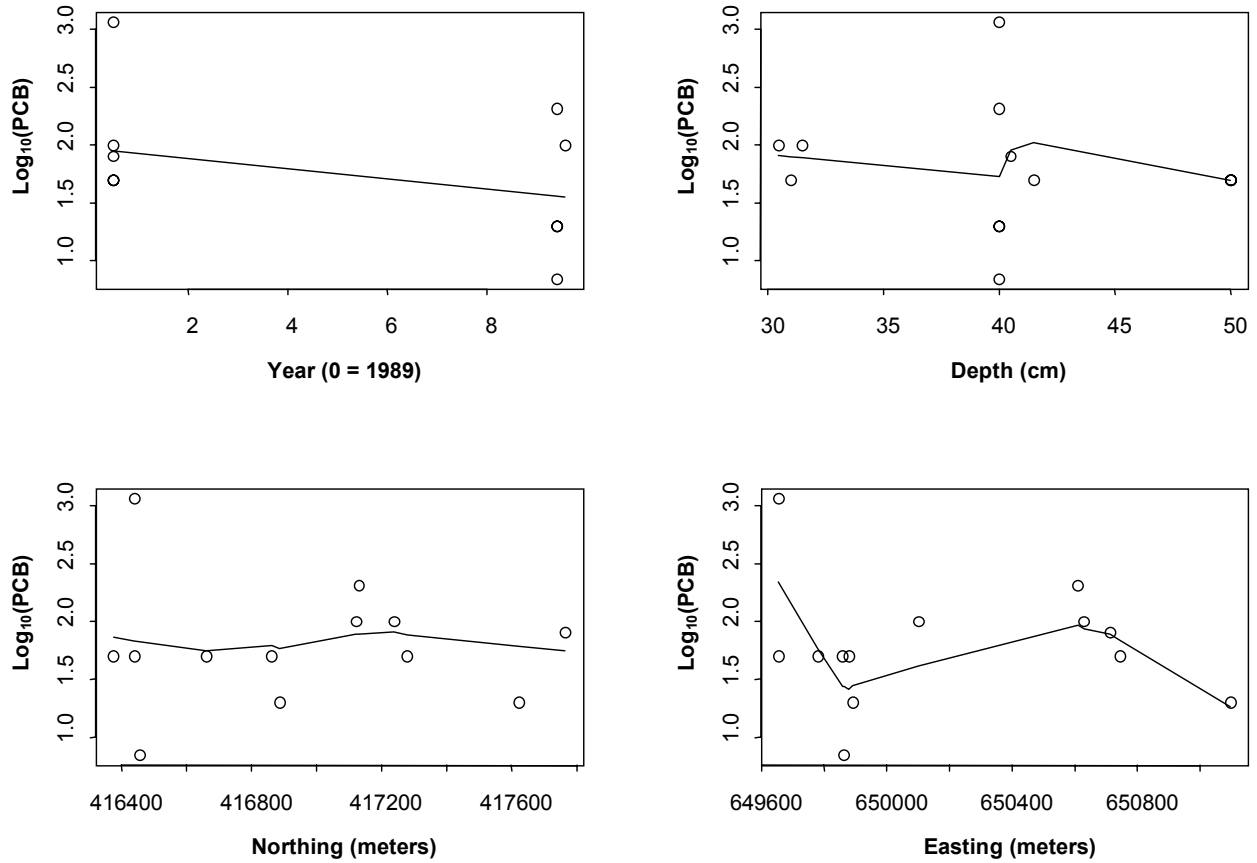


Figure A-64 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group Upper EE (30 to 50 cm) Including Fitted Smoothed Line

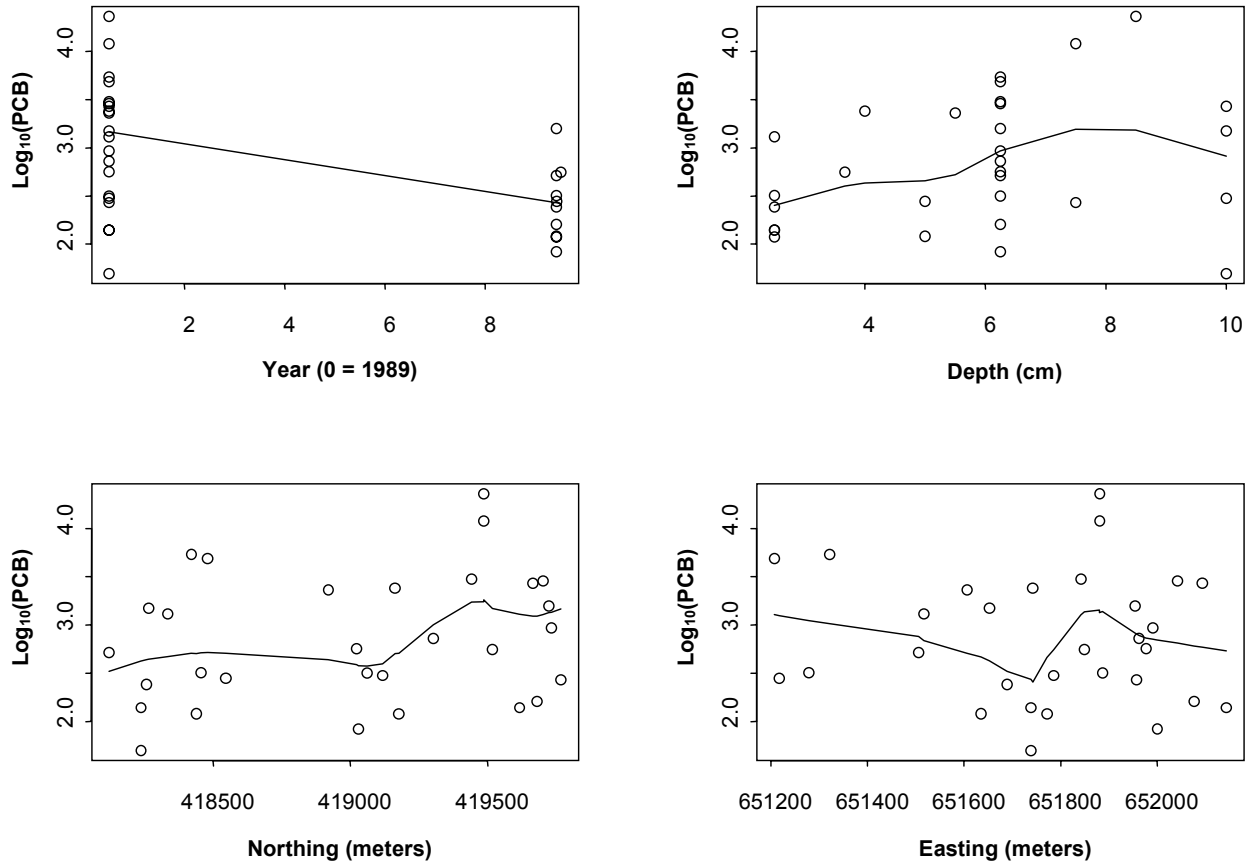


Figure A-65 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group Lower EE (0 to 10 cm) Including Fitted Smoothed Line

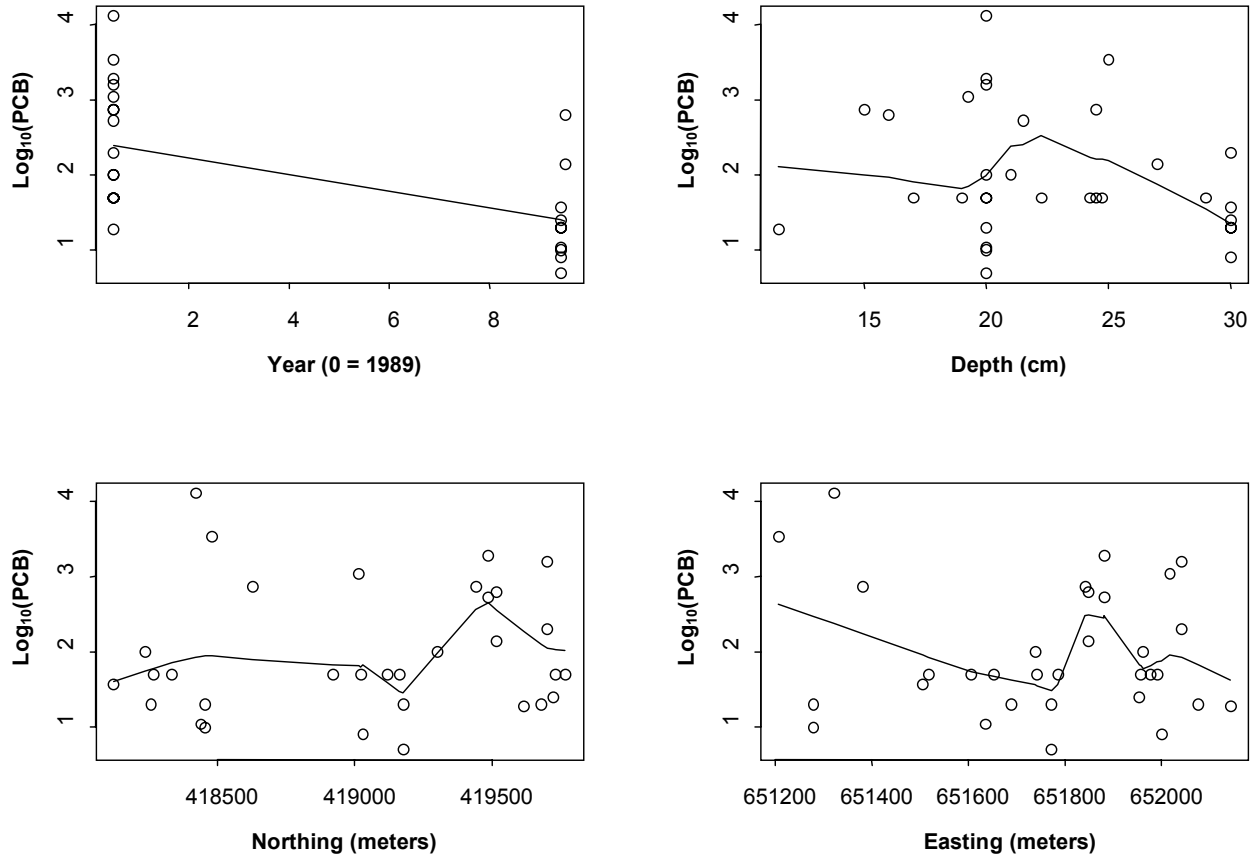


Figure A-66 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group Lower EE (10 to 30 cm) Including Fitted Smoothed Line

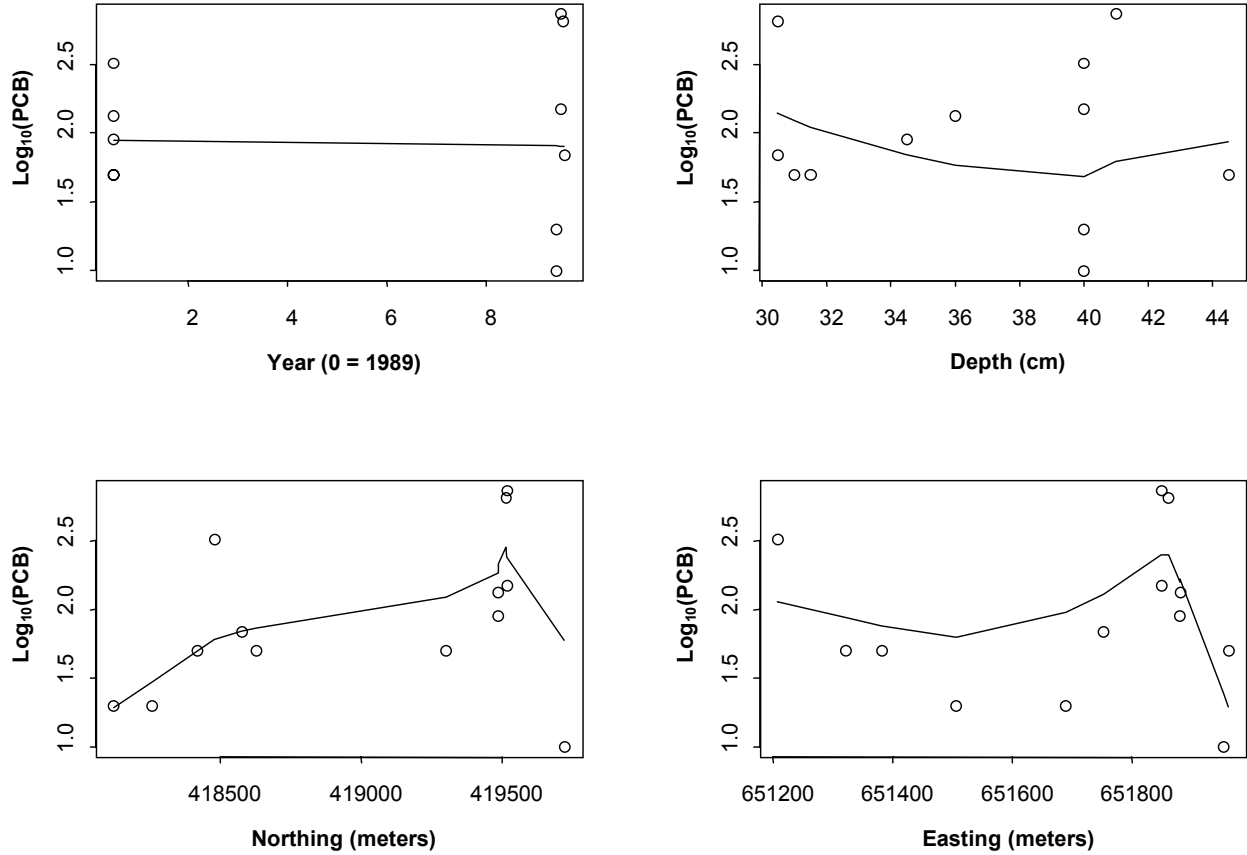


Figure A-67 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group Lower EE (30 to 50 cm) Including Fitted Smoothed Line

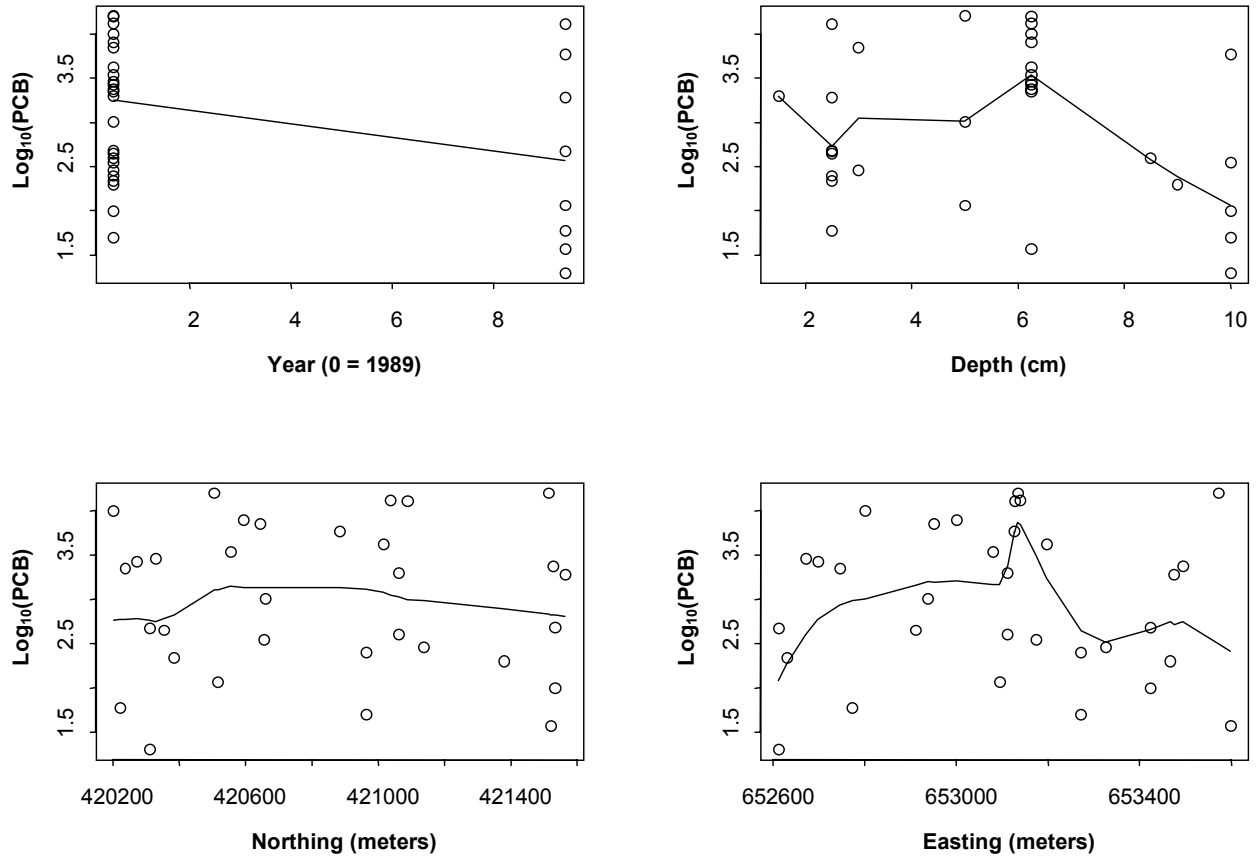


Figure A-68 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group FF (0 to 10 cm) Including Fitted Smoothed Line

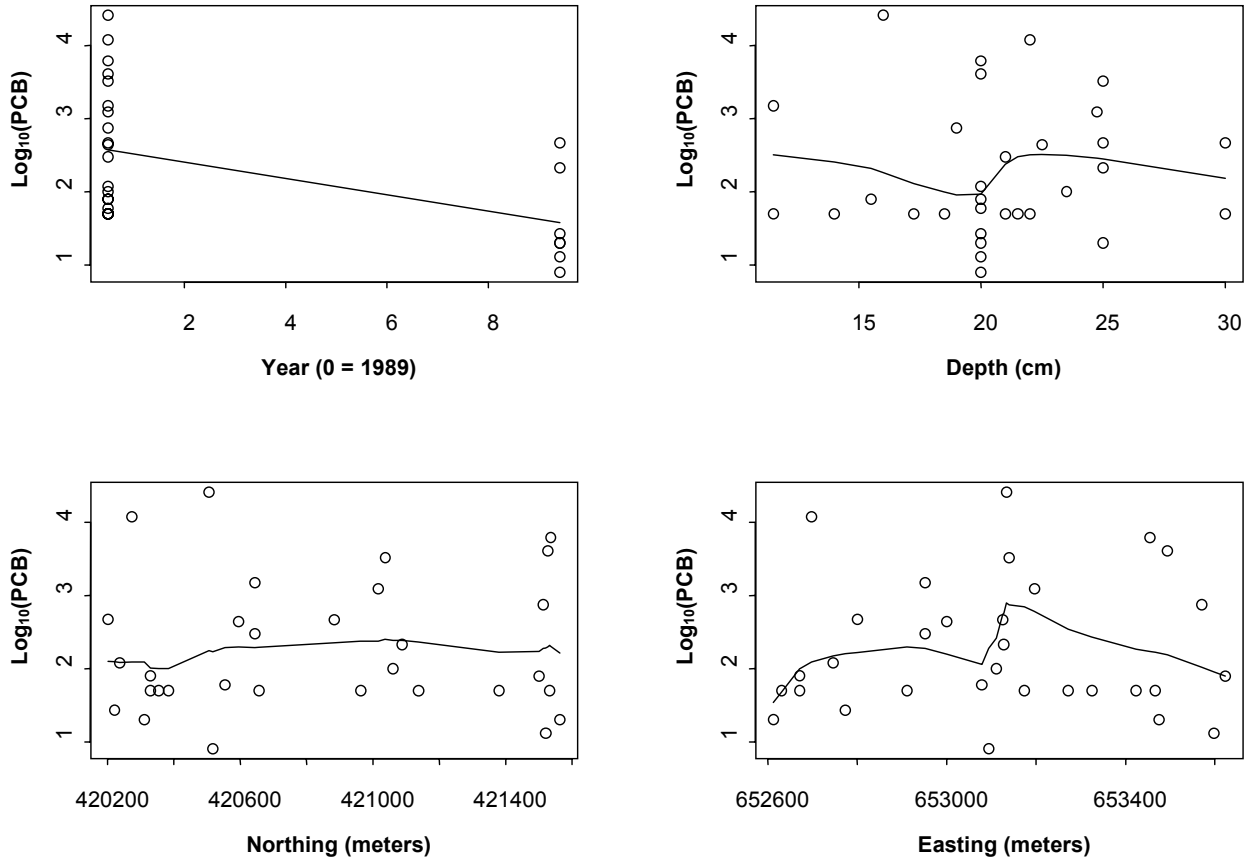


Figure A-69 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group FF (10 to 30 cm) Including Fitted Smoothed Line

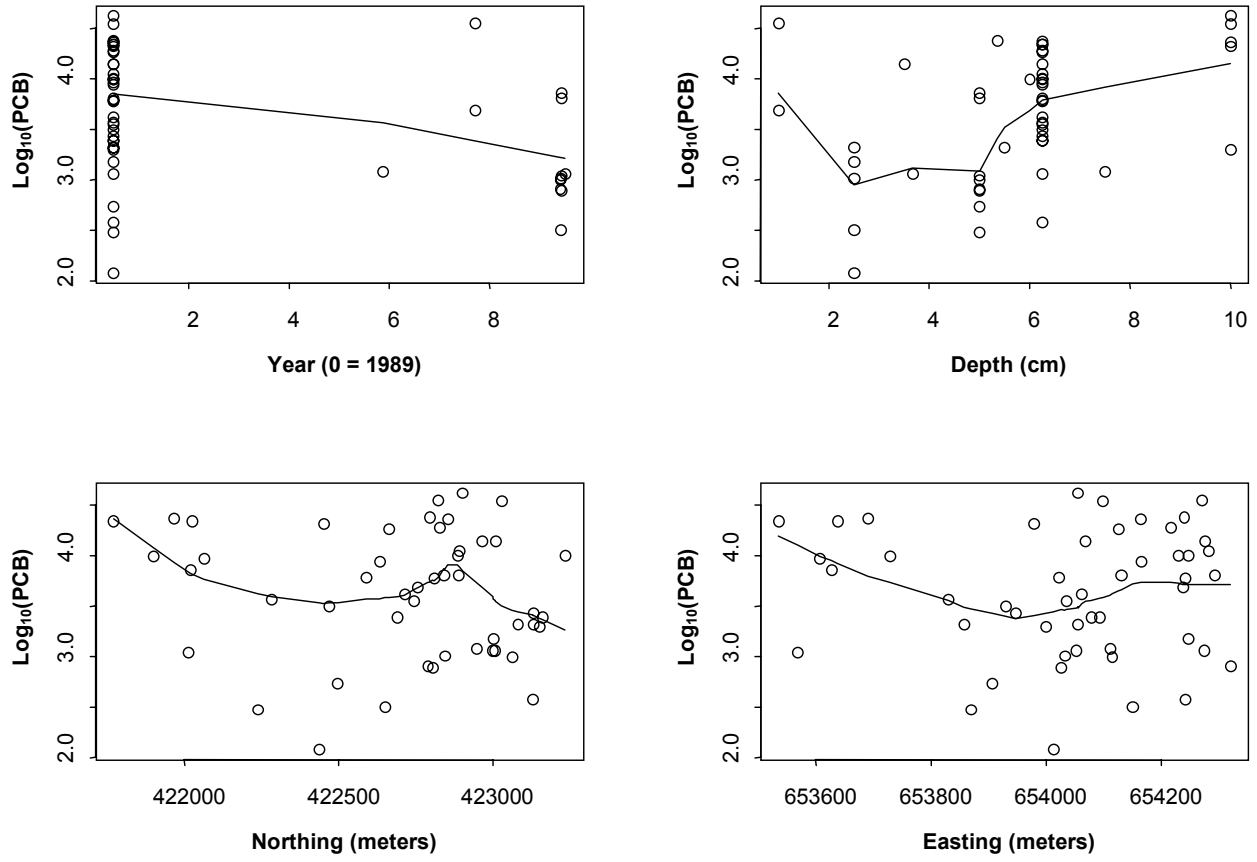


Figure A-70 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group GGHH (0 to 10 cm) Including Fitted Smoothed Line

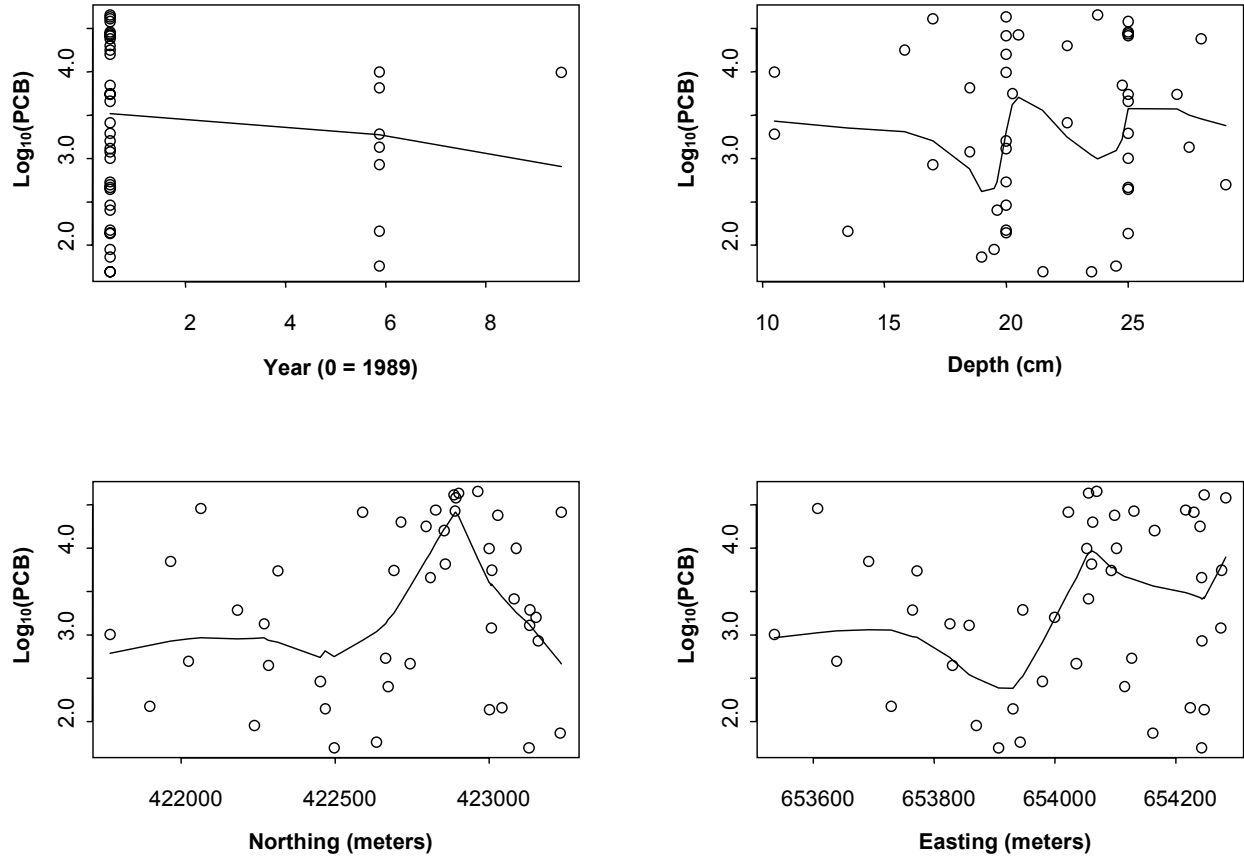


Figure A-71 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group GGHH (10 to 30 cm) Including Fitted Smoothed Line

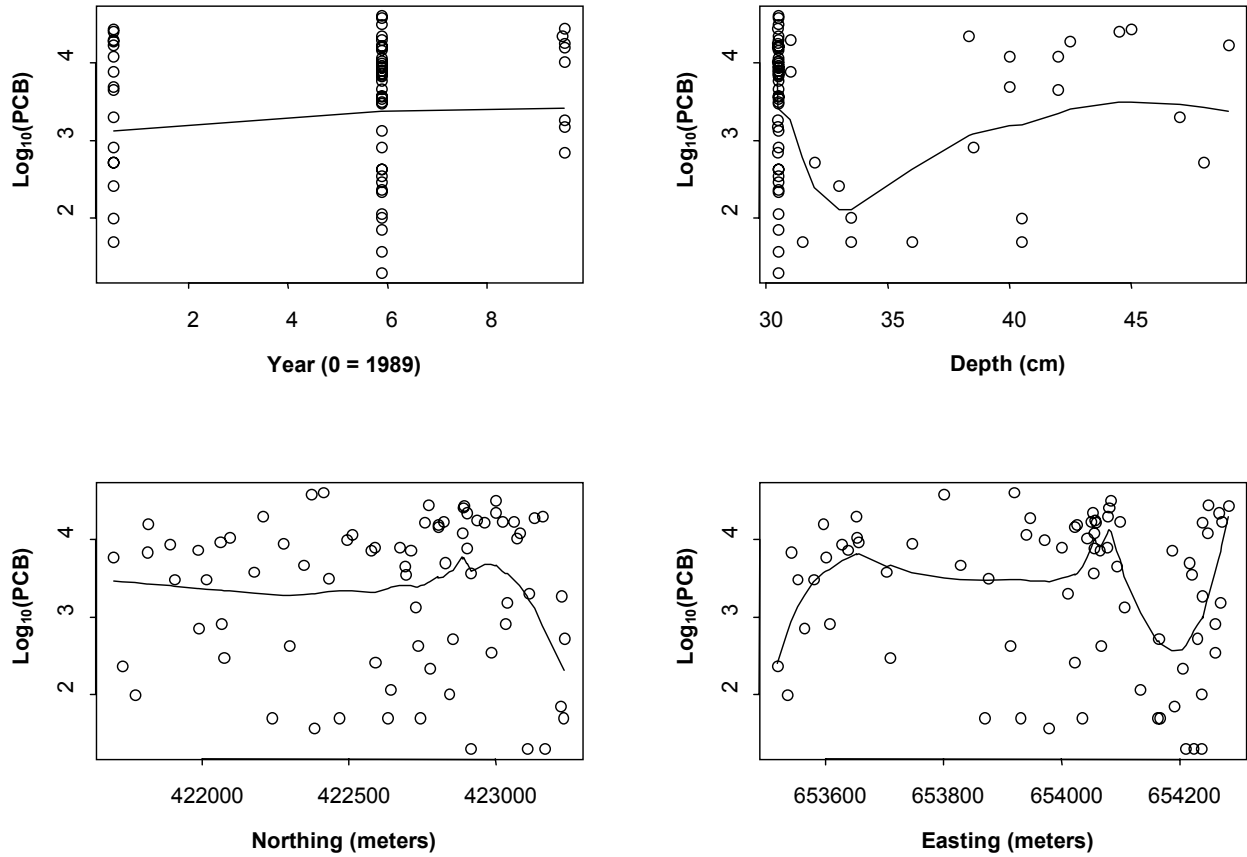


Figure A-72 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group GGHH (30 to 50 cm) Including Fitted Smoothed Line

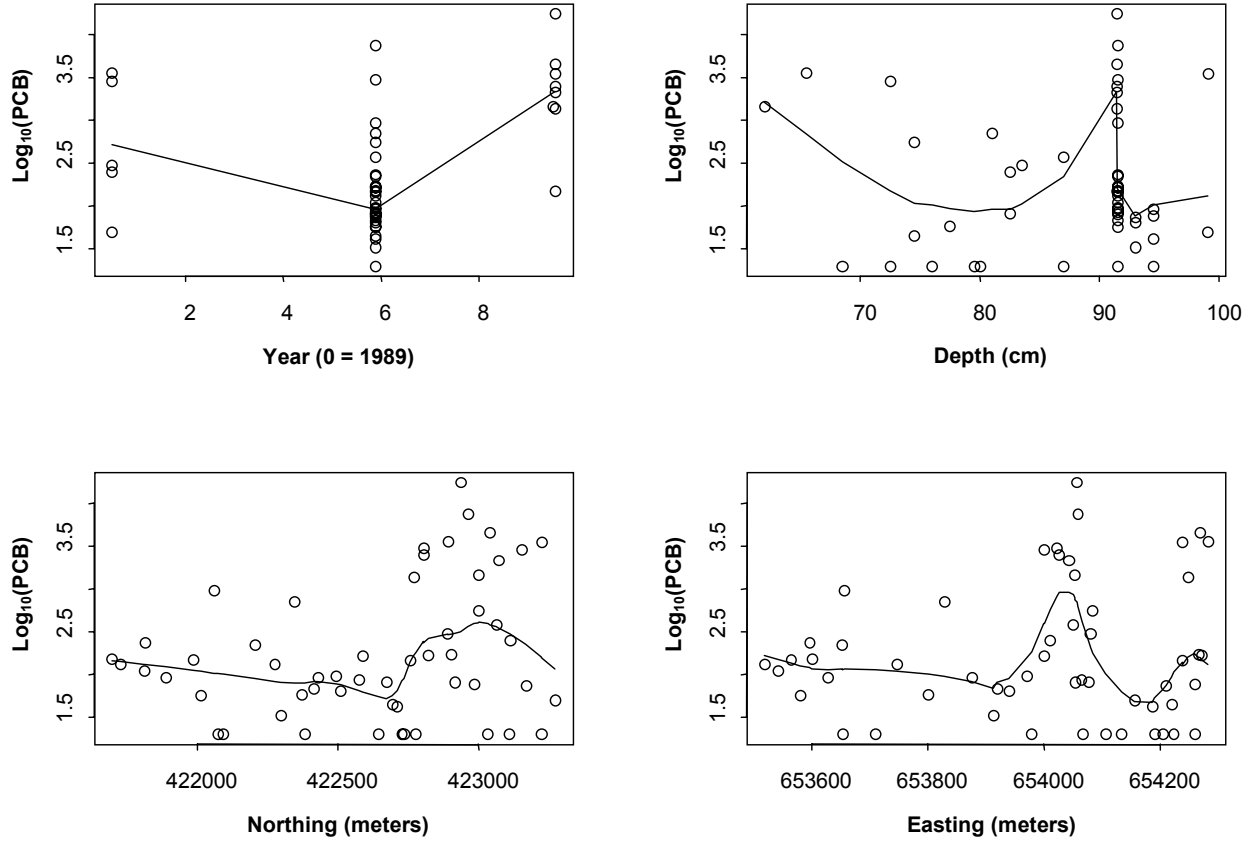


Figure A-73 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group GGHH (50 to 100 cm) Including Fitted Smoothed Line

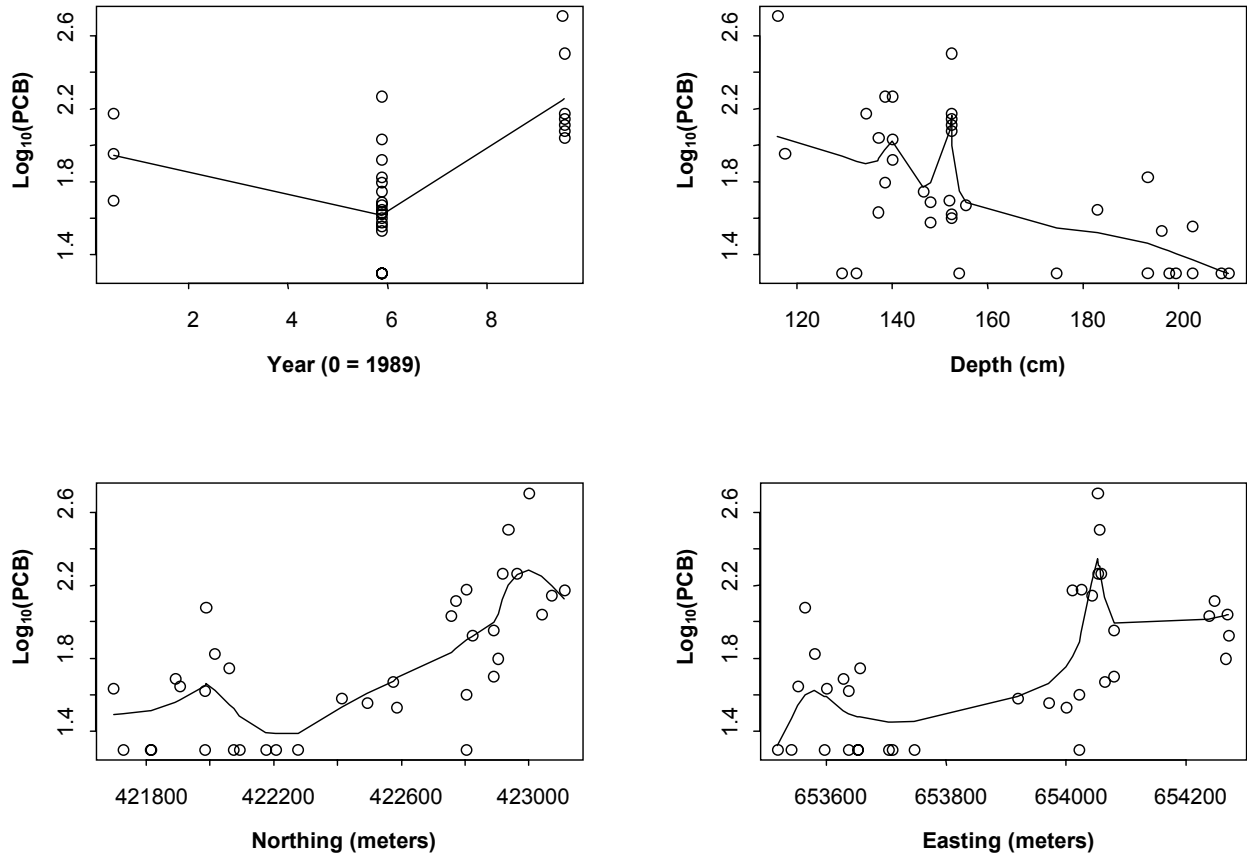


Figure A-74 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for Little Rapids Deposit Group GGHH (100+ cm) Including Fitted Smoothed Line

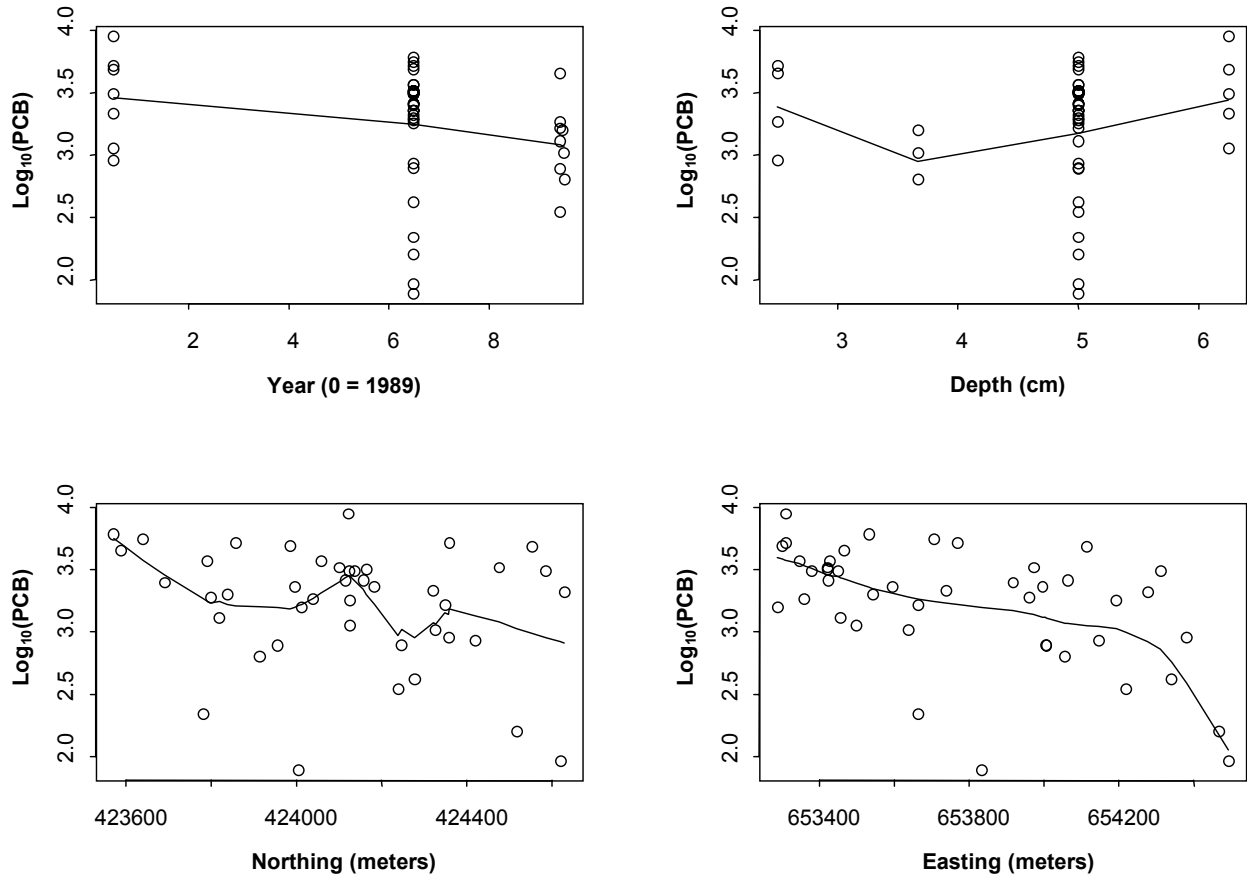


Figure A-75 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 2025 (0 to 10 cm) Including Fitted Smoothed Line

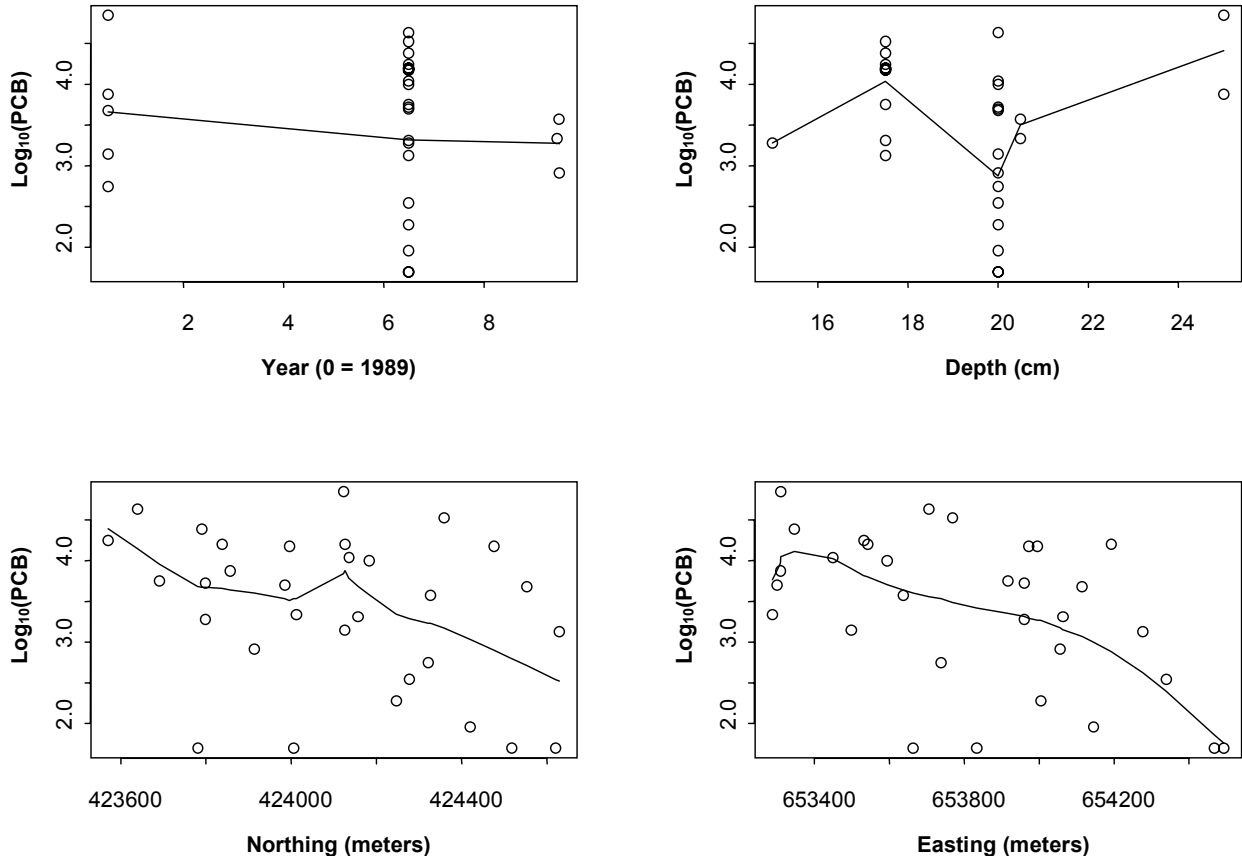


Figure A-76 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 2025 (10 to 30 cm) Including Fitted Smoothed Line

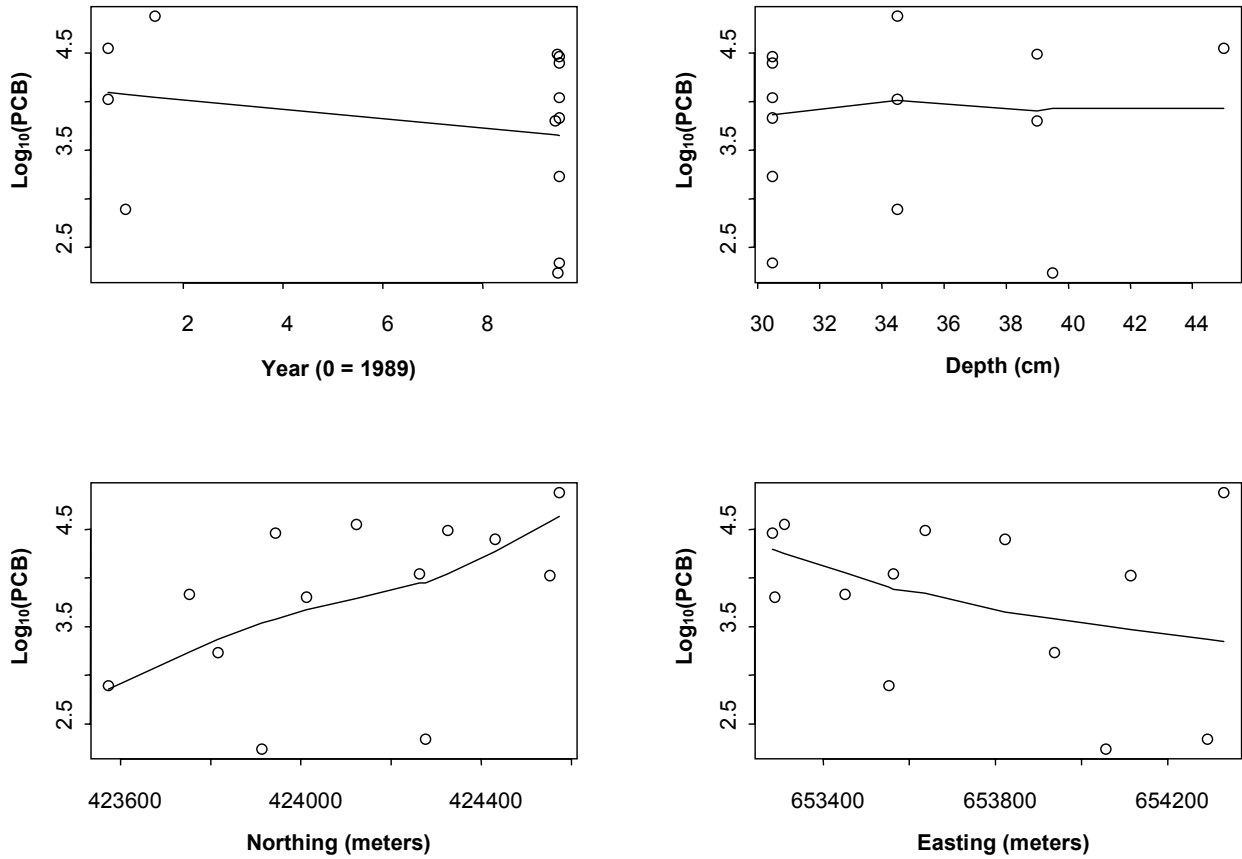


Figure A-77 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 2025 (30 to 50 cm) Including Fitted Smoothed Line

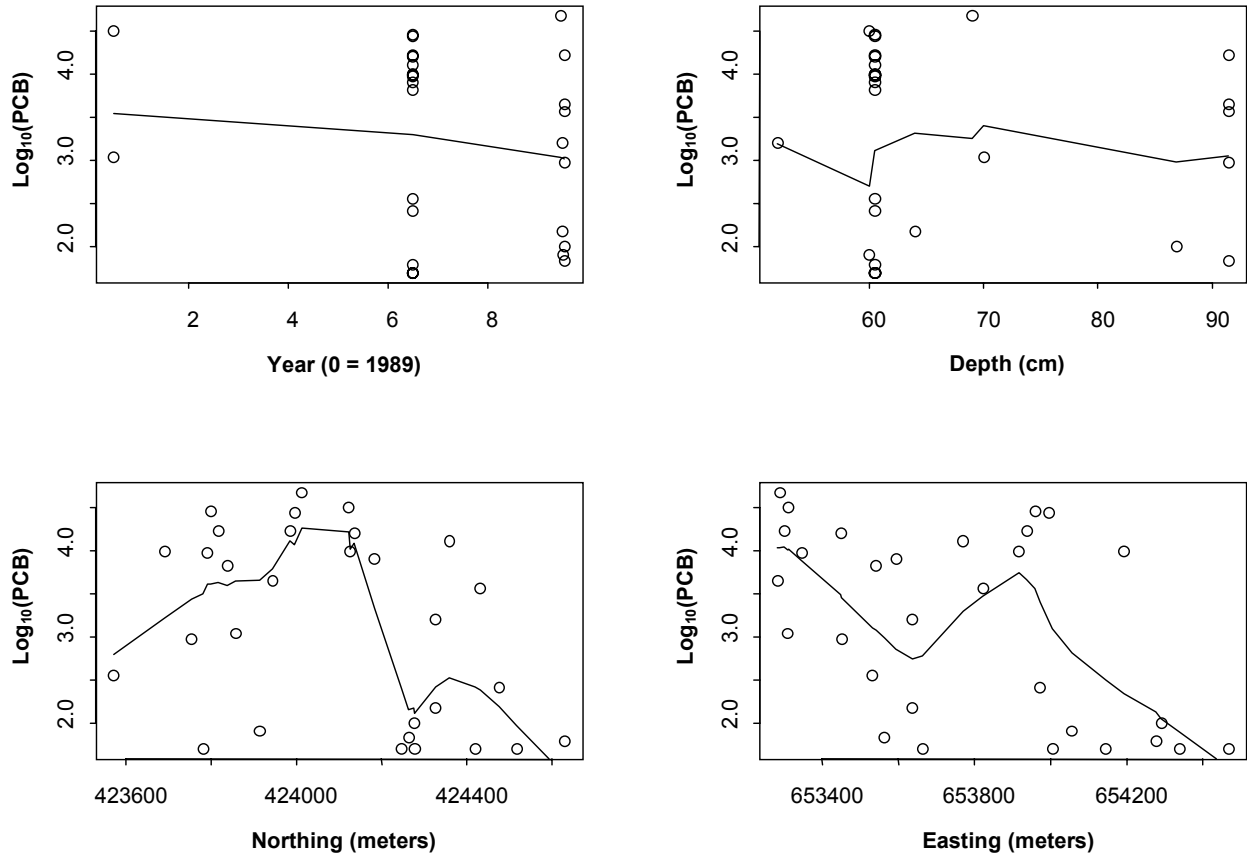


Figure A-78 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 2025 (50 to 100 cm) Including Fitted Smoothed Line

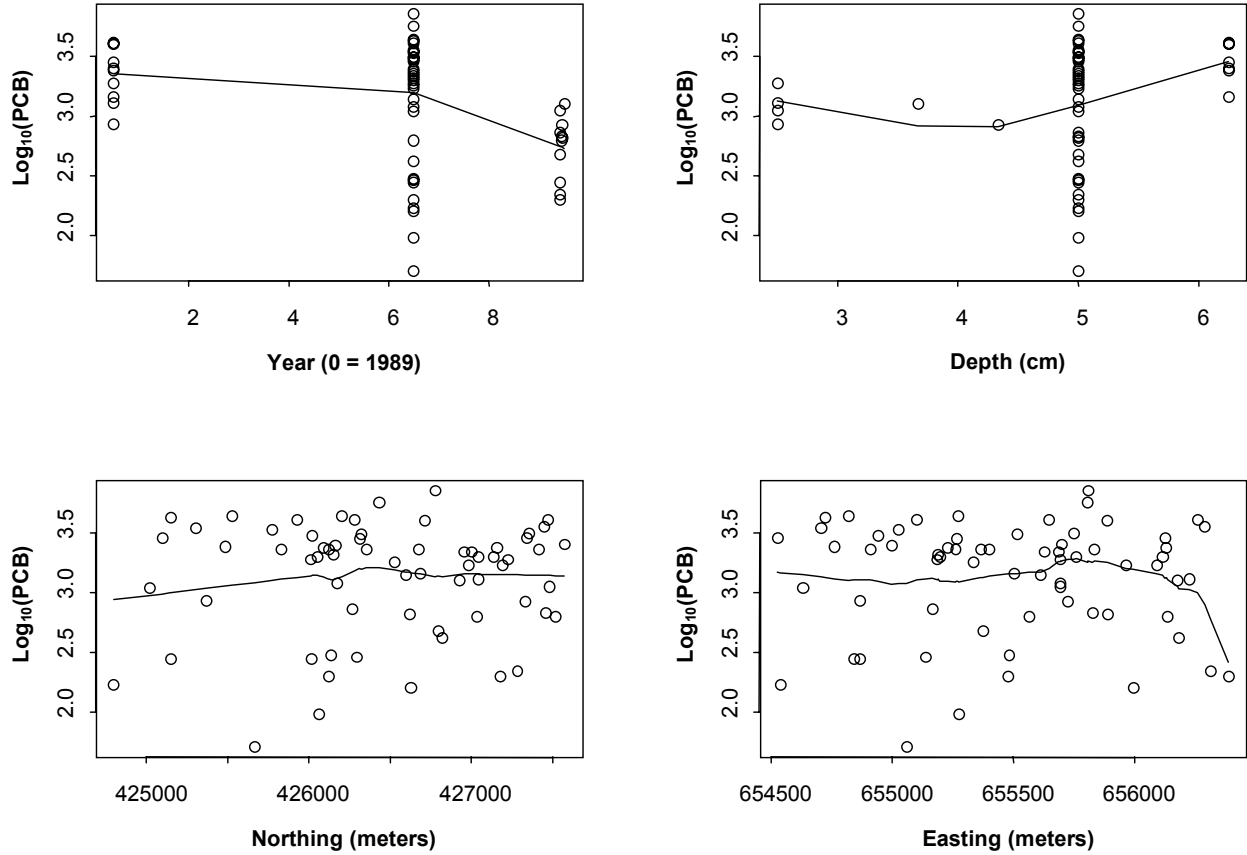


Figure A-79 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 2649 (0 to 10 cm) Including Fitted Smoothed Line

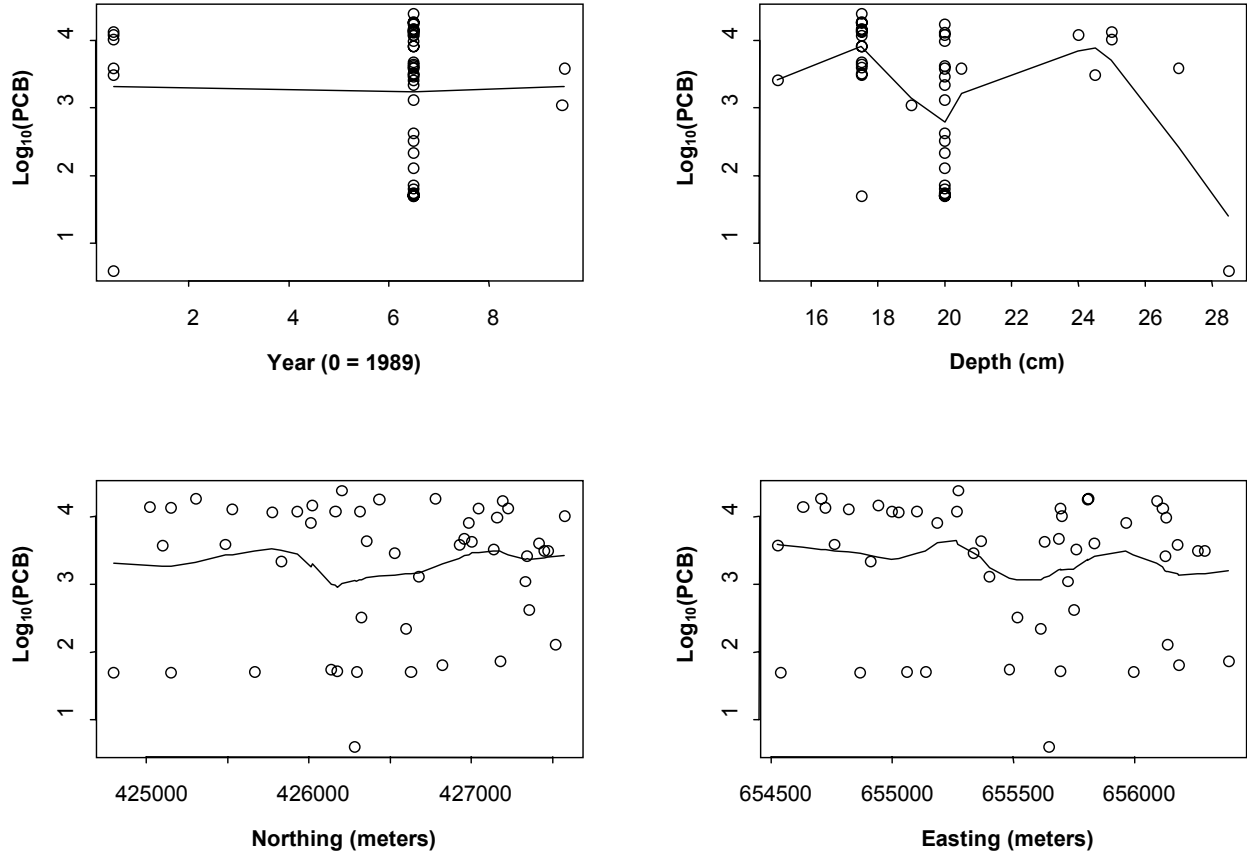


Figure A-80 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 2649 (10 to 30 cm) Including Fitted Smoothed Line

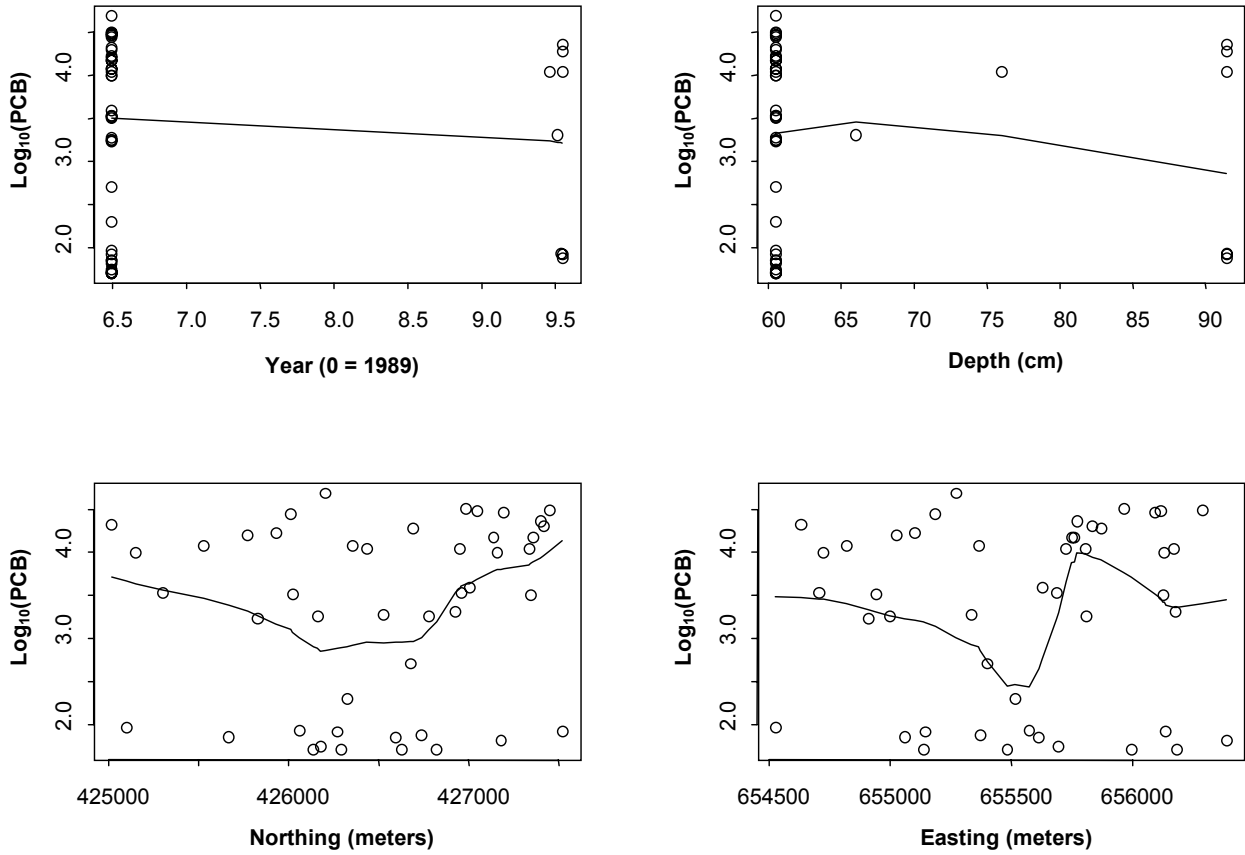


Figure A-81 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 2649 (50 to 100 cm) Including Fitted Smoothed Line

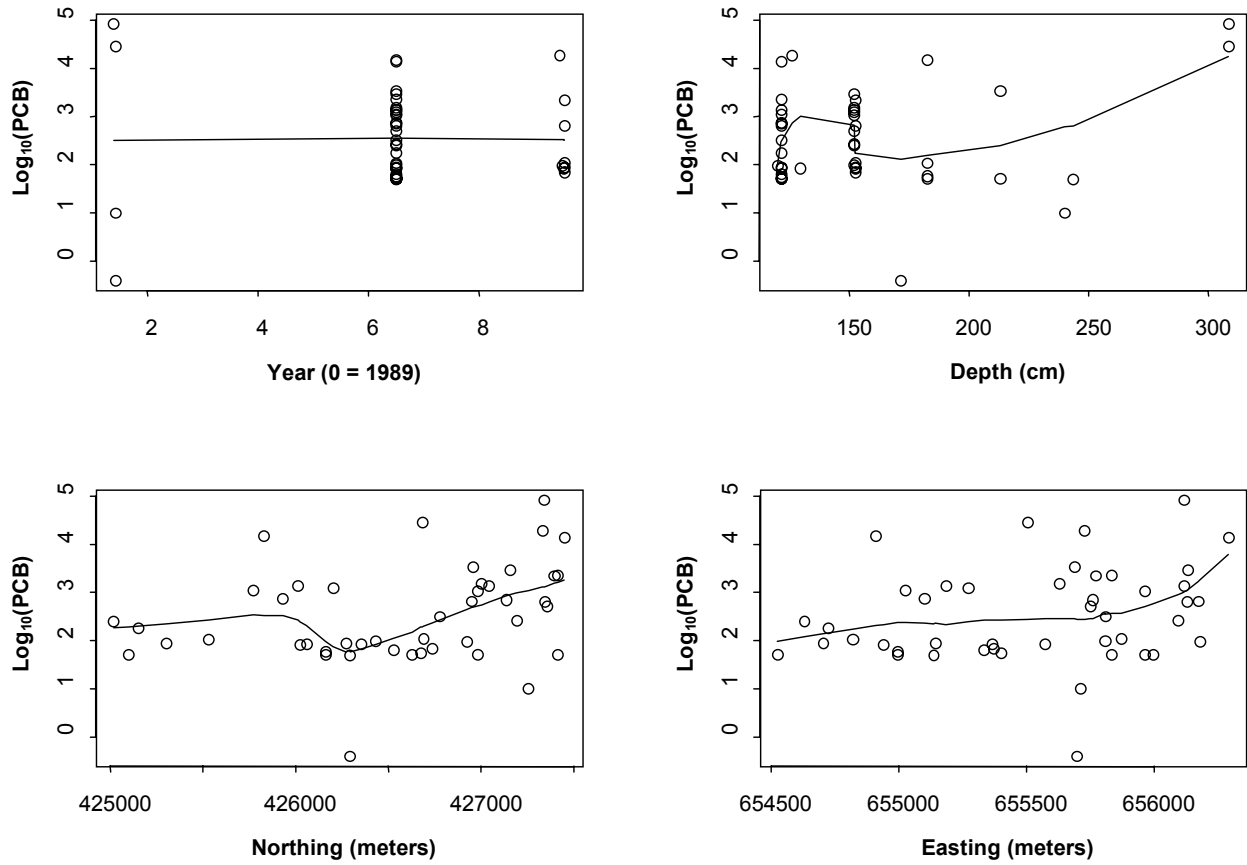


Figure A-82 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 2649 (100+ cm) Including Fitted Smoothed Line

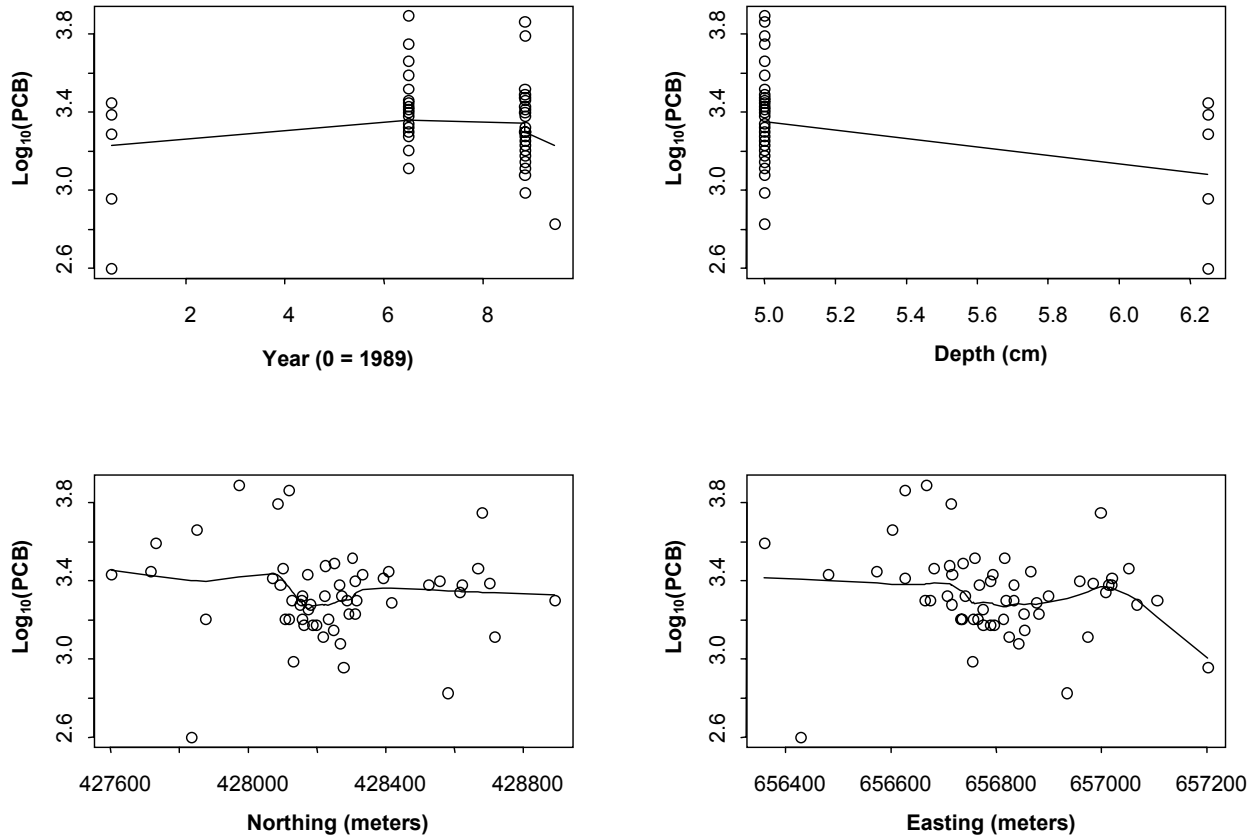


Figure A-83 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 5067 (0 to 10 cm) Including Fitted Smoothed Line

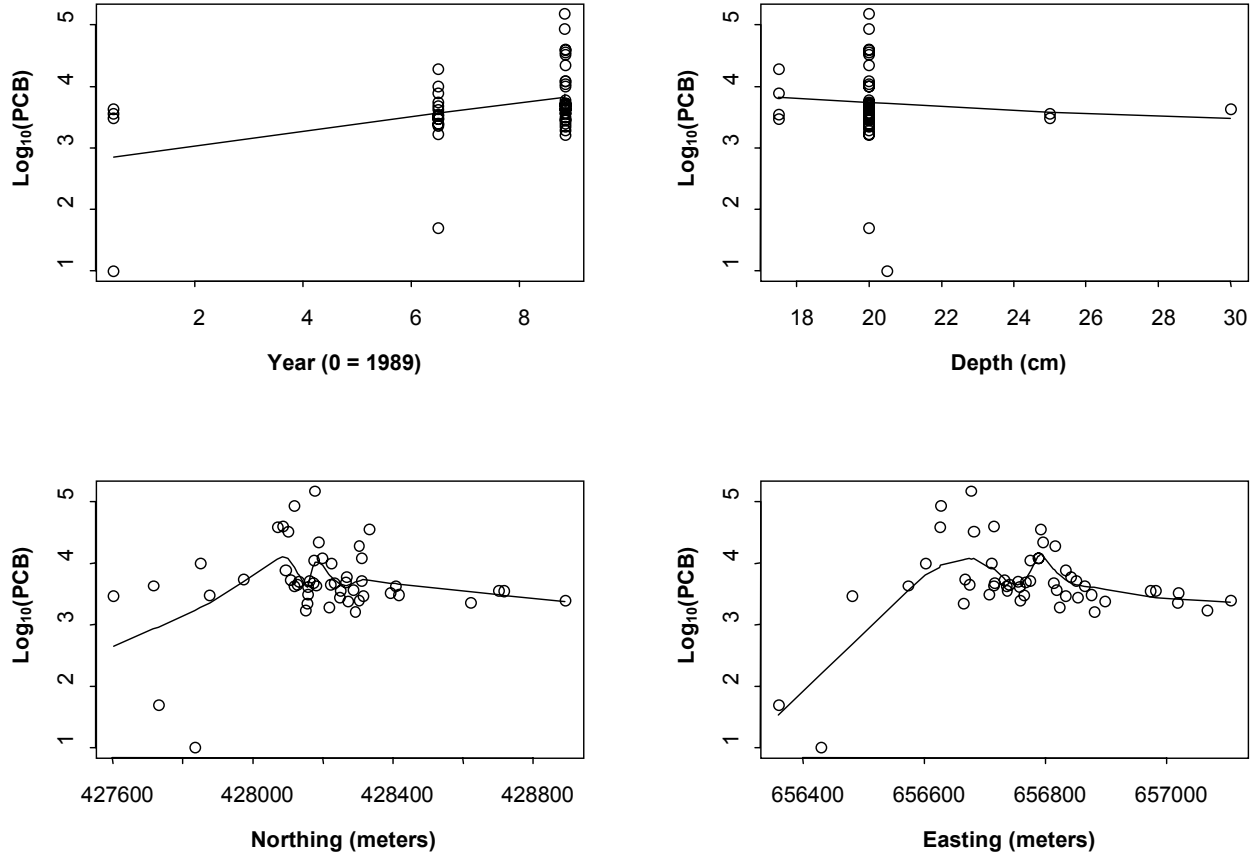


Figure A-84 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 5067 (10 to 30 cm) Including Fitted Smoothed Line

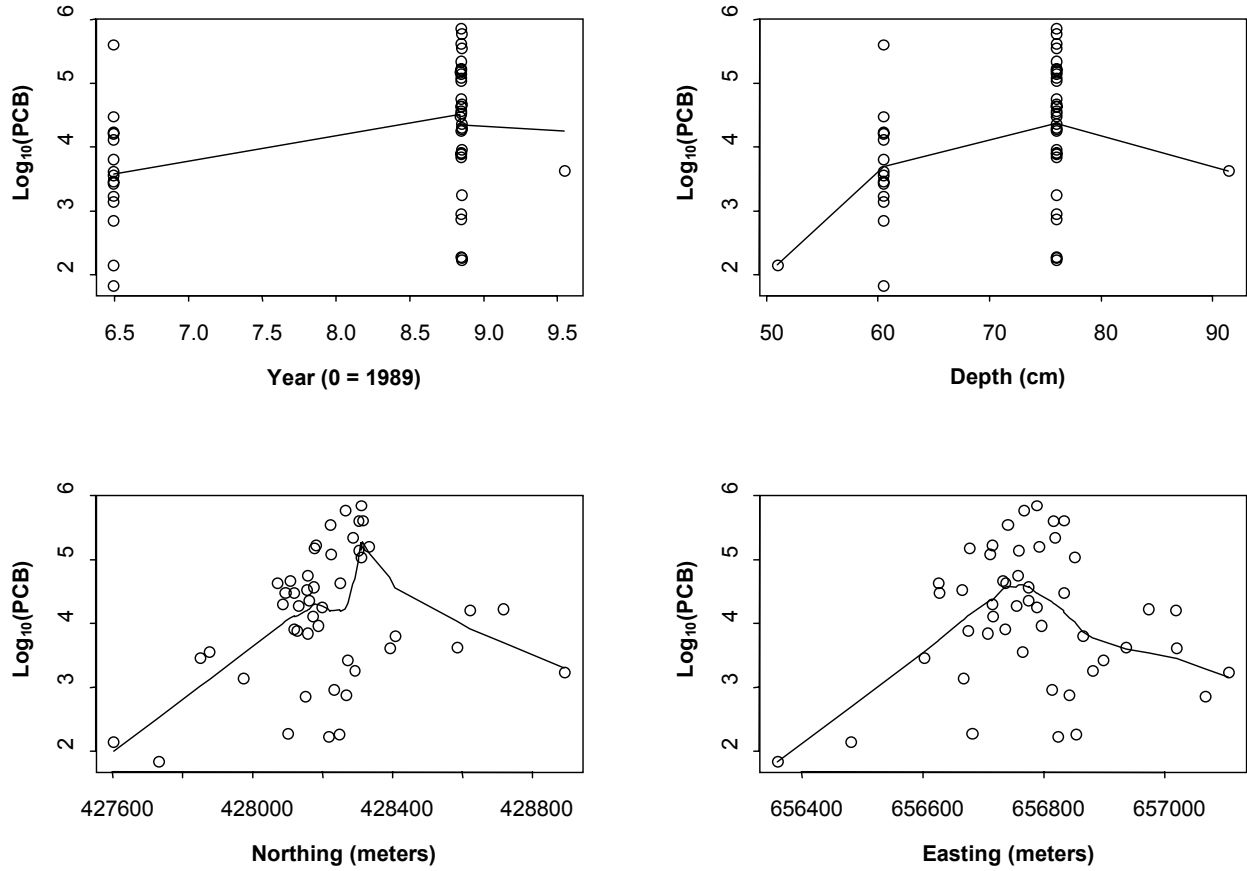


Figure A-85 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 5067 (50 to 100 cm) Including Fitted Smoothed Line

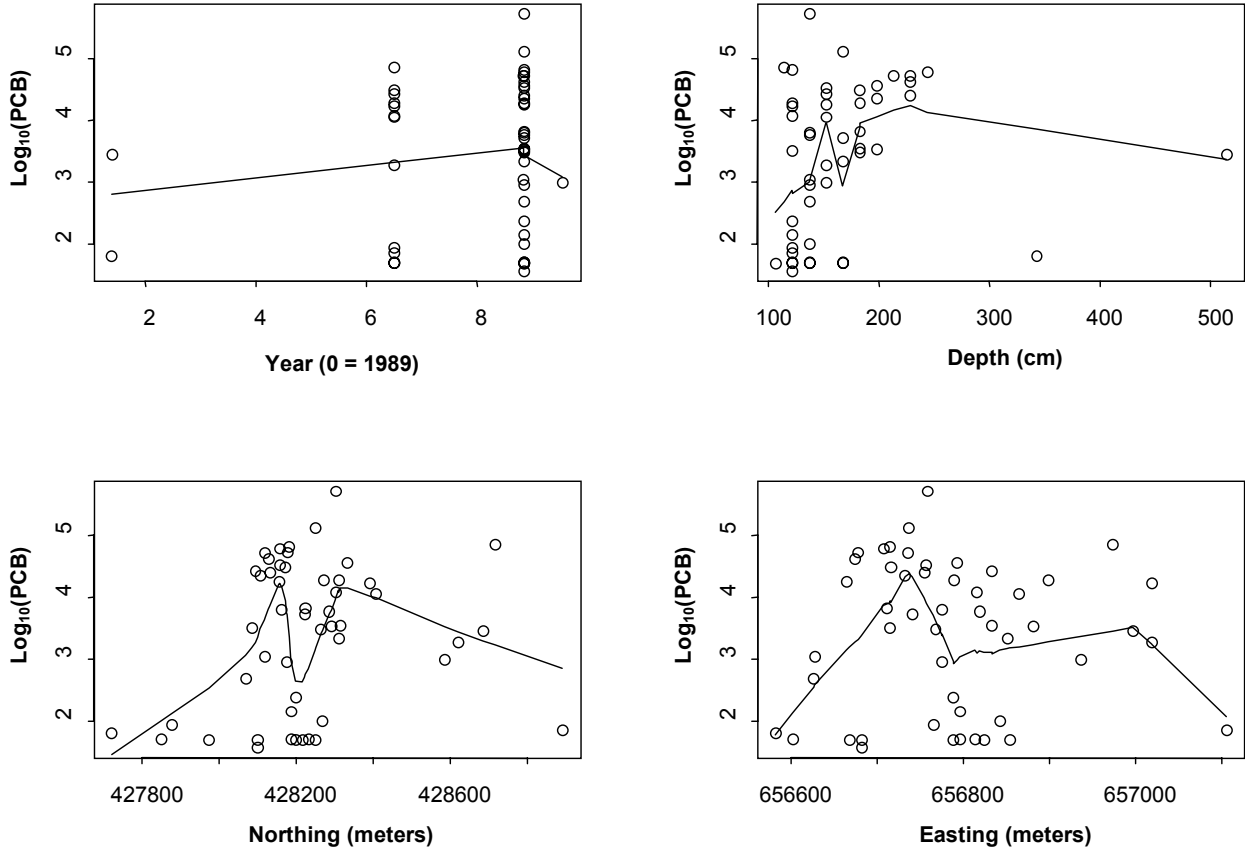


Figure A-86 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 5067 (100+ cm) Including Fitted Smoothed Line

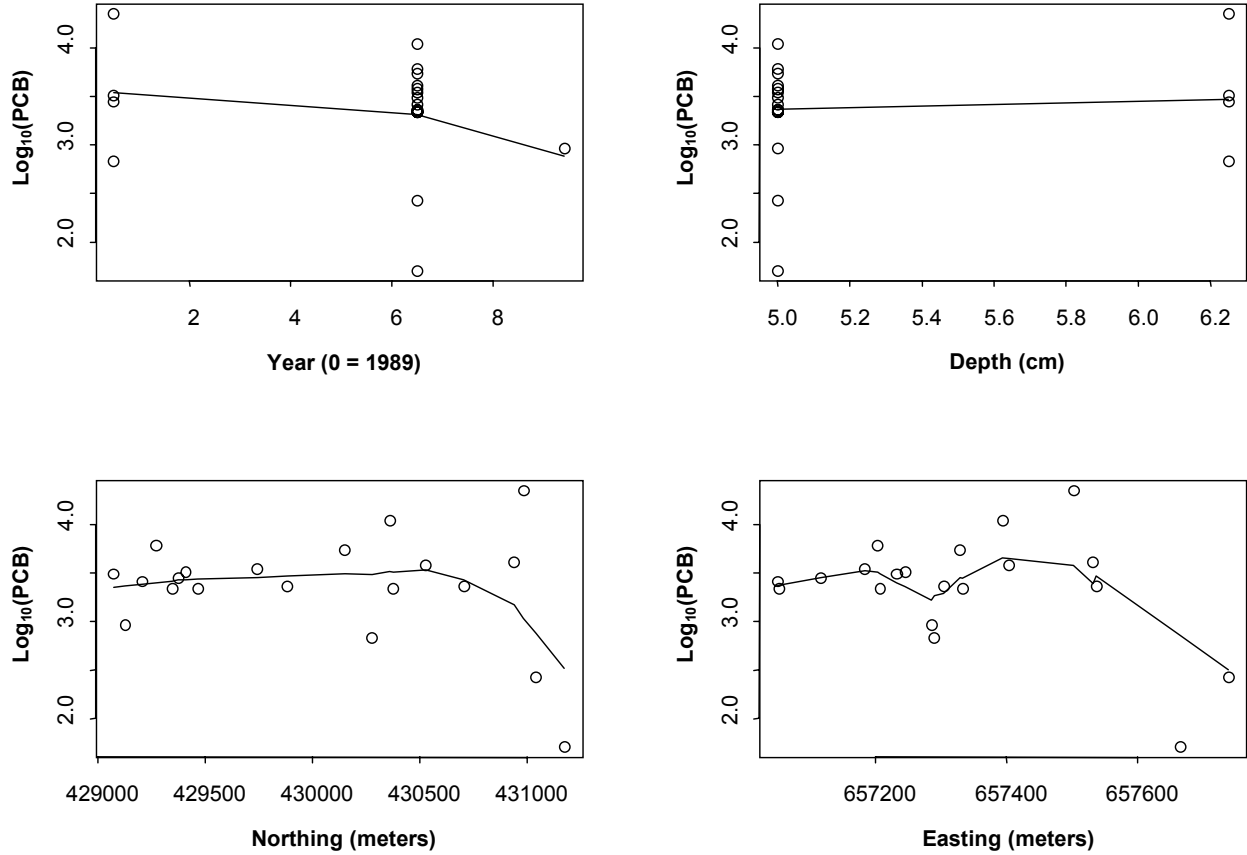


Figure A-87 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 6891 (0 to 10 cm) Including Fitted Smoothed Line

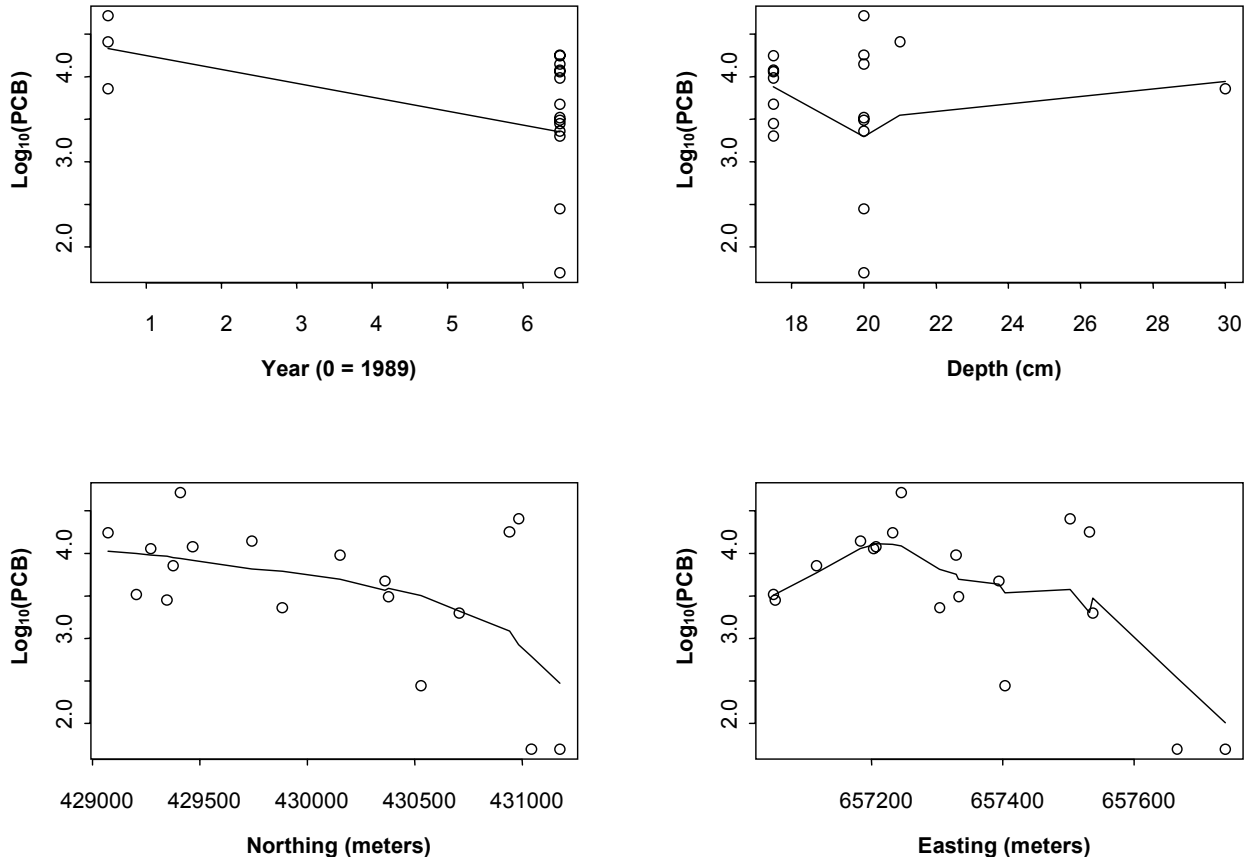


Figure A-88 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 6891 (10 to 30 cm) Including Fitted Smoothed Line

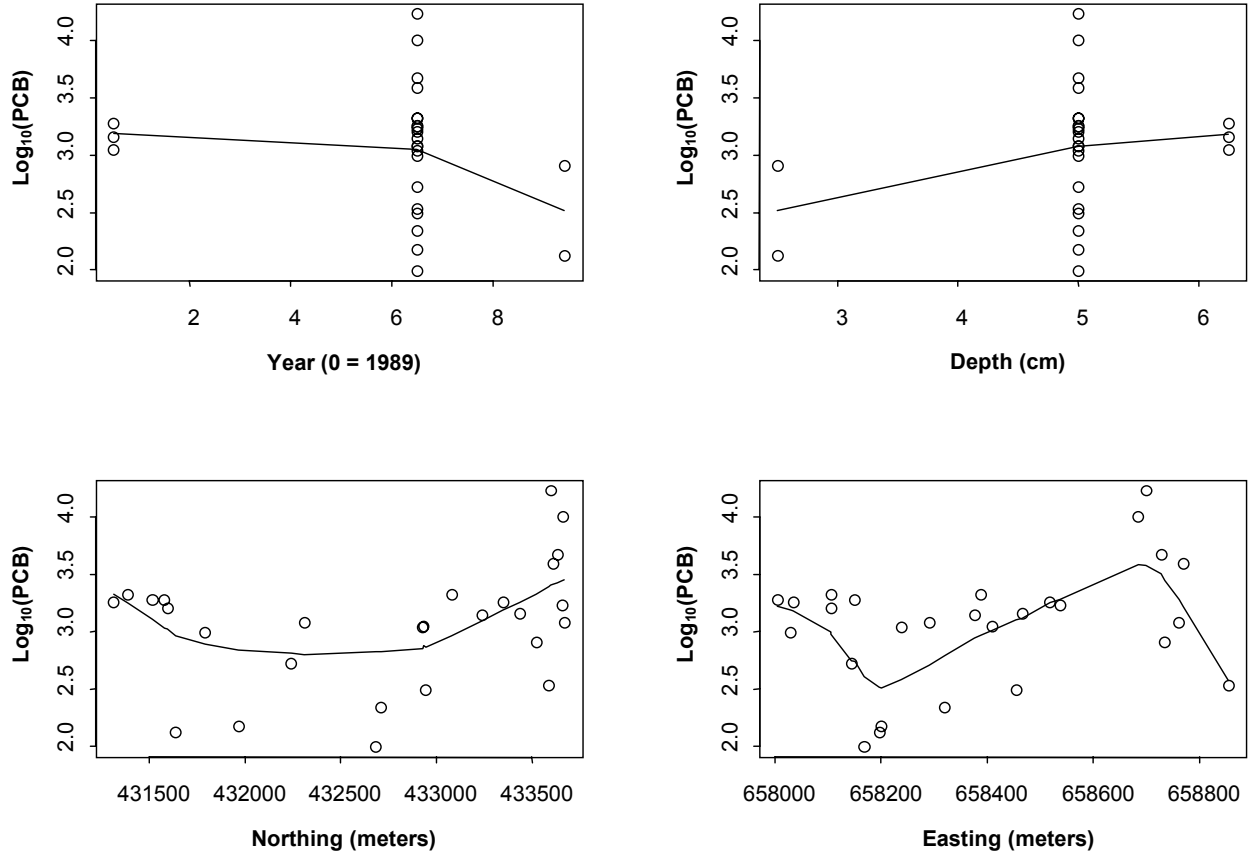


Figure A-89 Log_{10} PCB Concentration versus Time, Depth, Northing, and Easting for De Pere SMU Group 92115 (0 to 10 cm) Including Fitted Smoothed Line

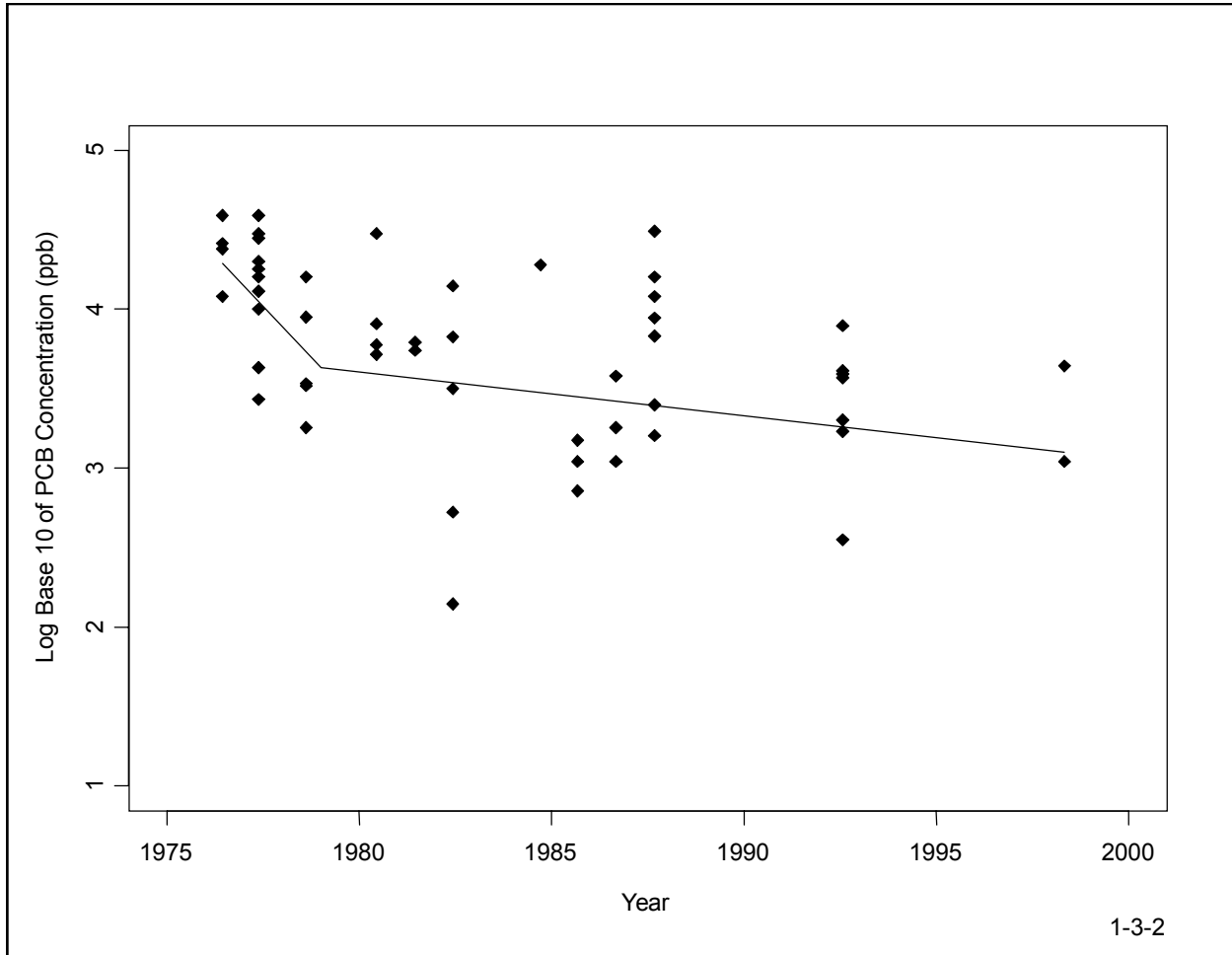


Figure A-90 Log_{10} PCB Concentration (ppb) in Little Lake Butte des Morts Carp, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ♦.

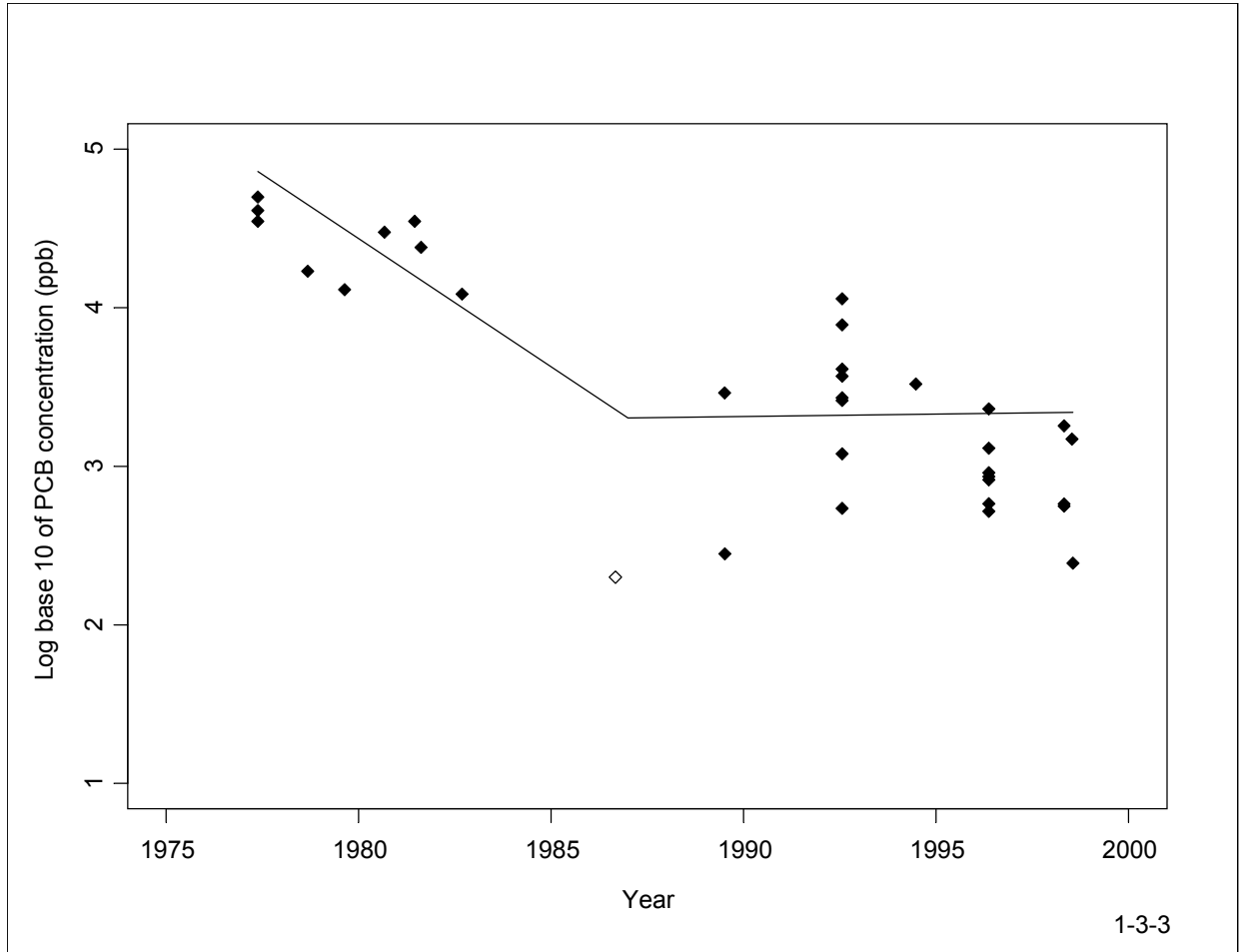


Figure A-91 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Carp, Whole Body, versus Time

Values at or above the detection limit are depicted as ◆. Any values below detection limit are depicted as ◇.

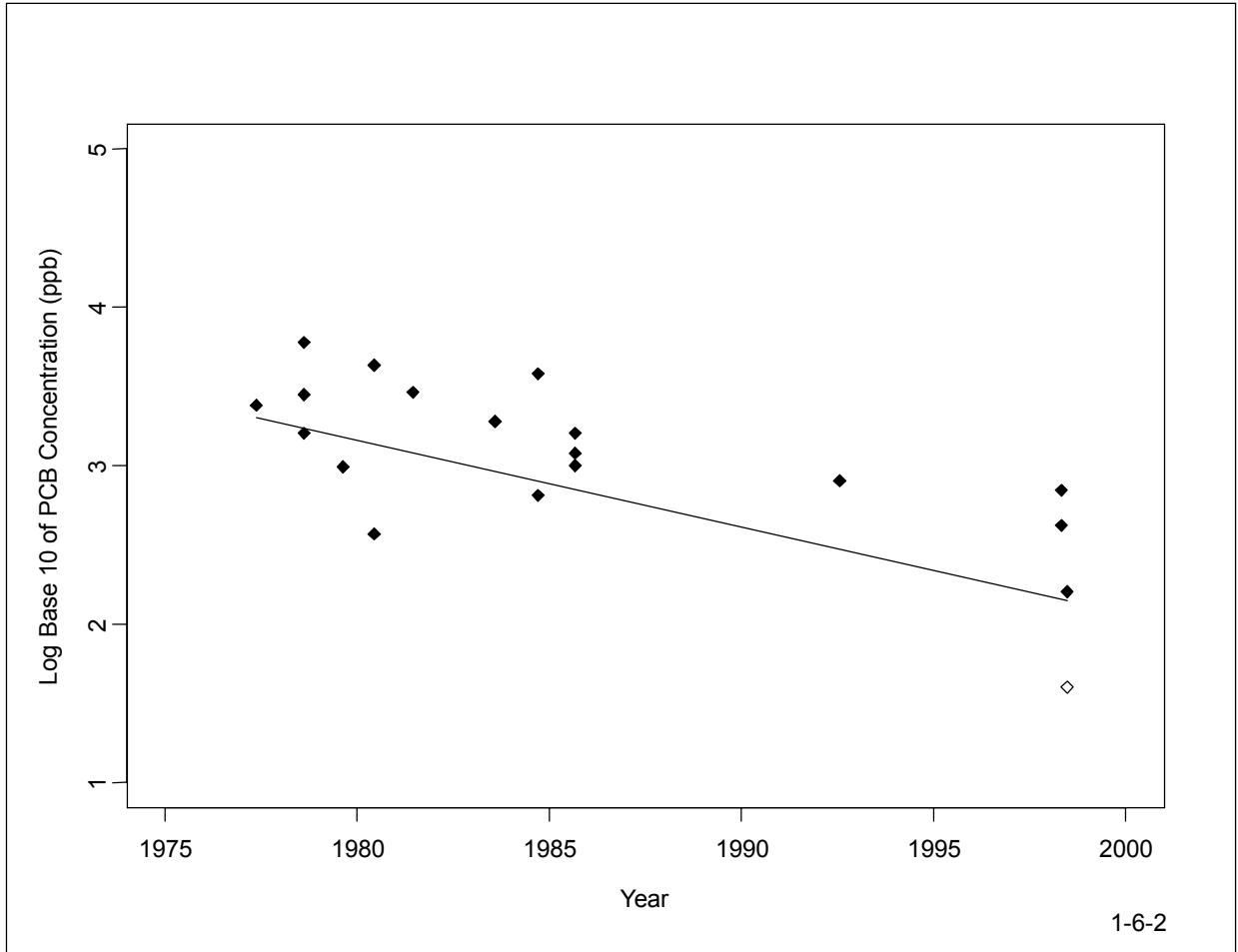


Figure A-92 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Northern Pike, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆. Any values below detection limit are depicted as ◇.

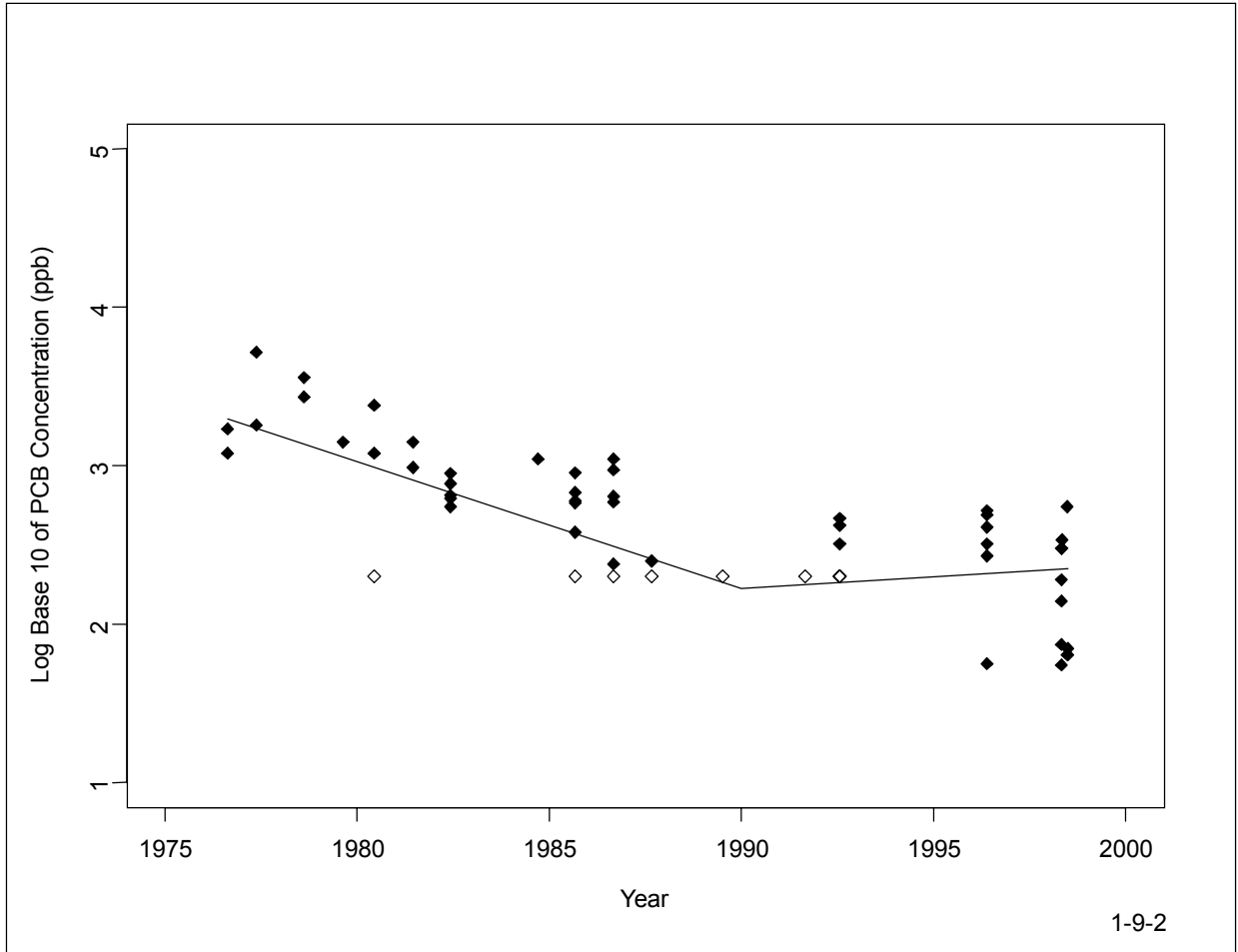


Figure A-93 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Walleye, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆. Any values below detection limit are depicted as ◇.

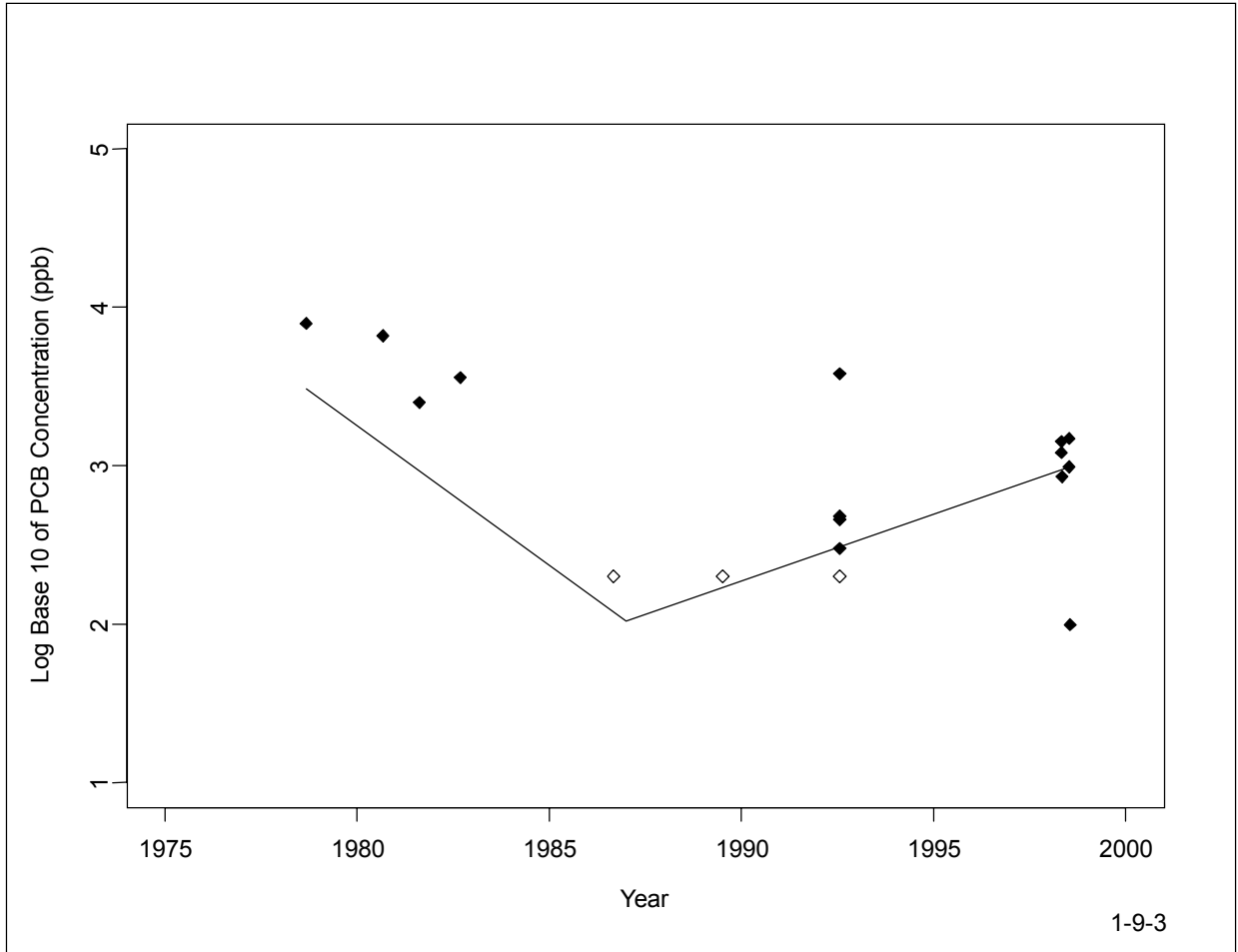


Figure A-94 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Walleye, Whole Body, versus Time

Values at or above the detection limit are depicted as ◆. Any values below detection limit are depicted as ◇.

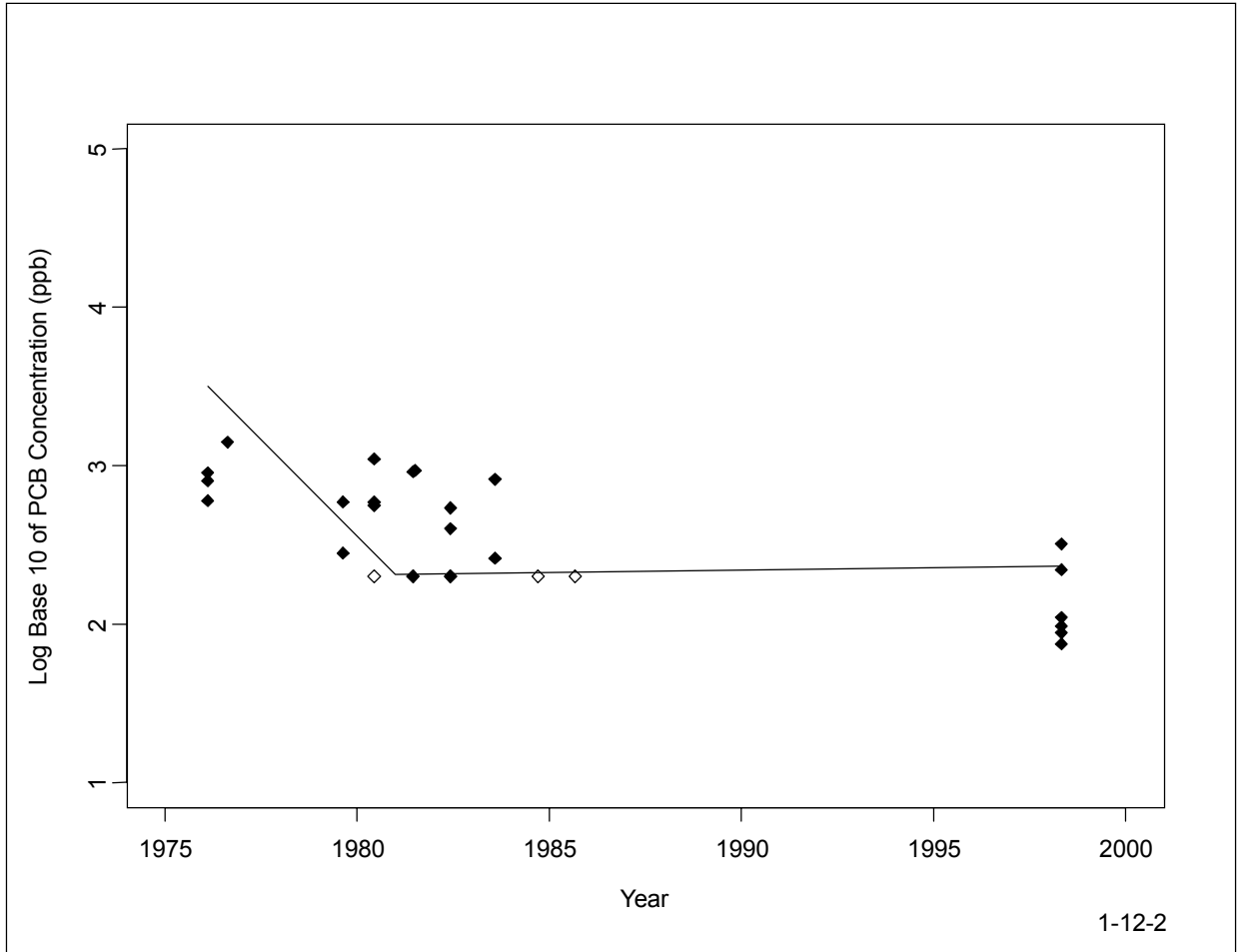


Figure A-95 Log₁₀ PCB Concentration (ppb) in Little Lake Butte des Morts Yellow Perch, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆. Any values below detection limit are depicted as ◇.

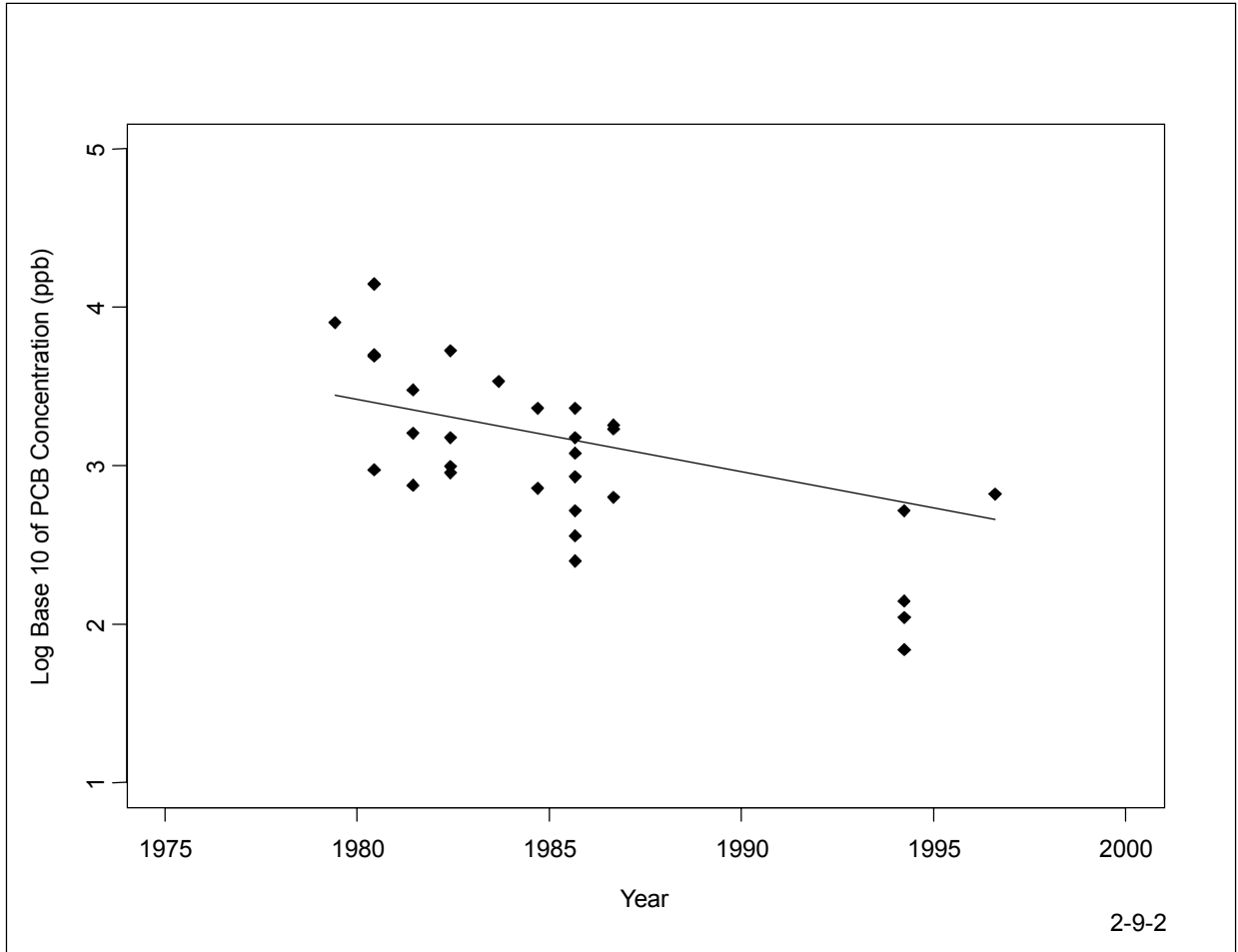


Figure A-96 Log₁₀ PCB Concentration (ppb) in Appleton to Little Rapids Walleye, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆.

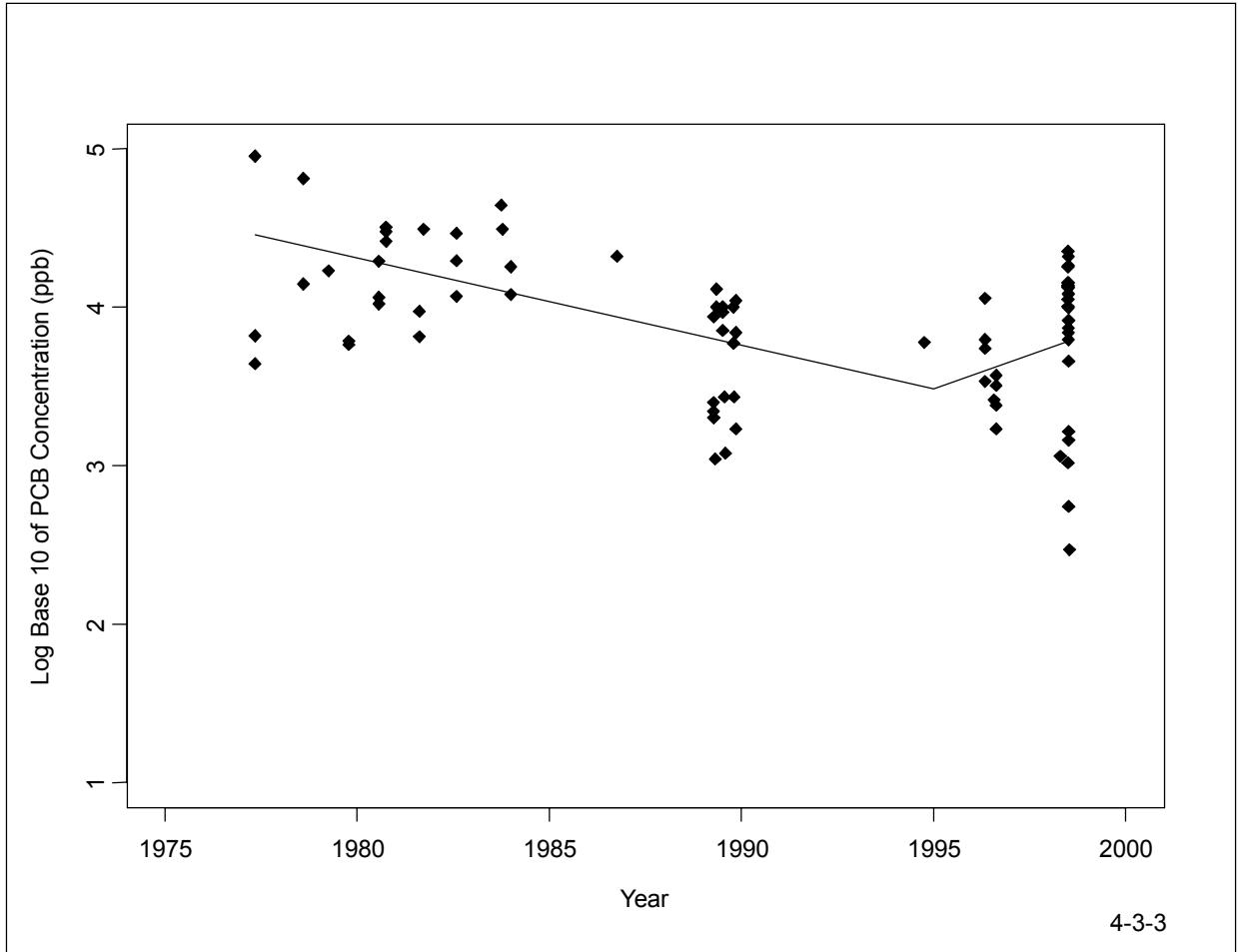


Figure A-97 Log₁₀ PCB Concentration (ppb) in De Pere to Green Bay Carp, Whole Body, versus Time

Values at or above the detection limit are depicted as ◆.

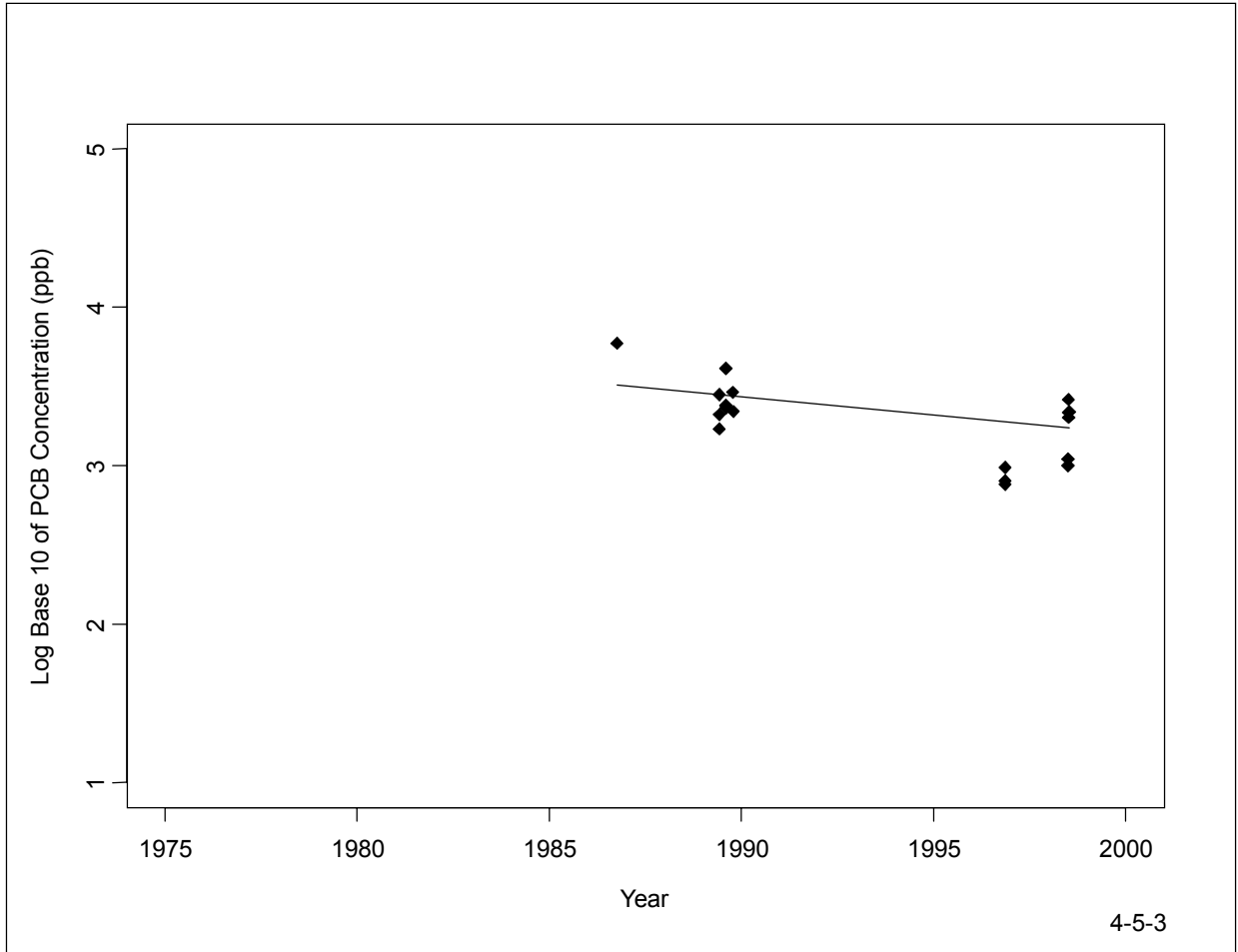


Figure A-98 Log_{10} PCB Concentration (ppb) in De Pere to Green Bay Gizzard Shad, Whole Body, versus Time

Values at or above the detection limit are depicted as \blacklozenge .

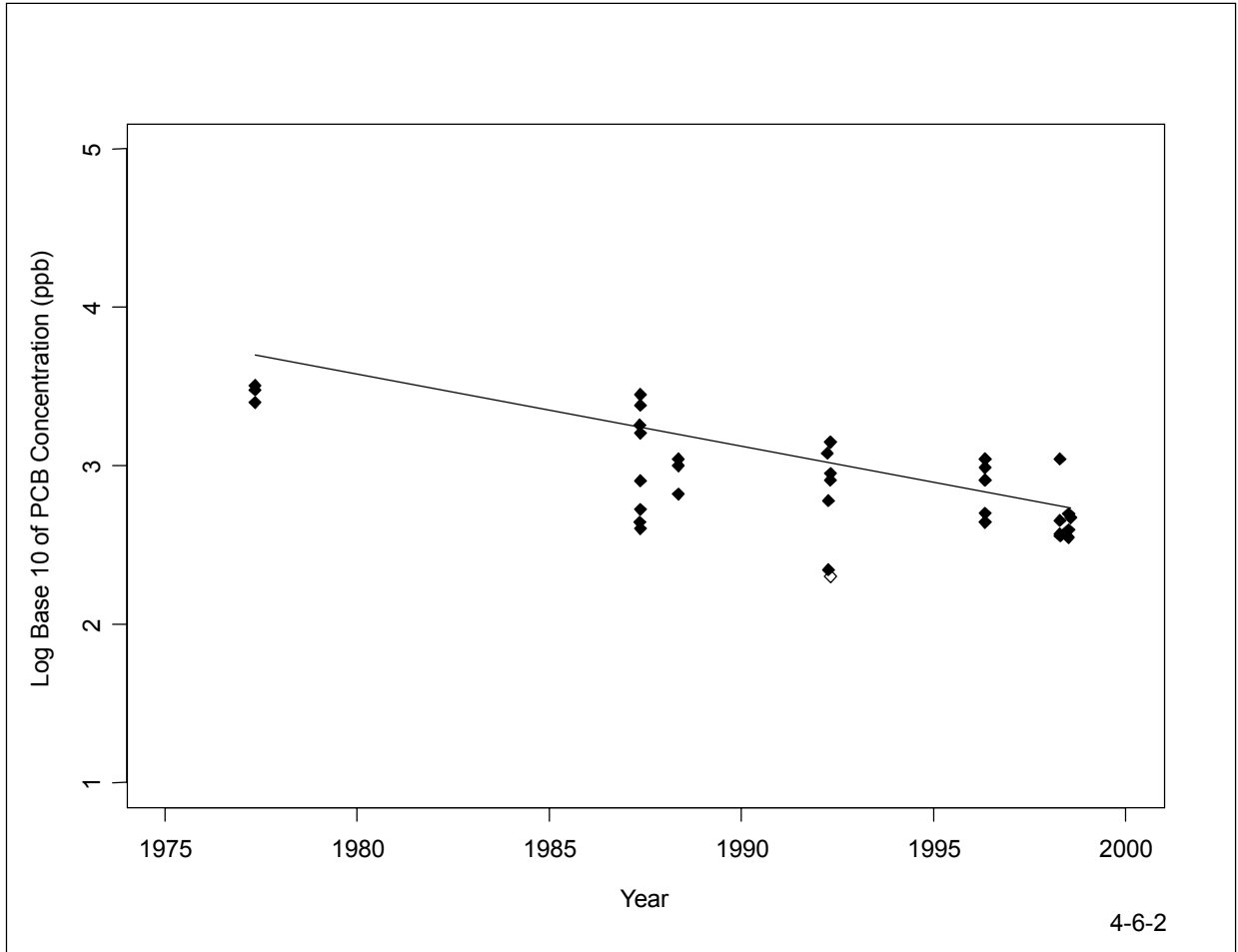


Figure A-99 Log₁₀ PCB Concentration (ppb) in De Pere to Green Bay Northern Pike, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆. Any values below detection limit are depicted as ◇.

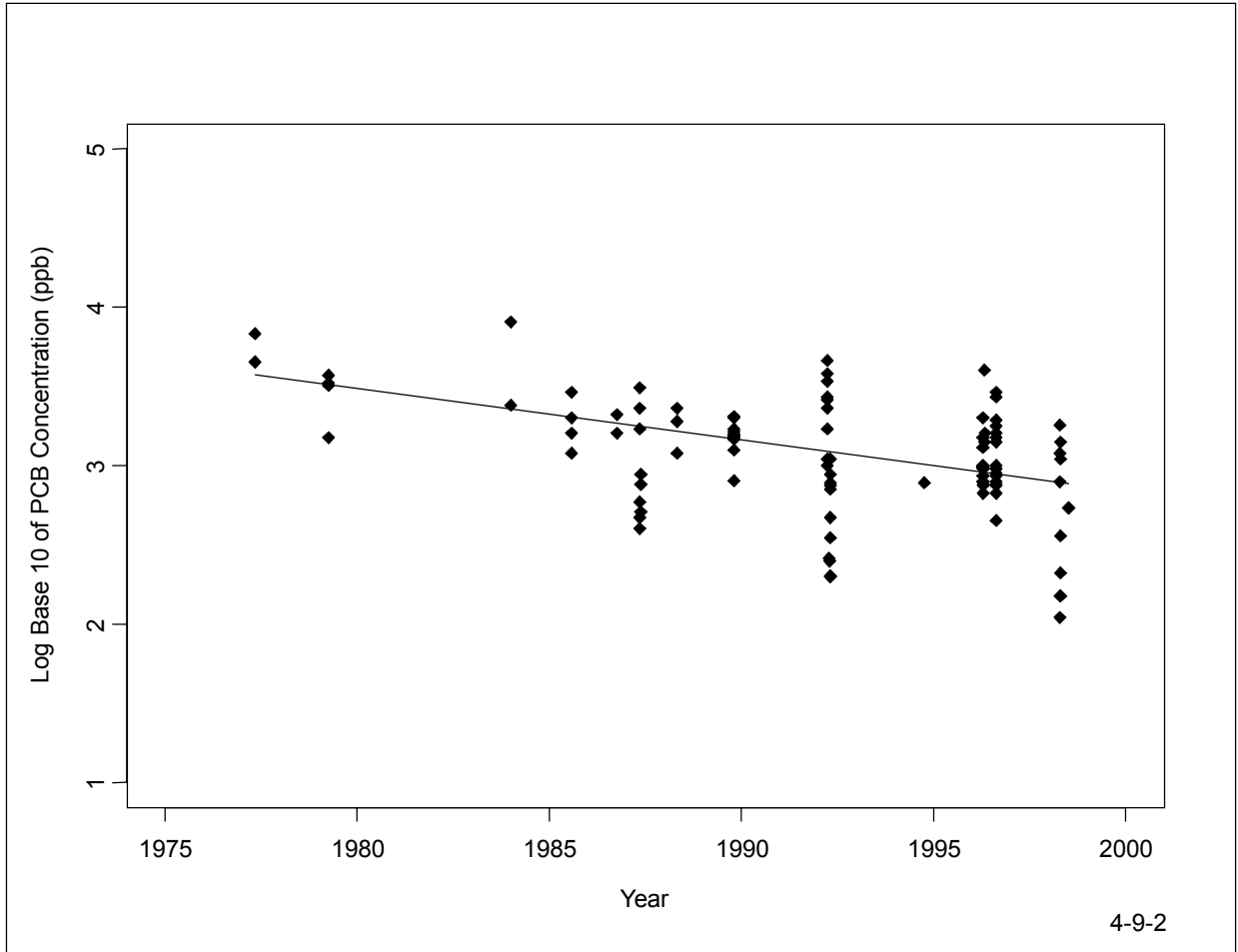


Figure A-100 Log₁₀ PCB Concentration (ppb) in De Pere to Green Bay Walleye, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆.

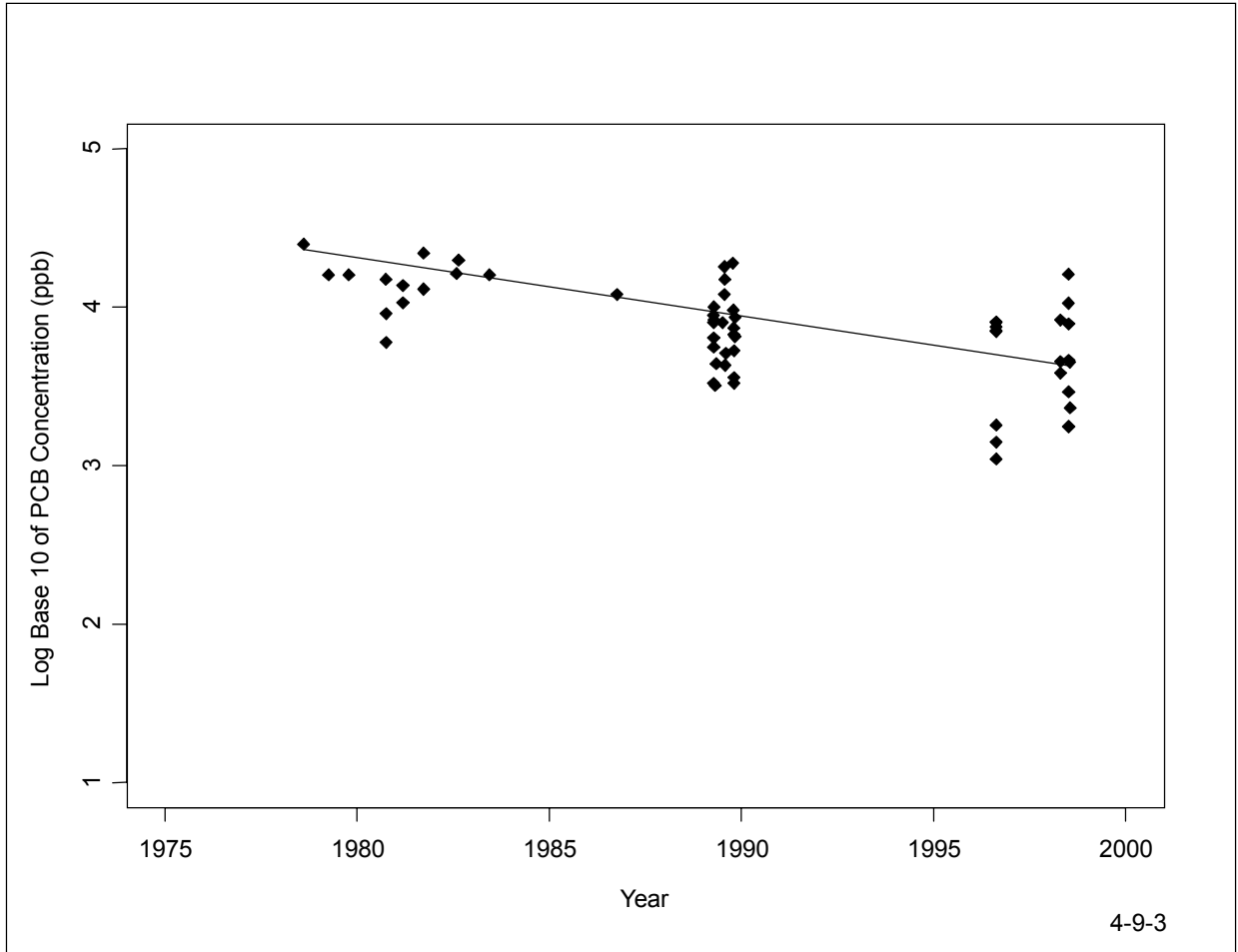


Figure A-101 Log₁₀ PCB Concentration (ppb) in De Pere to Green Bay Walleye, Whole Body, versus Time

Values at or above the detection limit are depicted as ◆.

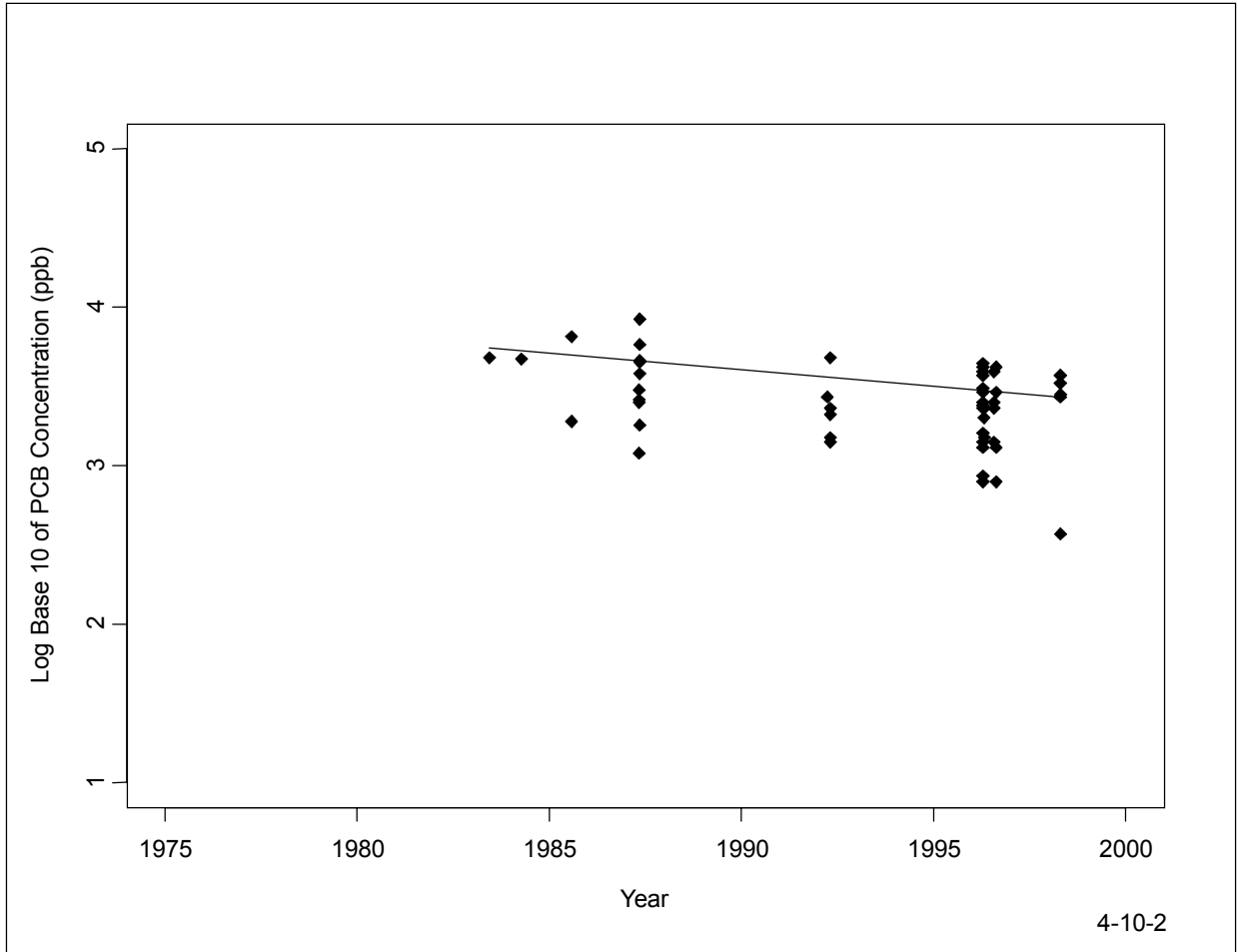


Figure A-102 Log₁₀ PCB Concentration (ppb) in De Pere to Green Bay White Bass, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆.

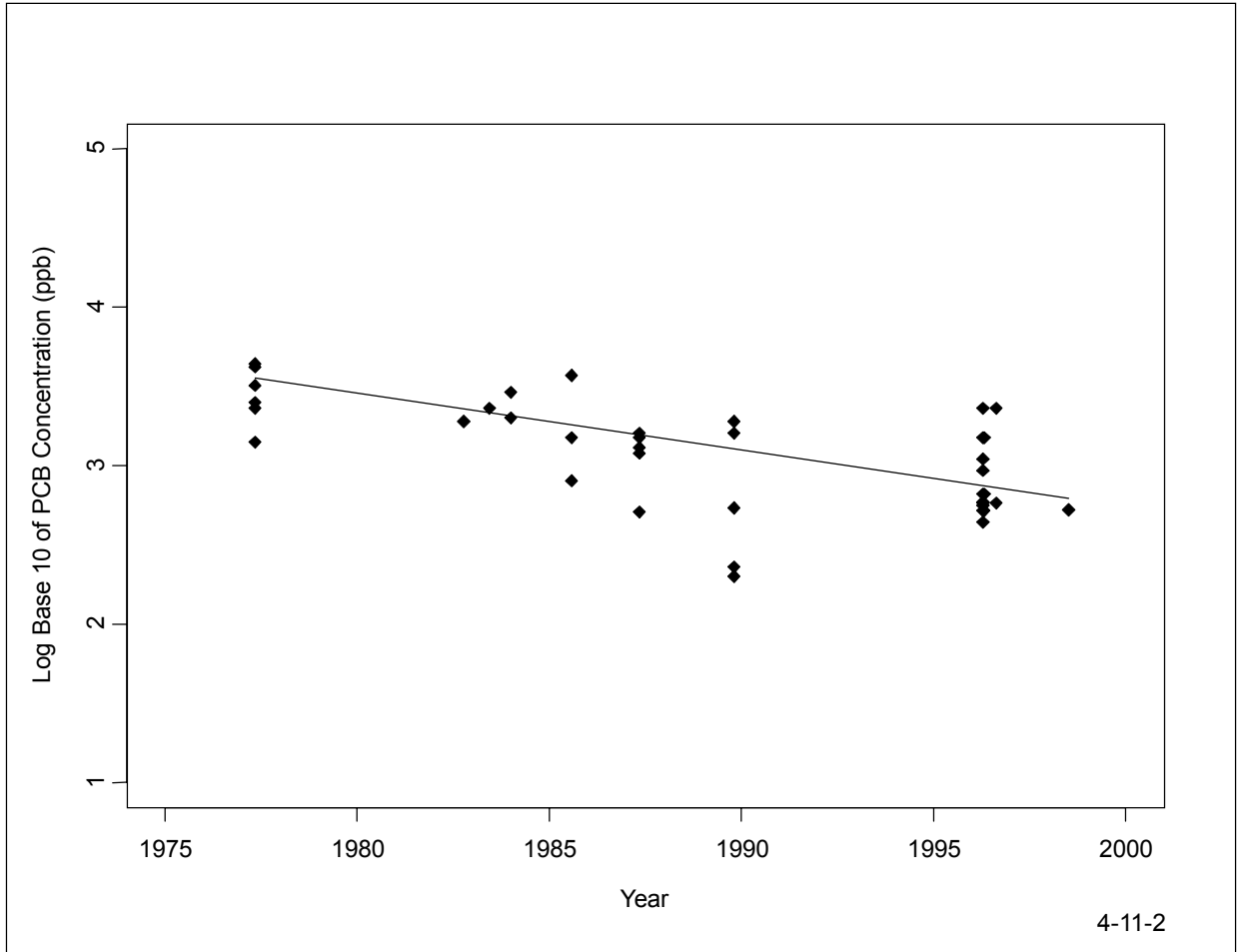


Figure A-103 Log₁₀ PCB Concentration (ppb) in De Pere to Green Bay White Sucker, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆.

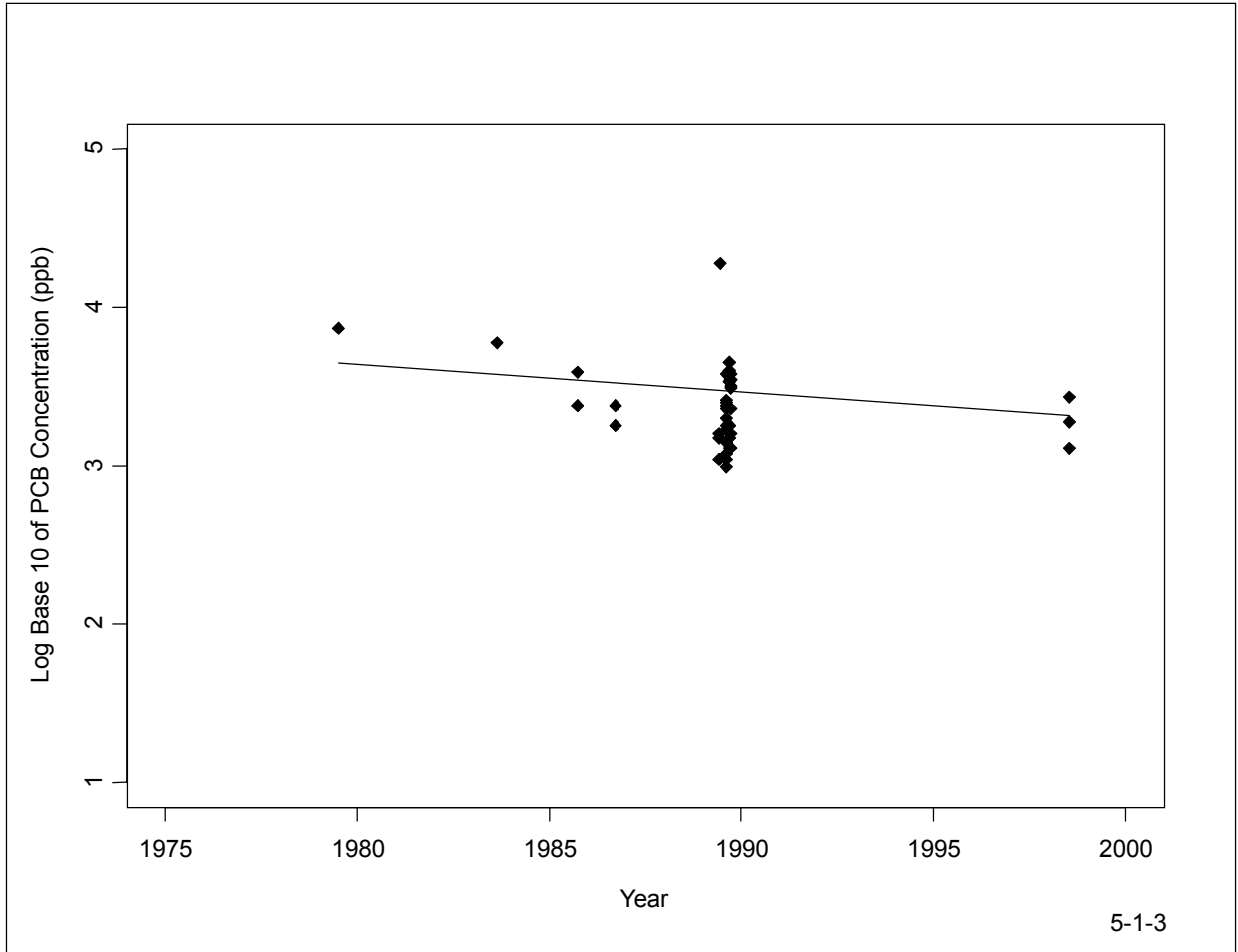


Figure A-104 Log₁₀ PCB Concentration (ppb) in Green Bay Zone 2 (2A and 2B) Alewife, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆.

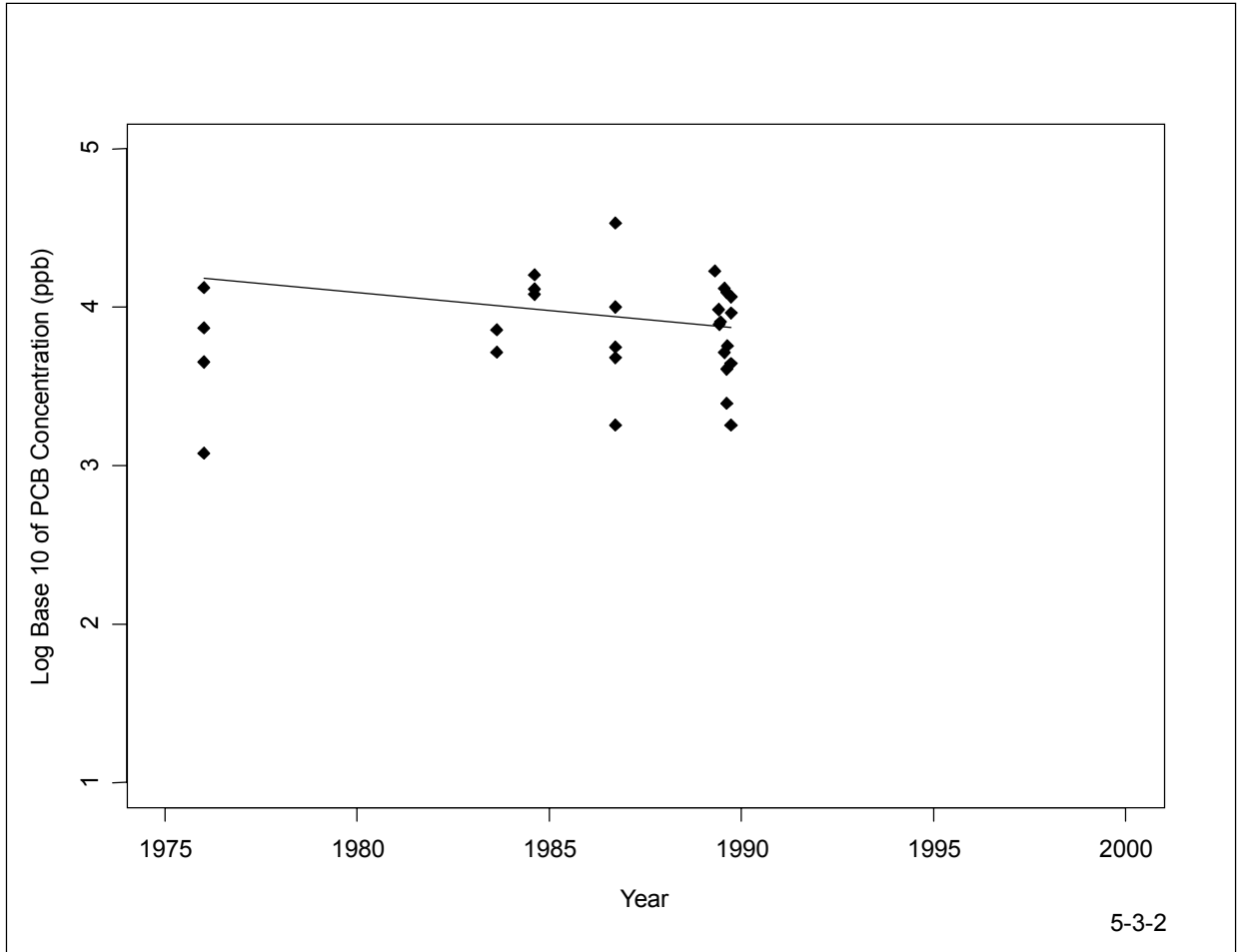


Figure A-105 Log₁₀ PCB Concentration (ppb) in Green Bay Zone 2 (2A and 2B) Carp, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆.

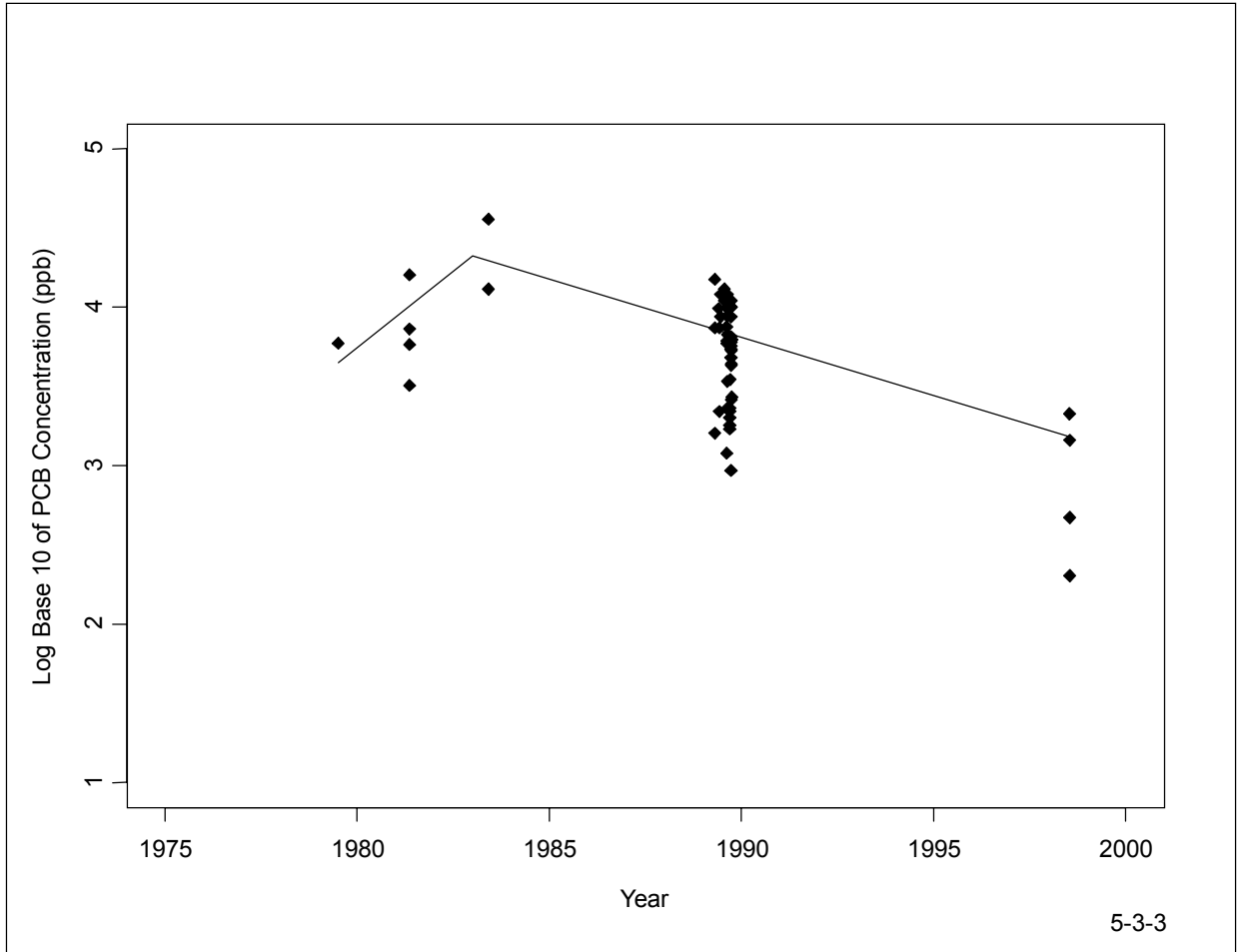


Figure A-106 Log₁₀ PCB Concentration (ppb) in Green Bay Zone 2 (2A and 2B) Carp, Whole Body, versus Time

Values at or above the detection limit are depicted as ◆.

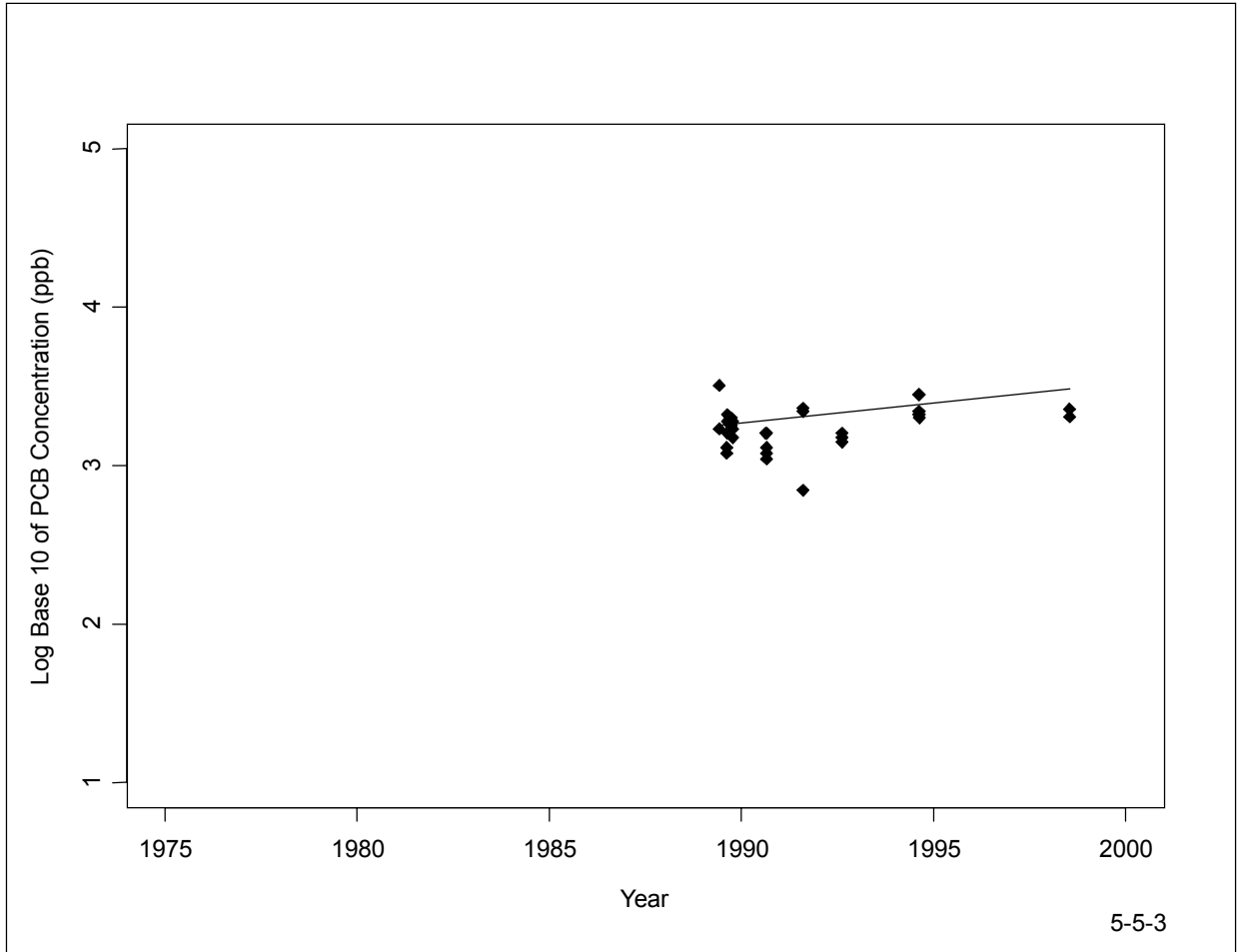


Figure A-107 Log₁₀ PCB Concentration (ppb) in Green Bay Zone 2 (2A and 2B) Gizzard Shad, Whole Body, versus Time

Values at or above the detection limit are depicted as ♦.

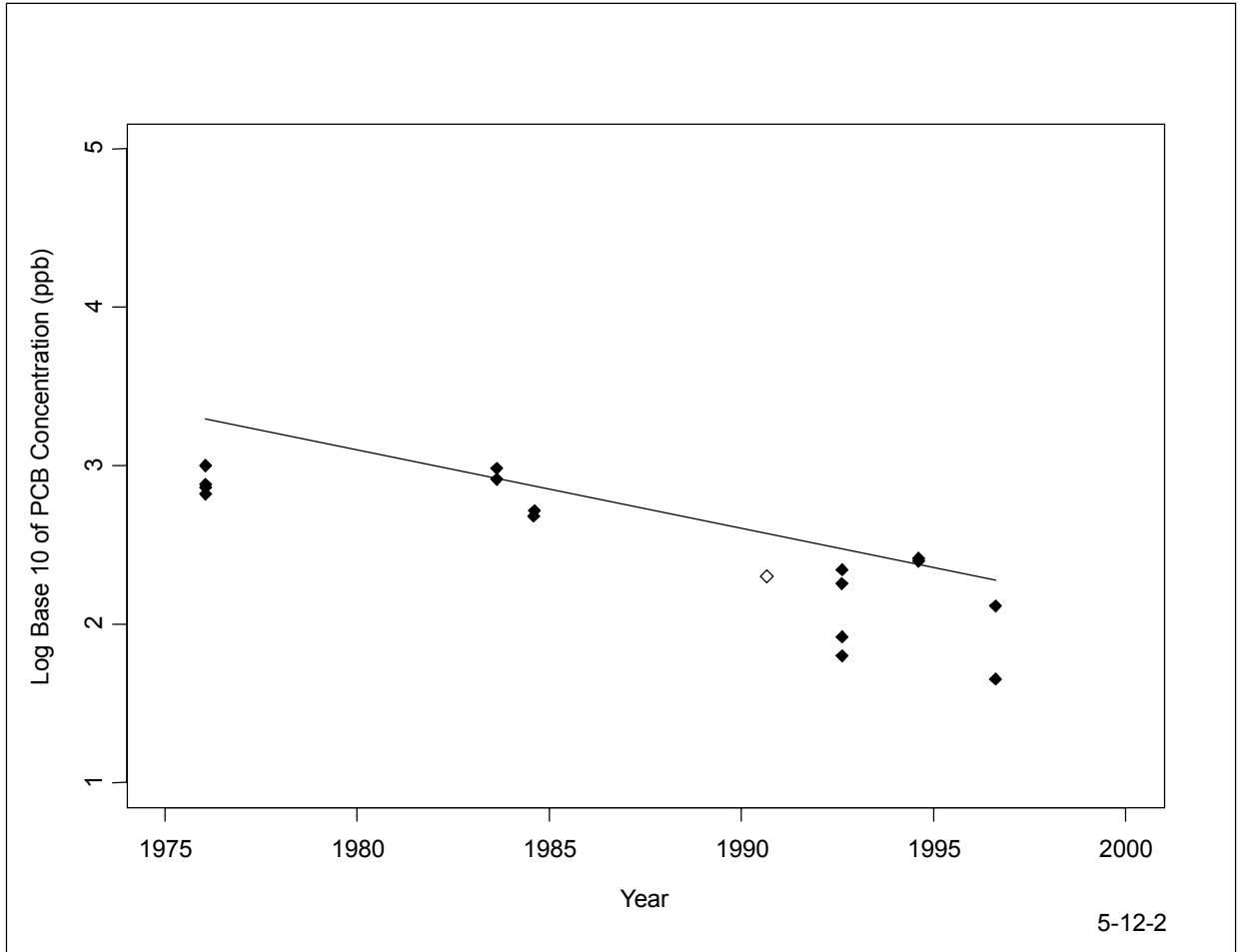


Figure A-108 Log₁₀ PCB Concentration (ppb) in Green Bay Zone 2 (2A and 2B) Yellow Perch, Skin-on Fillet, versus Time

Values at or above the detection limit are depicted as ◆. Any values below detection limit are depicted as ◇.

Table A-1 Details of Models Fitted to Time Trends in Sediment PCB Concentrations

| Reach and Deposit Group | Depth Range (cm) | R-squared | | | Sill Distance | Intercept Parameter Est. | WSEV Std. Err. of Intercept | Std. Err. of Intercept Based on Independence | Skewness of Untransformed PCB Conc. | Skewness of Log ₁₀ (PCB) Conc. | PCB Mass (kg) | Normal Scale (Std. Dev.) Est. |
|------------------------------------|------------------|---------------------------|-------------------|--------------------|---------------|--------------------------|-----------------------------|--|-------------------------------------|---|---------------|-------------------------------|
| | | Geographic Variables Only | Geographic + Time | Change Due to Time | | | | | | | | |
| <i>Little Lake Butte des Morts</i> | | | | | | | | | | | | |
| AB | 0-10 | 0.47 | 0.55 | 0.08 | 25 | 4.4461 | 0.3237 | 0.2788 | 1.74 | -1.13 | 71.7 | 0.465933 |
| | 10-30 | 0.17 | 0.17 | 0.00 | 25 | 4.0797 | 0.8357 | 0.6054 | 3.37 | -0.09 | 217.7 | 1.02534 |
| | 30-50 | 0.36 | 0.37 | 0.01 | 0 | 10.4324 | 2.8100 | 2.1917 | 4.52 | 1.03 | 328.3 | 1.1568 |
| C | 0-10 | 0.27 | 0.47 | 0.21 | 25 | 5.2096 | 1.2084 | 1.0586 | 1.97 | 0.19 | 25.4 | 0.336235 |
| | 10-30 | 0.55 | 0.69 | 0.14 | 25 | 5.0070 | 1.4441 | 1.3930 | 2.80 | 0.76 | 14.6 | 0.897603 |
| POG | 0-10 | 0.61 | 0.71 | 0.10 | 75 | 4.4765 | 0.5067 | 0.4769 | 1.33 | 0.34 | 113.5 | 0.425786 |
| | 10-30 | 0.67 | 0.78 | 0.10 | 0 | 3.8807 | 0.6868 | 0.3776 | 1.92 | -1.18 | 32.1 | 0.2376 |
| D | 0-10 | 0.19 | 0.80 | 0.61 | 0 | 2.2285 | 0.5288 | 0.5202 | 2.96 | -0.25 | 55.5 | 0.397127 |
| | 10-30 | 0.24 | 0.30 | 0.05 | 50 | 3.5528 | 0.3827 | 0.4099 | 2.40 | -0.52 | 142.5 | 0.520421 |
| F | 0-10 | 0.23 | 0.31 | 0.08 | 50 | 2.2040 | 1.3533 | 1.0844 | 5.15 | 0.97 | 180.1 | 0.789297 |
| | 10-30 | 0.02 | 0.61 | 0.59 | 0 | 3.1032 | 0.3153 | 0.3176 | 0.14 | -1.27 | 15.7 | 0.439535 |
| <i>Appleton</i> | | | | | | | | | | | | |
| IMOR | 0-10 | 0.09 | 0.41 | 0.32 | 0 | 3.1269 | 0.4735 | 0.4747 | 3.44 | 0.31 | 6.9 | 0.583018 |
| | 10-30 | 0.68 | 0.70 | 0.02 | 0 | 4.2292 | 0.4199 | 0.3549 | 1.14 | -0.52 | 6.9 | 0.326511 |
| | 30-50 | 0.43 | 0.48 | 0.05 | 50 | 3.7450 | 0.6539 | 0.6366 | 2.66 | -0.98 | 11.5 | 0.615759 |
| N Pre-dredge | 0-10 | 0.49 | 0.56 | 0.07 | 10 | 4.4070 | 1.5119 | 1.2267 | 1.00 | -2.56 | 4.9 | 0.570745 |
| | 10-30 | 0.14 | 0.31 | 0.17 | 0 | 3.2202 | 0.3490 | 0.2537 | 2.55 | 0.34 | 5.2 | 0.524406 |
| | 30-50 | 0.12 | 0.56 | 0.44 | 0 | 4.1303 | 0.6783 | 0.7806 | 4.76 | 0.99 | 2.9 | 0.734058 |
| VCC | 0-10 | 0.46 | 0.52 | 0.06 | 0 | 4.4304 | 0.5727 | 0.5713 | 1.05 | 0.06 | 0.9 | 0.11942 |
| | 10-30 | | | | | | | | | | | |
| | 30-50 | | | | | | | | | | | |
| <i>Little Rapids</i> | | | | | | | | | | | | |
| Upper EE | 0-10 | 0.09 | 0.16 | 0.06 | 0 | 3.2722 | 0.7469 | 0.4948 | 3.43 | 0.21 | 85.0 | 0.58418 |
| | 10-30 | 0.17 | 0.38 | 0.22 | 0 | 2.5703 | 1.1521 | 0.8651 | 4.06 | 0.51 | 46.4 | 0.822143 |
| | 30-50 | 0.03 | 0.24 | 0.22 | 200 | 4.7214 | 1.3448 | 1.7186 | 3.44 | 0.77 | 4.3 | 0.678349 |
| Lower EE | 0-10 | 0.36 | 0.52 | 0.16 | 0 | 2.9308 | 0.2663 | 0.3268 | 3.68 | 0.37 | 25.4 | 0.486326 |
| | 10-30 | 0.17 | 0.40 | 0.23 | 0 | 2.8576 | 0.7657 | 0.9180 | 4.97 | 0.80 | 13.2 | 0.96465 |
| | 30-50 | 0.47 | 0.56 | 0.09 | 0 | 5.0328 | 0.9549 | 1.1745 | 1.76 | 0.26 | 4.6 | 0.357574 |
| FF | 0-10 | 0.15 | 0.20 | 0.05 | 0 | 3.7208 | 0.3852 | 0.4231 | 1.52 | -0.24 | 36.7 | 0.83476 |
| | 10-30 | 0.07 | 0.25 | 0.18 | 0 | 2.1741 | 1.3609 | 1.2502 | 4.02 | 0.77 | 14.6 | 1.12086 |
| | 30-50 | 0.29 | 0.33 | 0.04 | 0 | 2.8846 | 0.7084 | 0.2893 | 1.47 | -0.38 | 131.6 | 0.50908 |
| GGHH | 0-10 | 0.12 | 0.12 | 0.00 | 0 | 3.3231 | 0.8171 | 0.9167 | 1.33 | -0.22 | 289.6 | 0.91031 |
| | 10-30 | 0.10 | 0.19 | 0.09 | 0 | 0.0821 | 2.8431 | 1.3045 | 1.33 | -0.74 | 271.4 | 0.964739 |
| | 30-50 | 0.16 | 0.23 | 0.07 | 0 | 1.4499 | 1.9204 | 1.2885 | 4.82 | 0.74 | 195.7 | 0.8449 |
| | 100+ | 0.62 | 0.72 | 0.09 | 0 | 2.3137 | 0.5420 | 0.4451 | 2.86 | 0.53 | 21.4 | 0.295787 |
| <i>De Pere</i> | | | | | | | | | | | | |
| SMU Group 2025 | 0-10 | 0.38 | 0.46 | 0.07 | 0 | 3.6631 | 0.4655 | 0.4255 | 1.11 | -1.18 | 225.6 | 0.350891 |
| | 10-30 | 0.35 | 0.37 | 0.02 | 100 | 6.3342 | 3.4691 | 2.2114 | 2.58 | -0.59 | 813.6 | 0.855251 |
| | 30-50 | 0.66 | 0.76 | 0.10 | 150 | 5.5480 | 0.9776 | 1.1642 | 1.76 | -0.76 | 950.3 | 0.430459 |
| | 50-100 | 0.35 | 0.36 | 0.01 | 50 | 4.0031 | 1.1675 | 1.2707 | 1.76 | -0.18 | 1569.3 | 1.13947 |
| 2649 | 0-10 | 0.06 | 0.17 | 0.11 | 0 | 3.2501 | 0.2065 | 0.4161 | 0.89 | -1.01 | 356.8 | 0.434768 |
| | 10-30 | 0.31 | 0.43 | 0.12 | 0 | 10.6240 | 3.2452 | 1.9813 | 0.80 | -0.90 | 1556.5 | 0.816451 |
| | 50-100 | 0.13 | 0.13 | 0.00 | 100 | 3.6653 | 2.3249 | 1.1267 | 1.32 | -0.47 | 3135.5 | 1.07814 |
| 5067 | 100+ | 0.20 | 0.22 | 0.02 | 0 | 1.2186 | 1.9141 | 1.2818 | 5.18 | 0.10 | 1717.6 | 1.05288 |
| | 0-10 | 0.13 | 0.27 | 0.14 | 0 | 7.6178 | 1.2394 | 1.1333 | 7.47 | 2.40 | 92.4 | 0.186359 |
| | 10-30 | 0.42 | 0.47 | 0.05 | 0 | 2.4000 | 1.4903 | 1.2775 | 4.35 | -1.43 | 353.7 | 0.472972 |
| | 50-100 | 0.42 | 0.43 | 0.01 | 0 | 6.5635 | 2.1819 | 1.5704 | 2.61 | -0.36 | 2764.9 | 0.778337 |
| 6891 | 100+ | 0.26 | 0.29 | 0.02 | 0 | 4.9240 | 2.3655 | 1.8648 | 5.97 | -0.22 | 4426.0 | 1.13022 |
| | 0-10 | 0.42 | 0.46 | 0.04 | 100 | 10.2963 | 4.2471 | 5.8601 | 3.04 | -1.34 | 72.1 | 0.422776 |
| 92115 | 10-30 | 0.63 | 0.74 | 0.11 | 100 | 6.4202 | 1.3240 | 1.3665 | 2.29 | -1.22 | 246.7 | 0.447153 |
| | 0-10 | 0.52 | 0.52 | 0.01 | 0 | 0.8839 | 0.9748 | 1.1169 | 3.37 | -0.12 | 37.1 | 0.359379 |

Table A-1 Details of Models Fitted to Time Trends in Sediment PCB Concentrations

| Reach and Deposit Group | Depth Range (cm) | Number of Samples | | | | | Mean of Within-core-avg. Sample Variances | Variance of Singleton Samples in Core-avg. Data Set | Parameter Estimates and Standard Errors | | | | | |
|------------------------------------|------------------|---------------------------------------|--------------------------------------|-----------------------|--|--|---|---|---|--------------------------|----------------------------------|---------------|---------------------|-----------------------------|
| | | Single Used in Core-averaged Analyses | Core-avg. Used in Core-avg. Analyses | Total Original Single | Total in Core-avg. Analyses (mixed, single, & core-avg.) | Avg. that Ended up in a Core-avg. Sample | | | Intercept Estimate | WSEV Std. Err. Intercept | Independence Std. Err. Intercept | Time Estimate | WSEV Std. Err. Time | Independence Std. Err. Time |
| <i>Little Lake Butte des Morts</i> | | | | | | | | | | | | | | |
| AB | 0-10 | 47 | 20 | 94 | 67 | 2.4 | 0.0632 | 0.4984 | 4.4461 | 0.3237 | 0.2788 | -0.0970 | 0.0348 | 0.0279 |
| | 10-30 | 87 | 18 | 134 | 105 | 2.6 | 0.4082 | 1.0782 | 4.0797 | 0.8357 | 0.6054 | -0.0213 | 0.0647 | 0.0501 |
| | 30-50 | 52 | 2 | 56 | 54 | 2.0 | 1.4924 | 0.7630 | 10.4324 | 2.8100 | 2.1917 | -0.0144 | 0.1113 | 0.0831 |
| C | 0-10 | 2 | 11 | 25 | 13 | 2.1 | 0.1032 | 0.0949 | 5.2096 | 1.2084 | 1.0586 | -0.0612 | 0.0342 | 0.0272 |
| | 10-30 | 12 | 3 | 18 | 15 | 2.0 | 1.1476 | 0.9889 | 5.0070 | 1.4441 | 1.3930 | 0.0317 | 0.0770 | 0.0709 |
| POG | 0-10 | 12 | 1 | 14 | 13 | 2.0 | 0.0311 | 0.6958 | 4.4765 | 0.5067 | 0.4769 | -0.0893 | 0.0567 | 0.0417 |
| D | 0-10 | 13 | 5 | 23 | 18 | 2.0 | 0.5476 | 0.2467 | 3.8807 | 0.6868 | 0.3776 | -0.0755 | 0.0317 | 0.0267 |
| | 10-30 | 13 | 2 | 17 | 15 | 2.0 | 0.3832 | 0.6341 | 2.2285 | 0.5288 | 0.5202 | 0.3168 | 0.0454 | 0.0526 |
| F | 0-10 | 12 | 17 | 49 | 29 | 2.2 | 0.1923 | 0.3178 | 3.5528 | 0.3827 | 0.4099 | -0.0373 | 0.0136 | 0.0266 |
| | 10-30 | 22 | 6 | 34 | 28 | 2.0 | 0.4242 | 0.5408 | 2.2040 | 1.3533 | 1.0844 | -0.0760 | 0.0749 | 0.0674 |
| GH | 0-10 | 9 | 6 | 21 | 15 | 2.0 | 0.0492 | 0.2345 | 3.1032 | 0.3153 | 0.3176 | -0.1244 | 0.0541 | 0.0389 |
| <i>Appleton</i> | | | | | | | | | | | | | | |
| IMOR | 0-10 | 12 | 6 | 24 | 18 | 2.0 | 0.0184 | 0.3153 | 3.1269 | 0.4735 | 0.4747 | 0.0412 | 0.0255 | 0.0458 |
| N Pre-dredge | 0-10 | 26 | 6 | 42 | 32 | 2.7 | 0.1282 | 0.4005 | 4.2292 | 0.4199 | 0.3549 | -0.0281 | 0.0065 | 0.0185 |
| | 10-30 | 23 | 4 | 32 | 27 | 2.3 | 0.0186 | 0.7645 | 3.7450 | 0.6539 | 0.6366 | 0.0572 | 0.0440 | 0.0334 |
| | 30-50 | 16 | 1 | 18 | 17 | 2.0 | 0.0006 | 0.7463 | 4.4070 | 1.5119 | 1.2267 | 0.0846 | 0.0932 | 0.0504 |
| VCC | 0-10 | 27 | 14 | 57 | 41 | 2.1 | 0.3692 | 0.3242 | 3.2202 | 0.3490 | 0.2537 | -0.0582 | 0.0275 | 0.0209 |
| | 10-30 | 31 | 3 | 37 | 34 | 2.0 | 0.1965 | 0.5572 | 4.1303 | 0.6783 | 0.7806 | -0.1537 | 0.0164 | 0.0420 |
| | 30-50 | 15 | 2 | 19 | 17 | 2.0 | 0.0041 | 0.0638 | 4.4304 | 0.5727 | 0.5713 | -0.0060 | 0.0151 | 0.0135 |
| <i>Little Rapids</i> | | | | | | | | | | | | | | |
| Upper EE | 0-10 | 13 | 18 | 51 | 31 | 2.1 | 0.2516 | 0.2396 | 3.2722 | 0.7469 | 0.4948 | -0.0447 | 0.0435 | 0.0291 |
| | 10-30 | 15 | 10 | 36 | 25 | 2.1 | 0.2717 | 0.3608 | 2.5703 | 1.1521 | 0.8651 | -0.0944 | 0.0429 | 0.0460 |
| | 30-50 | 13 | 0 | 13 | 13 | 0.0 | 0.2834 | 4.7214 | 1.3448 | 1.7186 | 1.7186 | -0.0712 | 0.0536 | 0.0659 |
| Lower EE | 0-10 | 15 | 15 | 49 | 30 | 2.3 | 0.2781 | 0.5693 | 2.9308 | 0.2663 | 0.3268 | -0.0682 | 0.0193 | 0.0232 |
| | 10-30 | 23 | 10 | 45 | 33 | 2.2 | 0.4506 | 0.6548 | 2.8576 | 0.7657 | 0.9180 | -0.0759 | 0.0390 | 0.0495 |
| | 30-50 | 11 | 2 | 15 | 13 | 2.0 | 0.1221 | 0.3792 | 5.0328 | 0.9549 | 1.1745 | 0.0900 | 0.0330 | 0.0364 |
| FF | 0-10 | 18 | 14 | 50 | 32 | 2.3 | 0.3690 | 0.6980 | 3.7208 | 0.3852 | 0.4231 | -0.0549 | 0.0557 | 0.0401 |
| | 10-30 | 24 | 7 | 39 | 31 | 2.1 | 0.3304 | 0.9190 | 2.1741 | 1.3609 | 1.2502 | -0.0962 | 0.0390 | 0.0606 |
| GGHH | 0-10 | 24 | 25 | 80 | 49 | 2.2 | 0.1169 | 0.5300 | 2.8846 | 0.7084 | 0.2893 | -0.0394 | 0.0231 | 0.0235 |
| | 10-30 | 27 | 18 | 71 | 45 | 2.4 | 0.3074 | 0.9414 | 3.3231 | 0.8171 | 0.9167 | -0.0182 | 0.0596 | 0.0665 |
| | 30-50 | 73 | 2 | 78 | 75 | 2.5 | 0.0008 | 0.9359 | 0.0821 | 2.8431 | 1.3045 | 0.1762 | 0.1008 | 0.0560 |
| | 50-100 | 51 | 3 | 57 | 54 | 2.0 | 0.8083 | 0.5186 | 1.4499 | 1.9204 | 1.2885 | 0.1012 | 0.0700 | 0.0572 |
| | 100+ | 33 | 3 | 39 | 36 | 2.0 | 0.0367 | 0.1512 | 2.3137 | 0.5420 | 0.4451 | 0.0365 | 0.0249 | 0.0259 |
| <i>De Pere</i> | | | | | | | | | | | | | | |
| SMU Group 2025 | 0-10 | 32 | 11 | 57 | 43 | 2.3 | 0.0271 | 0.2709 | 3.6631 | 0.4655 | 0.4255 | -0.0528 | 0.0231 | 0.0217 |
| | 10-30 | 16 | 15 | 54 | 31 | 2.5 | 0.0886 | 0.9893 | 6.3342 | 3.4691 | 2.2114 | -0.0556 | 0.0750 | 0.0726 |
| | 30-50 | 9 | 4 | 23 | 13 | 3.5 | 0.0925 | 0.6680 | 5.5480 | 0.9776 | 1.1642 | -0.0580 | 0.0322 | 0.0335 |
| | 50-100 | 28 | 2 | 34 | 30 | 3.0 | 0.1551 | 1.1742 | 4.0031 | 1.1675 | 1.2707 | -0.0847 | 0.1058 | 0.1163 |
| 2649 | 0-10 | 54 | 12 | 80 | 66 | 2.2 | 0.0153 | 0.2503 | 3.2501 | 0.2065 | 0.4161 | -0.0608 | 0.0109 | 0.0211 |
| | 10-30 | 25 | 23 | 73 | 48 | 2.1 | 0.1028 | 1.0853 | 10.6240 | 3.2452 | 1.9813 | -0.2882 | 0.1440 | 0.0956 |
| | 50-100 | 44 | 2 | 51 | 46 | 3.5 | 0.0433 | 1.1505 | 3.6653 | 2.3249 | 1.1267 | 0.1957 | 0.1419 | 0.3961 |
| 5067 | 100+ | 31 | 14 | 63 | 45 | 2.3 | 0.5315 | 1.0783 | 1.2186 | 1.9141 | 1.2818 | 0.0177 | 0.1548 | 0.1046 |
| | 0-10 | 53 | 5 | 63 | 58 | 2.0 | 0.0736 | 0.0919 | 7.6178 | 1.2394 | 1.1333 | -0.0998 | 0.0345 | 0.0307 |
| | 10-30 | 45 | 6 | 57 | 51 | 2.0 | 0.2006 | 0.4654 | 2.4000 | 1.4903 | 1.2775 | 0.0912 | 0.0649 | 0.0465 |
| | 50-100 | 47 | 1 | 49 | 48 | 2.0 | 0.1247 | 1.0992 | 6.5635 | 2.1819 | 1.5704 | 0.3677 | 0.0684 | 0.4775 |
| 6891 | 100+ | 13 | 37 | 176 | 50 | 4.4 | 0.6534 | 4.9240 | 1.1959 | 2.3655 | 1.8648 | -0.1963 | 0.1220 | 0.1720 |
| | 0-10 | 16 | 4 | 24 | 20 | 2.0 | 0.1259 | 0.3116 | 10.2963 | 4.2471 | 5.8601 | -0.2208 | 0.0944 | 0.1858 |
| 92115 | 10-30 | 11 | 7 | 25 | 18 | 2.0 | 0.0973 | 1.0964 | 6.4202 | 1.3240 | 1.3665 | -0.1685 | 0.0765 | 0.0689 |
| | 0-10 | 21 | 6 | 33 | 27 | 2.0 | 0.0284 | 0.3161 | 0.8839 | 0.9748 | 1.1169 | 0.0413 | 0.0426 | 0.0574 |

Table A-2 Green Bay Zones 1 and 2 Outliers

| | Database ID | Reach | Fish Type | Sample Type | Total PCBs |
|--|--------------|------------------|-----------|-------------|------------|
| Fish Data: Comparison of Green Bay Zones 1 and 2 | WDF209006BC1 | Green Bay Zone 2 | alewife | whole body | 19,000 |

Reason:

Large outlier. Other PCB values range from 990 to 4,500.

Table A-3 Detailed Data for All Fish Results

| Reach | Model | Species | Sample Type | Year of Break-point | Number of Samples | Number of Samples Below Detection Limit | Standard Deviation | Chi-squared | Intercept | | | Final | | | Early | | |
|------------------------------|----------------|----------------|----------------|---------------------|-------------------|---|--------------------|-------------|-----------|-----------|---------|---------|-----------|---------|------------------|-----------|---------|
| | | | | | | | | | Intercept | Std. Err. | p-value | Slope | Std. Err. | p-value | Slope Difference | Std. Err. | p-value |
| Little Lake Butte des Morts | No Break-point | carp | skin-on fillet | | 55 | | 49.63 | | 3.3515 | 0.1131 | | -0.0456 | 0.0095 | 0.0000 | | | |
| | | | whole body | | 40 | 1 | 36.67 | | 3.6775 | 0.1089 | | -0.0750 | 0.0106 | 0.0000 | | | |
| | | northern pike | skin-on fillet | | 19 | 1 | 12.83 | | 2.6670 | 0.1303 | 0.0000 | -0.0547 | 0.0115 | 0.0003 | | | |
| | | walleye | skin-on fillet | | 63 | 8 | 42.31 | | 2.5700 | 0.0737 | | -0.0465 | 0.0066 | 0.0000 | | | |
| | | whole body | | 18 | 3 | 26.16 | | 2.6490 | 0.4089 | 0.0000 | -0.0026 | 0.0429 | 0.9532 | | | | |
| | | skin-on fillet | | 34 | 10 | 27.99 | | 2.1767 | 0.0925 | | -0.0262 | 0.0097 | 0.0112 | | | | |
| | Best Fitting | carp | skin-on fillet | 1979 | 55 | | 42.91 | 6.72 | 3.3574 | 0.1064 | | -0.0276 | 0.0112 | 0.0177 | -0.2280 | 0.0853 | 0.0102 |
| | | | whole body | 1987 | 40 | 1 | 29.39 | 7.28 | 3.3104 | 0.1645 | | 0.0031 | 0.0295 | 0.9172 | -0.1647 | 0.0588 | 0.0084 |
| | | northern pike | skin-on fillet | | 19 | 1 | 12.83 | | 2.6670 | 0.1303 | 0.0000 | -0.0547 | 0.0115 | 0.0003 | | | |
| | | walleye | skin-on fillet | 1990 | 63 | 8 | 35.98 | 6.33 | 2.2105 | 0.1605 | | 0.0147 | 0.0249 | 0.5576 | -0.0945 | 0.0373 | 0.0140 |
| | | whole body | 1987 | 18 | 3 | 16.69 | 9.48 | 2.1870 | 0.3811 | 0.0001 | 0.0845 | 0.0454 | 0.0874 | -0.2608 | 0.0802 | 0.0069 | |
| yellow perch | | skin-on fillet | 1981 | 34 | 10 | 17.83 | 10.16 | 2.3384 | 0.0908 | | 0.0031 | 0.0125 | 0.8025 | -0.2467 | 0.0771 | 0.0034 | |
| Appleton to Little Rapids | No Break-point | walleye | skin-on fillet | | 30 | | -7.15 | | 3.0085 | 0.1256 | | -0.0456 | 0.0138 | 0.0028 | | | |
| | Best Fitting | walleye | skin-on fillet | | 30 | | -7.15 | | 3.0085 | 0.1256 | | -0.0456 | 0.0138 | 0.0028 | | | |
| De Pere to Green Bay | No Break-point | carp | whole body | | 90 | | 58.07 | | 4.0144 | 0.0542 | | -0.0341 | 0.0055 | 0.0000 | | | |
| | | gizzard shad | whole body | | 19 | | -42.45 | | 3.4553 | 0.0325 | | -0.0226 | 0.0045 | 0.0002 | | | |
| | | northern pike | skin-on fillet | | 40 | 1 | -11.40 | | 3.1688 | 0.0998 | | -0.0455 | 0.0073 | 0.0000 | | | |
| | | walleye | skin-on fillet | | 120 | 1 | -41.16 | | 3.1963 | 0.0435 | | -0.0324 | 0.0036 | 0.0000 | | | |
| | | | whole body | | 58 | | -12.22 | | 3.9812 | 0.0541 | | -0.0367 | 0.0054 | 0.0000 | | | |
| | | white bass | skin-on fillet | | 58 | | -41.00 | | 3.6259 | 0.0678 | | -0.0210 | 0.0065 | 0.0020 | | | |
| | | skin-on fillet | | 44 | | -3.92 | | 3.1349 | 0.0762 | | -0.0357 | 0.0056 | 0.0000 | | | | |
| | Best Fitting | carp | whole body | 1995 | 90 | | 48.59 | 9.48 | 2.9712 | 0.3339 | 0.0000 | 0.0855 | 0.0382 | 0.0277 | -0.1406 | 0.0445 | 0.0022 |
| | | gizzard shad | whole body | | 19 | | -42.45 | | 3.4553 | 0.0325 | | -0.0226 | 0.0045 | 0.0002 | | | |
| | | northern pike | skin-on fillet | | 40 | 1 | -11.40 | | 3.1688 | 0.0998 | | -0.0455 | 0.0073 | 0.0000 | | | |
| | | walleye | skin-on fillet | | 120 | 1 | -41.16 | | 3.1963 | 0.0435 | | -0.0324 | 0.0036 | 0.0000 | | | |
| | | | whole body | | 58 | | -12.22 | | 3.9812 | 0.0541 | | -0.0367 | 0.0054 | 0.0000 | | | |
| | | white bass | skin-on fillet | | 58 | | -41.00 | | 3.6259 | 0.0678 | | -0.0210 | 0.0065 | 0.0020 | | | |
| | | | skin-on fillet | | 44 | | -3.92 | | 3.1349 | 0.0762 | | -0.0357 | 0.0056 | 0.0000 | | | |
| | | whole body | | 44 | | -3.92 | | 3.1349 | 0.0762 | | -0.0357 | 0.0056 | 0.0000 | | | | |
| Green Bay Zone 2 (2A and 2B) | No Break-point | alewife | whole body | | 44 | | -30.42 | | 3.4844 | 0.0544 | | -0.0176 | 0.0087 | 0.0497 | | | |
| | | carp | skin-on fillet | | 28 | | -4.77 | | 3.8869 | 0.0803 | | -0.0226 | 0.0154 | 0.1557 | | | |
| | | | whole body | | 57 | | -11.66 | | 3.7679 | 0.0530 | | -0.0414 | 0.0090 | 0.0000 | | | |
| | | gizzard shad | whole body | | 32 | | -51.90 | | 3.2444 | 0.0535 | | 0.0249 | 0.0095 | 0.0144 | | | |
| | | skin-on fillet | | 19 | 3 | -8.96 | | 2.6539 | 0.4357 | 0.0000 | -0.0494 | 0.0143 | 0.0038 | | | | |
| | Best Fitting | alewife | whole body | | 44 | | -30.42 | | 3.4844 | 0.0544 | | -0.0176 | 0.0087 | 0.0497 | | | |
| | | carp | skin-on fillet | | 28 | | -4.77 | | 3.8869 | 0.0803 | | -0.0226 | 0.0154 | 0.1557 | | | |
| | | | whole body | 1983 | 57 | | -29.32 | 17.66 | 3.8825 | 0.0519 | | -0.0733 | 0.0104 | 0.0000 | 0.2664 | 0.0585 | 0.0000 |
| | | gizzard shad | whole body | | 32 | | -51.90 | | 3.2444 | 0.0535 | | 0.0249 | 0.0095 | 0.0144 | | | |
| | | | skin-on fillet | | 19 | 3 | -8.96 | | 2.6539 | 0.4357 | 0.0000 | -0.0494 | 0.0143 | 0.0038 | | | |
| yellow perch | | skin-on fillet | | 19 | 3 | -8.96 | | 2.6539 | 0.4357 | 0.0000 | -0.0494 | 0.0143 | 0.0038 | | | | |

Note:

In the fitted models, amplitude and month of peak can be ignored if \log_{10} PCB concentration estimates are needed for July 1 of any year. For other times of year, let M be the \log_{10} of the estimated concentration on July 1, A = amplitude, t_{max} = ("month of peak" - 1)/12, and t = the specified time of year as a value between zero (1 January) and 1.0 (31 December). Define $Q(t) = -A \cdot \cos[2\pi(0.5 - t_{max})] + A \cdot \cos[2\pi(t - t_{max})]$. Then the estimated mean concentration (ppb) at time-of-year t is $M \cdot 10^{Q(t)}$.

Table A-3 Detailed Data for All Fish Results

| Reach | Model | Species | Sample Type | Year of Break-point | Fat | | | Month Peak | Amplitude | | Covariate Intercept Time | Mean Squared Error | Percent Change per Year | T-squared | | |
|------------------------------|----------------|----------------|----------------|---------------------|-------------------|-----------|---------|------------|-----------|---------|--------------------------|--------------------|-------------------------|-----------|-----------|---------|
| | | | | | Log ₁₀ | Std. Err. | p-value | | Amplitude | p-value | | | | T-squared | Std. Err. | p-value |
| Little Lake Butte des Morts | No Break-point | carp | skin-on fillet | | 0.8927 | 0.1611 | 0.0000 | 1.328 | 0.5316 | 0.2260 | 0.0006 | 0.1444 | -9.9650 | 0.00231 | 0.00190 | 0.2292 |
| | | | whole body | | 0.8753 | 0.3590 | 0.0200 | 6.356 | 0.6174 | 0.0965 | -0.0004 | 0.1374 | -15.8538 | 0.00360 | 0.00229 | 0.1249 |
| | | northern pike | skin-on fillet | | 0.4469 | 0.2976 | 0.1554 | 1.311 | 0.6671 | 0.1594 | 0.0005 | 0.1034 | -11.8315 | -0.00334 | 0.00242 | 0.1904 |
| | | walleye | skin-on fillet | | 0.3898 | 0.1444 | 0.0091 | 1.558 | 0.1861 | 0.6458 | 0.0001 | 0.0934 | -10.1572 | 0.00285 | 0.00123 | 0.0241 |
| | Best Fitting | carp | skin-on fillet | 1979 | 0.8675 | 0.1519 | 0.0000 | 12.904 | 0.3939 | 0.0078 | 0.0006 | 0.1277 | -6.1477 | -0.00137 | 0.00236 | 0.5645 |
| | | | whole body | 1987 | 0.8626 | 0.3293 | 0.0131 | 7.013 | 0.8307 | 0.0025 | -0.0039 | 0.1156 | 0.7139 | -0.01442 | 0.00670 | 0.0388 |
| | | northern pike | skin-on fillet | | 0.4469 | 0.2976 | 0.1554 | 1.311 | 0.6671 | 0.1594 | 0.0005 | 0.1034 | -11.8315 | -0.00334 | 0.00242 | 0.1904 |
| | | walleye | skin-on fillet | 1990 | 0.5012 | 0.1455 | 0.0011 | 11.638 | 0.2005 | 0.0273 | -0.0034 | 0.0857 | 3.4395 | -0.00949 | 0.00939 | 0.3167 |
| | | whole body | 1987 | 0.9858 | 0.3619 | 0.0185 | 11.562 | 0.4627 | 0.0040 | -0.0157 | 0.1410 | 21.4715 | -0.02024 | 0.01008 | 0.0698 | |
| | | yellow perch | skin-on fillet | 1981 | 0.4946 | 0.2067 | 0.0236 | 7.033 | 0.2185 | 0.0007 | 0.0005 | 0.0719 | 0.7276 | -0.00211 | 0.00587 | 0.7217 |
| Appleton to Little Rapids | No Break-point | walleye | skin-on fillet | | 1.0801 | 0.1555 | 0.0000 | 8.121 | 0.4280 | 0.0010 | 0.0015 | 0.0461 | -9.9680 | -0.00472 | 0.00405 | 0.2554 |
| | Best Fitting | walleye | skin-on fillet | | 1.0801 | 0.1555 | 0.0000 | 8.121 | 0.4280 | 0.0010 | 0.0015 | 0.0461 | -9.9680 | -0.00472 | 0.00405 | 0.2554 |
| De Pere to Green Bay | No Break-point | carp | whole body | | 0.8225 | 0.1180 | 0.0000 | 6.889 | 0.1825 | 0.0471 | -0.0001 | 0.1116 | -7.5413 | 0.00214 | 0.00103 | 0.0411 |
| | | | gizzard shad | whole body | | 0.5055 | 0.0897 | 0.0001 | 8.558 | 0.5814 | 0.0000 | -0.0001 | 0.0063 | -5.0657 | 0.00318 | 0.00289 |
| | | northern pike | skin-on fillet | | 0.7224 | 0.1664 | 0.0001 | 10.122 | 0.1730 | 0.3531 | -0.0004 | 0.0407 | -9.9517 | 0.00093 | 0.00079 | 0.2489 |
| | | walleye | skin-on fillet | | 0.8509 | 0.0673 | | 9.454 | 0.0172 | 0.7566 | -0.0001 | 0.0406 | -7.1920 | -0.00051 | 0.00062 | 0.4177 |
| | | | whole body | | 0.4449 | 0.1231 | 0.0007 | 6.973 | 0.1190 | 0.2038 | -0.0001 | 0.0474 | -8.1055 | -0.00003 | 0.00082 | 0.9712 |
| | | white bass | skin-on fillet | | 0.8170 | 0.1134 | 0.0000 | 6.750 | 0.3258 | 0.1043 | 0.0001 | 0.0289 | -4.7229 | 0.00152 | 0.00183 | 0.4104 |
| | white sucker | skin-on fillet | | 0.4255 | 0.1496 | 0.0071 | 6.923 | 0.0827 | 0.5528 | 0.0000 | 0.0536 | -7.8956 | 0.00110 | 0.00104 | 0.2996 | |
| | Best Fitting | carp | whole body | 1995 | 0.7871 | 0.1125 | 0.0000 | 6.657 | 0.0642 | 0.0004 | -0.0126 | 0.1005 | 21.7626 | 0.01676 | 0.03616 | 0.6442 |
| | | | gizzard shad | whole body | | 0.5055 | 0.0897 | 0.0001 | 8.558 | 0.5814 | 0.0000 | -0.0001 | 0.0063 | -5.0657 | 0.00318 | 0.00289 |
| | | northern pike | skin-on fillet | | 0.7224 | 0.1664 | 0.0001 | 10.122 | 0.1730 | 0.3531 | -0.0004 | 0.0407 | -9.9517 | 0.00093 | 0.00079 | 0.2489 |
| | | walleye | skin-on fillet | | 0.8509 | 0.0673 | | 9.454 | 0.0172 | 0.7566 | -0.0001 | 0.0406 | -7.1920 | -0.00051 | 0.00062 | 0.4177 |
| | | | whole body | | 0.4449 | 0.1231 | 0.0007 | 6.973 | 0.1190 | 0.2038 | -0.0001 | 0.0474 | -8.1055 | -0.00003 | 0.00082 | 0.9712 |
| | | white bass | skin-on fillet | | 0.8170 | 0.1134 | 0.0000 | 6.750 | 0.3258 | 0.1043 | 0.0001 | 0.0289 | -4.7229 | 0.00152 | 0.00183 | 0.4104 |
| | | white sucker | skin-on fillet | | 0.4255 | 0.1496 | 0.0071 | 6.923 | 0.0827 | 0.5528 | 0.0000 | 0.0536 | -7.8956 | 0.00110 | 0.00104 | 0.2996 |
| Green Bay Zone 2 (2A and 2B) | | No Break-point | alewife | whole body | | 0.9126 | 0.1409 | 0.0000 | 6.054 | 0.1664 | 0.0335 | -0.0001 | 0.0293 | -3.9623 | 0.00191 | 0.00113 |
| | carp | | | skin-on fillet | | 0.7643 | 0.1515 | 0.0000 | 3.941 | 0.2377 | 0.0288 | -0.0001 | 0.0494 | -5.0631 | -0.00608 | 0.00349 |
| | gizzard shad | | whole body | | 0.9578 | 0.1099 | 0.0000 | 6.794 | 0.1308 | 0.2408 | 0.0000 | 0.0477 | -9.1004 | -0.00275 | 0.00118 | 0.0238 |
| | | | yellow perch | skin-on fillet | | -0.1295 | 0.1177 | 0.2811 | 2.645 | 0.3356 | 0.0300 | -0.0002 | 0.0116 | 5.9098 | -0.00074 | 0.00319 |
| | Best Fitting | alewife | whole body | | 0.9126 | 0.1409 | 0.0000 | 6.054 | 0.1664 | 0.0335 | -0.0001 | 0.0293 | -3.9623 | 0.00191 | 0.00113 | 0.0992 |
| | | | carp | skin-on fillet | 1983 | 0.7643 | 0.1515 | 0.0000 | 3.941 | 0.2377 | 0.0288 | -0.0001 | 0.0494 | -5.0631 | -0.00608 | 0.00349 |
| | | gizzard shad | whole body | | 0.8981 | 0.0950 | 0.0000 | 6.864 | 0.2382 | 0.0000 | -0.0002 | 0.0350 | -15.5359 | 0.00335 | 0.00175 | 0.0616 |
| | | | yellow perch | skin-on fillet | | -0.1295 | 0.1177 | 0.2811 | 2.645 | 0.3356 | 0.0300 | -0.0002 | 0.0116 | 5.9098 | -0.00074 | 0.00319 |
| | | yellow perch | skin-on fillet | | 1.0912 | 0.4683 | 0.0353 | 4.726 | 0.4459 | 0.5489 | -0.0020 | 0.0316 | -10.7477 | 0.01258 | 0.00339 | 0.0026 |
| | | | whole body | | 0.9126 | 0.1409 | 0.0000 | 6.054 | 0.1664 | 0.0335 | -0.0001 | 0.0293 | -3.9623 | 0.00191 | 0.00113 | 0.0992 |

Note:

In the fitted models, amplitude and month of peak can be concentration on July 1, A = amplitude, t_{max} = ("month of peak" - 1)/12, and t = the specified time of year (1 January) and 1.0 (31 December). Define $Q(t) = -A \cdot \cos[2\pi(0.5 - t_{max})] + A \cdot \cos[2\pi(t - t_{max})]$.

ignored if \log_{10} PCB concentration estimates are needed for July 1 of any year. For other times of year, let M be the \log_{10} of the estimate. Then the estimated mean concentration (ppb) at time-of-year t is $M \cdot 10^{Q(t)}$.

Table A-4 Testing the Null Hypothesis that a Straight Line Fits As Well As a Spline Model with a Breakpoint

| Reach | Species | Sample Type | Year of Best-fitting Breakpoint | Sample Size | p-value for Breakpoint | p < 0.05 | Final (post-break) Slope | p-value for Final Slope | Pre-break Slope Minus Final Slope | p-value for Slope Difference | Pre-break Slope |
|------------------------------|-------------------|-------------------|---------------------------------|-------------|------------------------|---------------|--------------------------|-------------------------|-----------------------------------|------------------------------|-----------------|
| Little Lake Butte des Morts | Carp | skin-on fillet | 1979 | 55 | 0.0347 | * | -0.028 | 0.0177 | -0.228 | 0.0102 | -0.256 |
| | | whole body | 1987 | 40 | 0.0263 | * | 0.003 | 0.9172 | -0.165 | 0.0084 | -0.162 |
| | Northern Pike | skin-on fillet | 1996 | 19 | 0.2723 | | -0.325 | 0.0685 | 0.301 | 0.1214 | -0.024 |
| | Walleye | skin-on fillet | 1990 | 63 | 0.0423 | * | 0.015 | 0.5576 | -0.095 | 0.0140 | -0.080 |
| | | whole body | 1987 | 18 | 0.0088 | * | 0.084 | 0.0874 | -0.261 | 0.0069 | -0.176 |
| | Yellow Perch | skin-on fillet | 1981 | 34 | 0.0062 | * | 0.003 | 0.8025 | -0.247 | 0.0034 | -0.244 |
| | Combined++ | | | 229 | 0.0000 | * | | | | | |
| Appleton to Little Rapids | Walleye | skin-on fillet | 1983 | 30 | 0.4526 | | -0.056 | 0.0015 | 0.103 | 0.2142 | 0.047 |
| De Pere to Green Bay | Carp | whole body | 1995 | 90 | 0.0087 | * | 0.086 | 0.0277 | -0.141 | 0.0022 | -0.055 |
| | Gizzard Shad | whole body | 1990 | 19 | 0.4672 | | -0.020 | 0.0018 | -0.042 | 0.2303 | -0.062 |
| | Northern Pike | skin-on fillet | 1996 | 40 | 0.1421 | | 0.060 | 0.2616 | -0.117 | 0.0514 | -0.056 |
| | Walleye | skin-on fillet | 1993 | 120 | 0.5680 | | -0.046 | 0.0006 | 0.019 | 0.2885 | -0.027 |
| | | whole body | 1996 | 58 | 0.5550 | | 0.010 | 0.8196 | -0.052 | 0.2805 | -0.042 |
| | White Bass | skin-on fillet | 1996 | 58 | 0.6059 | | 0.019 | 0.6373 | -0.045 | 0.3193 | -0.025 |
| | White Sucker | skin-on fillet | 1990 | 44 | 0.1986 | | -0.006 | 0.7235 | -0.049 | 0.0749 | -0.055 |
| | Combined++ | | | 429 | 0.0906 | | | | | | |
| Green Bay Zone 2 (2A and 2B) | Alewife | whole body | 1986 | 44 | 0.0863 | | -0.001 | 0.9394 | -0.076 | 0.0285 | -0.077 |
| | Carp | skin-on fillet | 1985 | 28 | 0.1811 | | -0.063 | 0.0226 | 0.105 | 0.0698 | 0.042 |
| | | whole body | 1983 | 57 | 0.0001 | * | -0.073 | 0.0000 | 0.266 | 0.0000 | 0.193 |
| | Gizzard Shad | whole body | 1996 | 32 | 0.6655 | | -0.014 | 0.7556 | 0.047 | 0.3721 | 0.033 |
| | Yellow Perch | skin-on fillet | 1986 | 19 | 0.0008 | * | 0.062 | 0.0325 | -0.573 | 0.0004 | -0.511 |
| | | Combined++ | | | 180 | 0.0000 | * | | | | |

Note:

++ Indicates p-value for the test that all fish categories in a reach do not have a breakpoint.

Table A-5 Breakpoint, Final Slope, and Percent Change per Year of PCB Concentration from Best-fitting Model

| Reach | Species | Sample Type | Year of Breakpoint | Number of Samples | Final (post-break) Slope | p-value for Final Slope (versus zero) | Percent per Year |
|------------------------------|----------------|----------------|--------------------|-------------------|--------------------------|---------------------------------------|------------------|
| Little Lake Butte des Morts | Carp | skin-on fillet | 1979 | 55 | -0.028 | 0.0177 | -6.1 |
| | | whole body | 1987 | 40 | 0.003 | 0.9172 | 0.7 |
| | Northern Pike | skin-on fillet | 0 | 19 | -0.055 | 0.0003 | -11.8 |
| | Walleye | skin-on fillet | 1990 | 63 | 0.015 | 0.5576 | 3.4 |
| | | whole body | 1987 | 18 | 0.084 | 0.0874 | 21.5 |
| Yellow Perch | skin-on fillet | 1981 | 34 | 0.003 | 0.8025 | 0.7 | |
| Appleton to Little Rapids | Walleye | skin-on fillet | 0 | 30 | -0.046 | 0.0028 | -10.0 |
| De Pere to Green Bay | Carp | whole body | 1995 | 90 | 0.086 | 0.0277 | 21.8 |
| | Gizzard Shad | whole body | 0 | 19 | -0.023 | 0.0002 | -5.1 |
| | Northern Pike | skin-on fillet | 0 | 40 | -0.046 | 0.0000 | -10.0 |
| | Walleye | skin-on fillet | 0 | 120 | -0.032 | 0.0000 | -7.2 |
| | | whole body | 0 | 58 | -0.037 | 0.0000 | -8.1 |
| | White Bass | skin-on fillet | 0 | 58 | -0.021 | 0.0020 | -4.7 |
| White Sucker | skin-on fillet | 0 | 44 | -0.036 | 0.0000 | -7.9 | |
| Green Bay Zone 2 (2A and 2B) | Alewife | whole body | 0 | 44 | -0.018 | 0.0497 | -4.0 |
| | Carp | skin-on fillet | 0 | 28 | -0.023 | 0.1557 | -5.1 |
| | | whole body | 1983 | 57 | -0.073 | 0.0000 | -15.5 |
| | Gizzard Shad | whole body | 0 | 32 | 0.025 | 0.0144 | 5.9 |
| Yellow Perch | skin-on fillet | 0 | 19 | -0.049 | 0.0038 | -10.7 | |

Table A-6 Model Parameters and Other Statistics for the Best-fitting Model

| Reach | Species | Sample Type | Year of Breakpoint | Number of Samples | | | Final (post-break) Slope | |
|------------------------------------|----------------|----------------|--------------------|-------------------|-----------|----------------|--------------------------|----------------|
| | | | | <i>n</i> | Intercept | Standard Error | Final | Standard Error |
| Little Lake Butte des Morts | Carp | skin-on fillet | 1979 | 55 | 3.36 | 0.11 | -0.028 | 0.011 |
| | | whole body | 1987 | 40 | 3.31 | 0.16 | 0.003 | 0.030 |
| | Northern Pike | skin-on fillet | 0 | 19 | 2.67 | 0.13 | -0.055 | 0.011 |
| | Walleye | skin-on fillet | 1990 | 63 | 2.21 | 0.16 | 0.015 | 0.025 |
| | | whole body | 1987 | 18 | 2.19 | 0.38 | 0.084 | 0.045 |
| Yellow Perch | skin-on fillet | 1981 | 34 | 2.34 | 0.09 | 0.003 | 0.012 | |
| Appleton to Little Rapids | Walleye | skin-on fillet | 0 | 30 | 3.01 | 0.13 | -0.046 | 0.014 |
| De Pere to Green Bay | Carp | whole body | 1995 | 90 | 2.97 | 0.33 | 0.086 | 0.038 |
| | Gizzard Shad | whole body | 0 | 19 | 3.46 | 0.03 | -0.023 | 0.005 |
| | Northern Pike | skin-on fillet | 0 | 40 | 3.17 | 0.10 | -0.046 | 0.007 |
| | Walleye | skin-on fillet | 0 | 120 | 3.20 | 0.04 | -0.032 | 0.004 |
| | | whole body | 0 | 58 | 3.98 | 0.05 | -0.037 | 0.005 |
| | White Bass | skin-on fillet | 0 | 58 | 3.63 | 0.07 | -0.021 | 0.006 |
| White Sucker | skin-on fillet | 0 | 44 | 3.13 | 0.08 | -0.036 | 0.006 | |
| Green Bay Zone 2 (2A and 2B) | Alewife | whole body | 0 | 44 | 3.48 | 0.05 | -0.018 | 0.009 |
| | Carp | skin-on fillet | 0 | 28 | 3.89 | 0.08 | -0.023 | 0.015 |
| | | whole body | 1983 | 57 | 3.88 | 0.05 | -0.073 | 0.010 |
| | Gizzard Shad | whole body | 0 | 32 | 3.24 | 0.05 | 0.025 | 0.010 |
| Yellow Perch | skin-on fillet | 0 | 19 | 2.65 | 0.44 | -0.049 | 0.014 | |

Notes:

MSE - Mean square error.

* An estimate of the residual variance.

** An estimate of residual standard deviation.

Table A-6 Model Parameters and Other Statistics for the Best-fitting Model

| Reach | Species | Sample Type | Year of Breakpoint | p-value for Final Slope | | | | p-value for Early Slope Difference | | |
|------------------------------|----------------|----------------|--------------------|-------------------------|------------------|-----------------------------------|----------------|------------------------------------|-----------------------------|----------------|
| | | | | p-value | Percent per Year | Pre-break Slope Minus Final Slope | Standard Error | p-value | Coefficient of Log(% lipid) | Standard Error |
| Little Lake Butte des Morts | Carp | skin-on fillet | 1979 | 0.0177 | -6.1 | -0.228 | 0.085 | 0.0102 | 0.87 | 0.15 |
| | | whole body | 1987 | 0.9172 | 0.7 | -0.165 | 0.059 | 0.0084 | 0.86 | 0.33 |
| | Northern Pike | skin-on fillet | 0 | 0.0003 | -11.8 | | | | 0.45 | 0.30 |
| | Walleye | skin-on fillet | 1990 | 0.5576 | 3.4 | -0.095 | 0.037 | 0.0140 | 0.50 | 0.15 |
| | | whole body | 1987 | 0.0874 | 21.5 | -0.261 | 0.080 | 0.0069 | 0.99 | 0.36 |
| Yellow Perch | skin-on fillet | 1981 | 0.8025 | 0.7 | -0.247 | 0.077 | 0.0034 | 0.49 | 0.21 | |
| Appleton to Little Rapids | Walleye | skin-on fillet | 0 | 0.0028 | -10.0 | | | | 1.08 | 0.16 |
| De Pere to Green Bay | Carp | whole body | 1995 | 0.0277 | 21.8 | -0.141 | 0.044 | 0.0022 | 0.79 | 0.11 |
| | Gizzard Shad | whole body | 0 | 0.0002 | -5.1 | | | | 0.51 | 0.09 |
| | Northern Pike | skin-on fillet | 0 | 0.0000 | -10.0 | | | | 0.72 | 0.17 |
| | Walleye | skin-on fillet | 0 | 0.0000 | -7.2 | | | | 0.85 | 0.07 |
| | | whole body | 0 | 0.0000 | -8.1 | | | | 0.44 | 0.12 |
| | White Bass | skin-on fillet | 0 | 0.0020 | -4.7 | | | | 0.82 | 0.11 |
| White Sucker | skin-on fillet | 0 | 0.0000 | -7.9 | | | | 0.43 | 0.15 | |
| Green Bay Zone 2 (2A and 2B) | Alewife | whole body | 0 | 0.0497 | -4.0 | | | | 0.91 | 0.14 |
| | Carp | skin-on fillet | 0 | 0.1557 | -5.1 | | | | 0.76 | 0.15 |
| | | whole body | 1983 | 0.0000 | -15.5 | 0.266 | 0.059 | 0.0000 | 0.90 | 0.10 |
| | Gizzard Shad | whole body | 0 | 0.0144 | 5.9 | | | | -0.13 | 0.12 |
| | Yellow Perch | skin-on fillet | 0 | 0.0038 | -10.7 | | | | 1.09 | 0.47 |

Notes:

MSE - Mean square error.

* An estimate of the residual variance.

** An estimate of residual standard deviation.

Table A-6 Model Parameters and Other Statistics for the Best-fitting Model

| Reach | Species | Sample Type | Year of Breakpoint | p-value for Log(% lipid) | Month of Seasonal Peak | Amplitude of Seasonal Peak | p-value for Seasonal Effect | Mean Square Error* | Square Root of MSE** |
|------------------------------|----------------|----------------|--------------------|--------------------------|------------------------|----------------------------|-----------------------------|--------------------|----------------------|
| Little Lake Butte des Morts | Carp | skin-on fillet | 1979 | 0.0000 | 12.9 | 0.39 | 0.0078 | 0.128 | 0.357 |
| | | whole body | 1987 | 0.0131 | 7.0 | 0.83 | 0.0025 | 0.116 | 0.340 |
| | Northern Pike | skin-on fillet | 0 | 0.1554 | 1.3 | 0.67 | 0.1594 | 0.103 | 0.322 |
| | Walleye | skin-on fillet | 1990 | 0.0011 | 11.6 | 0.20 | 0.0273 | 0.086 | 0.293 |
| | | whole body | 1987 | 0.0185 | 11.6 | 0.46 | 0.0040 | 0.141 | 0.376 |
| Yellow Perch | skin-on fillet | 1981 | 0.0236 | 7.0 | 0.22 | 0.0007 | 0.072 | 0.268 | |
| Appleton to Little Rapids | Walleye | skin-on fillet | 0 | 0.0000 | 8.1 | 0.43 | 0.0010 | 0.046 | 0.215 |
| De Pere to Green Bay | Carp | whole body | 1995 | 0.0000 | 6.7 | 0.06 | 0.0004 | 0.100 | 0.317 |
| | Gizzard Shad | whole body | 0 | 0.0001 | 8.6 | 0.58 | 0.0000 | 0.006 | 0.079 |
| | Northern Pike | skin-on fillet | 0 | 0.0001 | 10.1 | 0.17 | 0.3531 | 0.041 | 0.202 |
| | Walleye | skin-on fillet | 0 | 0.0000 | 9.5 | 0.02 | 0.7566 | 0.041 | 0.201 |
| | | whole body | 0 | 0.0007 | 7.0 | 0.12 | 0.2038 | 0.047 | 0.218 |
| | White Bass | skin-on fillet | 0 | 0.0000 | 6.7 | 0.33 | 0.1043 | 0.029 | 0.170 |
| White Sucker | skin-on fillet | 0 | 0.0071 | 6.9 | 0.08 | 0.5528 | 0.054 | 0.231 | |
| Green Bay Zone 2 (2A and 2B) | Alewife | whole body | 0 | 0.0000 | 6.1 | 0.17 | 0.0335 | 0.029 | 0.171 |
| | Carp | skin-on fillet | 0 | 0.0000 | 3.9 | 0.24 | 0.0288 | 0.049 | 0.222 |
| | | whole body | 1983 | 0.0000 | 6.9 | 0.24 | 0.0000 | 0.035 | 0.187 |
| | Gizzard Shad | whole body | 0 | 0.2811 | 2.6 | 0.34 | 0.0300 | 0.012 | 0.108 |
| | Yellow Perch | skin-on fillet | 0 | 0.0353 | 4.7 | 0.45 | 0.5489 | 0.032 | 0.178 |

Notes:

MSE - Mean square error.

* An estimate of the residual variance.

** An estimate of residual standard deviation.

Table A-7 Final Slope and Percent Change per Year for Best-fitting Model, and Sensitivity Analysis

| Reach | Species | Sample Type | Sample Size | Year of Breakpoint—Best Model | | | Year of Breakpoint—Earliest | | | Year of Breakpoint—Latest | | | Year of Breakpoint—1985 | | |
|------------------------------|-----------------|-----------------|-------------|-------------------------------|-------------------------|---------------------|-----------------------------|-------------------------|---------------------|---------------------------|-------------------------|---------------------|-------------------------|-------------------------|----------|
| | | | | Year | Percent Change per Year | p-value (for % = 0) | Year | Percent Change per Year | p-value (for % = 0) | Year | Percent Change per Year | p-value (for % = 0) | Year | Percent Change per Year | p-value* |
| Little Lake Butte des Morts | Carp | skin-on fillet | 55 | 1979 | -6.15 | 0.0177 | 1979 | -6.15 | 0.0177 | 1985 | -1.56 | 0.7419 | 1985 | -1.56 | 0.7419 |
| | | whole body | 40 | 1987 | 0.71 | 0.9172 | 1985 | -4.04 | 0.5264 | 1990 | -0.25 | 0.9765 | 1985 | -4.04 | 0.5264 |
| | Northern Pike | skin-on fillet | 19 | 0 | -11.83 | 0.0003 | | | | | | | | | |
| | Walleye | skin-on fillet | 63 | 1990 | 3.44 | 0.5576 | 1979 | -8.37 | 0.0000 | 1994 | 8.82 | 0.4482 | 1985 | -5.83 | 0.0379 |
| | | whole body | 18 | 1987 | 21.47 | 0.0874 | 1984 | 15.10 | 0.2024 | 1990 | 21.11 | 0.1324 | 1985 | 18.49 | 0.1285 |
| | Yellow Perch | skin-on fillet | 34 | 1981 | 0.73 | 0.8025 | 1979 | 0.27 | 0.9252 | 1996 | 333.61 | 0.0122 | 1985 | 4.33 | 0.3297 |
| | Combined | | | | -4.86 | 0.0055 | | | | | | | | | |
| Appleton to Little Rapids | Walleye | skin-on fillet | 30 | 0 | -9.97 | 0.0028 | | | | | | | | | |
| De Pere to Green Bay | Carp | whole body | 90 | 1995 | 21.76 | 0.0277 | 1990 | -0.69 | 0.8232 | 1996 | 29.80 | 0.0191 | 1985 | -5.63 | 0.0238 |
| | Gizzard Shad | whole body | 19 | 0 | -5.07 | 0.0002 | | | | | | | | | |
| | Northern Pike | skin-on fillet | 40 | 0 | -9.95 | 0.0000 | | | | | | | | | |
| | | whole body | 58 | 0 | -8.11 | 0.0000 | | | | | | | | | |
| | Walleye | skin-on fillet | 120 | 0 | -7.19 | 0.0000 | | | | | | | | | |
| | | whole body | 58 | 0 | -8.11 | 0.0000 | | | | | | | | | |
| | White Bass | skin-on fillet | 58 | 0 | -4.72 | 0.0020 | | | | | | | | | |
| White Sucker | skin-on fillet | 44 | 0 | -7.90 | 0.0000 | | | | | | | | | | |
| | Combined | | | | -6.89 | 0.0000 | | | | | | | -6.92 | 0.0000 | |
| Green Bay Zone 2 (2A and 2B) | Alewife | whole body | 44 | 0 | -3.96 | 0.0497 | | | | | | | | | |
| | Carp | skin-on fillet | 28 | 0 | -5.06 | 0.1557 | | | | | | | | | |
| | | whole body | 57 | 1983 | -15.54 | 0.0000 | 1983 | -15.54 | 0.0000 | 1984 | -16.15 | 0.0000 | 1985 | -15.90 | 0.0000 |
| | Gizzard Shad | whole body | 32 | 0 | 5.91 | 0.0144 | | | | | | | | | |
| | Yellow Perch | skin-on fillet | 19 | 0 | -10.75 | 0.0038 | | | | | | | | | |
| | | Combined | | | | -5.11 | 0.0000 | | | | | | | -5.99 | 0 |

Note:

* For testing whether percent change per year is different from zero.

Table A-8 Computing Whole Body PCB Concentrations*

| Species | Convert | Modify PCB Target by this Factor |
|----------------|----------------|---|
| Carp | 0.59 | 1.69 |
| Northern Pike | 0.1 | 10.00 |
| Walleye | 0.1 | 10.00 |
| White Bass | 0.43 | 2.33 |
| White Sucker | 0.59 | 1.69 |
| Yellow Perch | 0.04 | 25.00 |

Note:

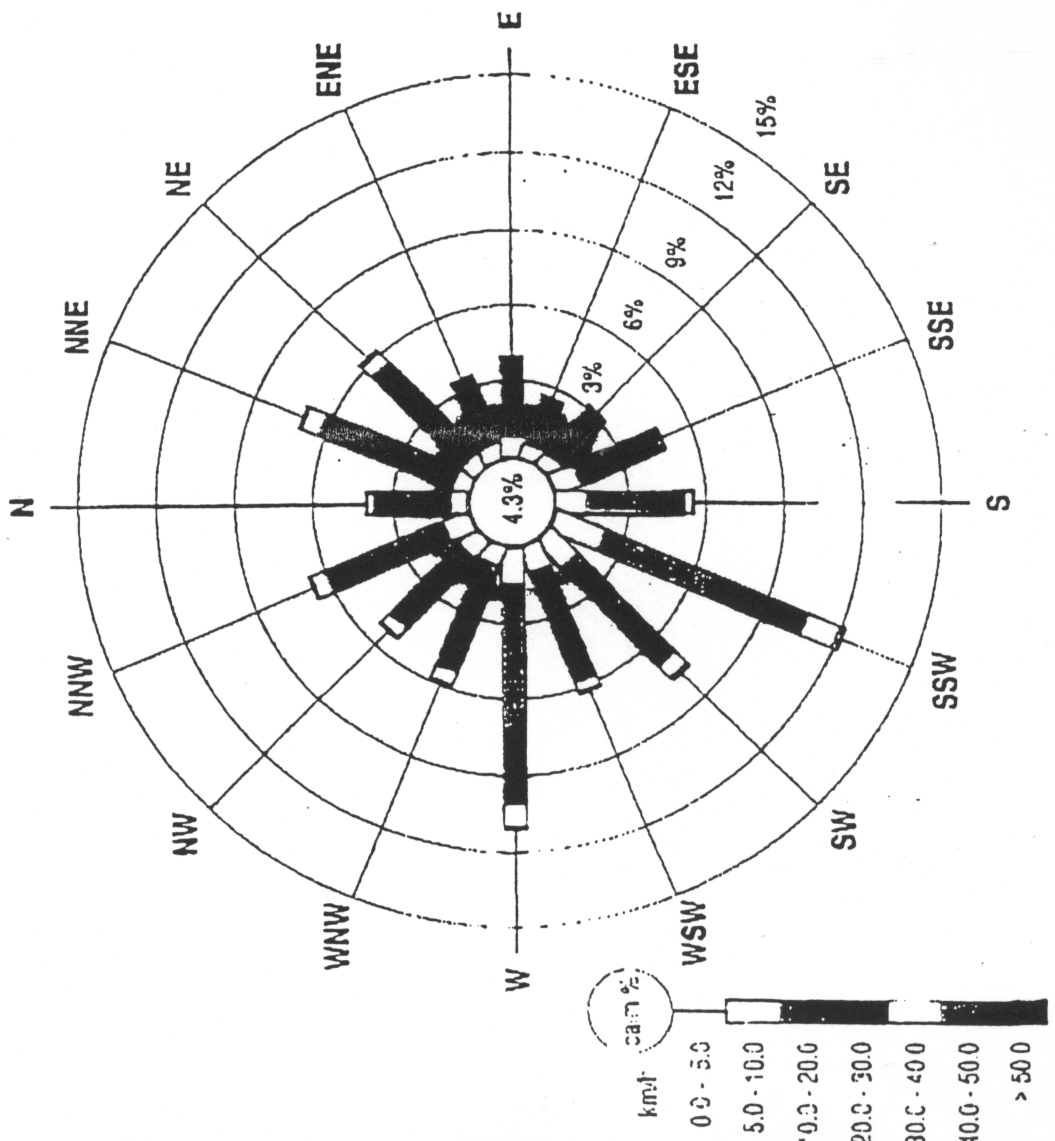
* Based on fillet-to-whole body conversion factors. These conversion factors were used to multiply specified skin-on fillet PCB concentrations to yield the corresponding expected concentration in a whole-body sample—used in analyses of time to reach specified PCB concentrations.

APPENDIX C

**A WINDROSE DIAGRAM, DEVELOPED FROM THE NATIONAL
OCEANIC AND ATMOSPHERIC ADMINISTRATION**

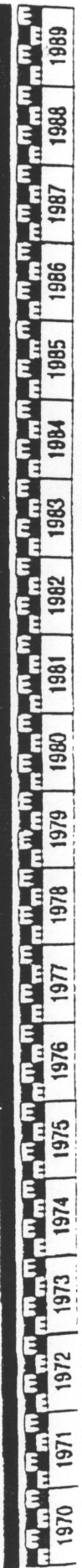
Wind Station: Green Bay

Wind Rose



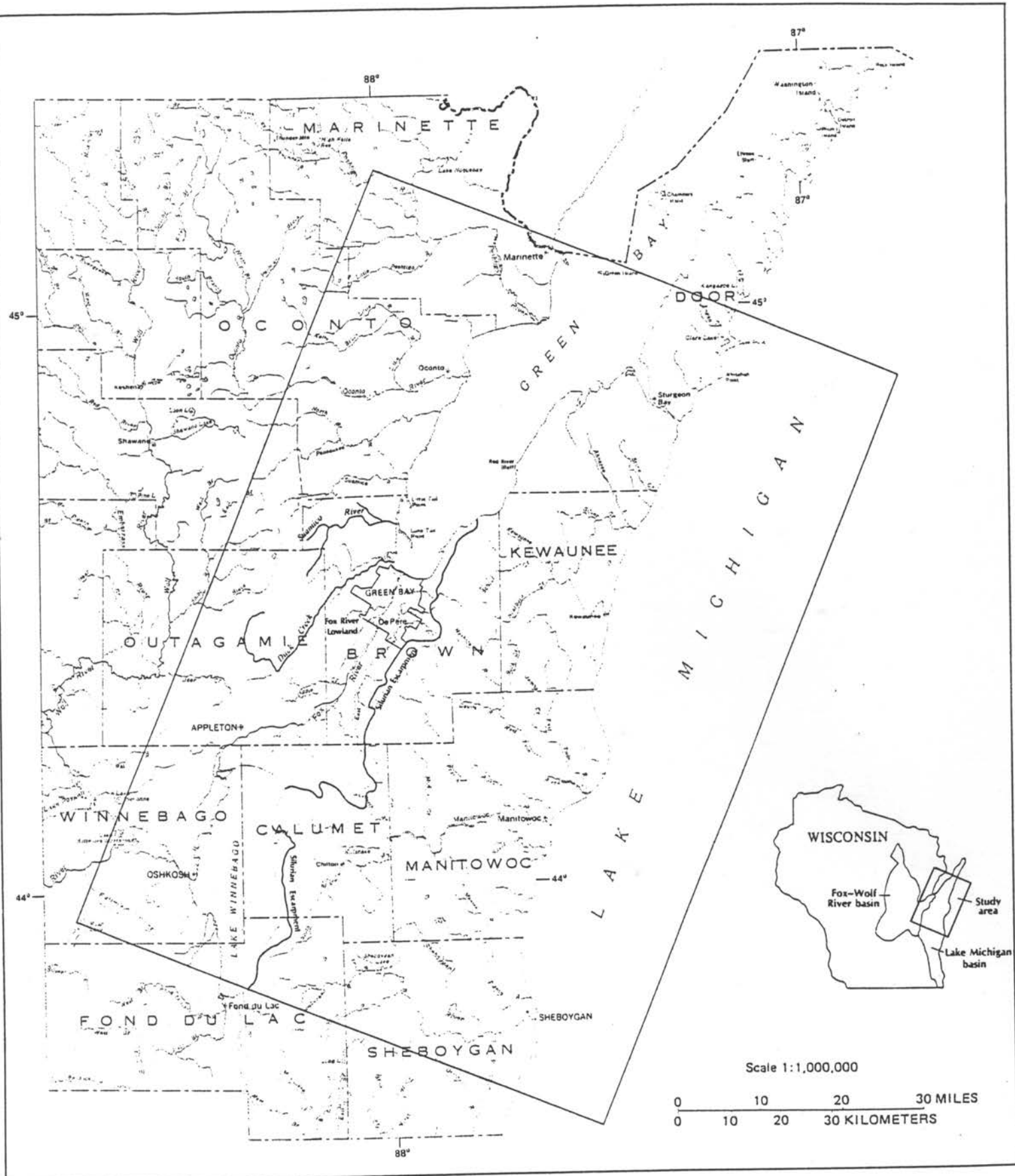
Cumulative Distribution (%)

| | 0.0 | 5.0 | 10.0 | 20.0 | 30.0 | 40.0 | 50.0 | km/h |
|-----|-----|-----|------|------|------|------|------|------|
| NNE | 0.0 | 0.8 | 4.0 | 6.3 | 6.9 | 7.0 | | |
| NE | 0.0 | 0.7 | 3.7 | 5.6 | 6.1 | 6.3 | | 6.3 |
| ENE | 0.0 | 0.7 | 2.6 | 3.3 | 3.4 | 3.4 | | |
| E | 0.1 | 0.9 | 3.1 | 3.8 | 3.9 | | | |
| ESE | 0.0 | 0.6 | 2.0 | 2.5 | 2.6 | | | |
| SE | 0.0 | 0.7 | 2.4 | 3.0 | 3.1 | | | |
| SSE | 0.1 | 1.1 | 3.6 | 4.6 | 4.7 | 4.8 | | |
| S | 0.1 | 1.3 | 4.1 | 5.2 | 5.5 | 5.5 | | |
| SSW | 0.1 | 2.2 | 7.4 | 10.6 | 11.9 | 12.1 | | 12.1 |
| SW | 0.1 | 1.5 | 5.0 | 6.9 | 7.5 | 7.7 | | 7.7 |
| WSW | 0.0 | 1.1 | 4.4 | 5.7 | 6.1 | 6.2 | | 6.2 |
| W | 0.1 | 1.5 | 7.1 | 10.1 | 11.0 | 11.1 | | 11.1 |
| WNW | 0.0 | 0.8 | 3.6 | 5.3 | 5.8 | 5.9 | | 5.9 |
| NW | 0.1 | 0.9 | 3.4 | 4.7 | 5.2 | 5.3 | | 5.3 |
| NNW | 0.1 | 1.2 | 4.3 | 6.0 | 6.6 | 6.7 | | 6.7 |
| N | 0.1 | 0.8 | 2.8 | 3.7 | 4.0 | 4.1 | | 4.1 |



APPENDIX D

**STRATIGRAPHIC CROSS-SECTION AND OTHER PERTINENT
INFORMATION CONCERNING THE REGIONAL GEOLOGY OF
THE AREA (KROHELSKI AND BROWN, 1986)**



Base from U.S.G.S.
 State base 1:1,000,000, 1968

Figure 1. Location of the study area.

HYDROGEOLOGY GEOLOGY

The descriptions of rock units presented in this report are based on drill cuttings obtained from Brown County wells. Lithologies of many of the rock formations are not uniform areally and, in fact, can differ over short distances. The formations include aquifers and confining units. The lithology and areal extent of the rocks and sediments in Brown County are summarized in table 1. The stratigraphy and nomenclature used in this report is that of Mudrey, Brown, and Greenberg (1982).

Bedrock Geology

By *B. A. Brown*¹

Brown County is underlain by Paleozoic sedimentary rocks that range in age from Cambrian to Silurian. The rocks rest directly on Precambrian basement rocks that consist predominantly of red granite. The Paleozoic rocks and the Precambrian surface slope to the east beneath Lake Michigan toward the Michigan Basin at about 30 to 40 ft/mi (fig. 2). Erosion has removed the Silurian rocks and the Maquoketa Formation in the western part of the county (fig. 3). The total thickness of the Paleozoic rocks ranges from 200 ft in the west to about 1,600 ft in eastern Brown County.

Cambrian System

The basal unit of the Cambrian is the Elk Mound Group, which overlies the Precambrian. The group normally consists of, in ascending order, the Mount Simon, Eau Claire, and Wonewoc Formations. The group name is used because the Eau Claire Formation cannot be identified in Brown County, and the sandstones of the Mount Simon and Wonewoc Formations commonly cannot be distinguished from one another.

In areas where these formations are distinguishable, the Mount Simon Formation consists of poorly cemented, subangular, fine to very fine-grained sandstone, which may locally be silty. The Wonewoc Formation consists of poorly cemented, subrounded medium to coarse-grained sandstone.

The Tunnel City Group overlies the Elk Mound Group and includes the Lone Rock and the Mazomanie Formations. The Mazomanie Formation is a fine to medium-grained, feldspathic sandstone. The Lone Rock Formation ranges from a dolomitic, feldspathic, glauconitic siltstone or sandstone to a sandy glauconitic dolomite. The Mazomanie and Lone Rock Formations are laterally equivalent facies and either or both facies may be present in the same well. Where fine-grained dolomite of the Lone Rock facies is present, it is difficult to identify the upper contact of the Tunnel City Group because of the similarity of these rocks to the overlying St. Lawrence Formation.

The Trempealeau Group, which consists of the St. Lawrence Formation and Jordan Formations, overlies the Tunnel City Group. The St. Lawrence Formation is a silty, shaly dolomite that commonly contains glauconite. The Jordan Formation can locally be subdivided into the Van Oser

and Coon Valley Members. The Van Oser Member consists of very fine to very coarse sandstone, commonly dolomitic that contains minor glauconite. The Coon Valley Member consists of dolomite that contains variable amounts of sand, shale, and minor glauconite. This member is difficult to identify from drill cuttings. The Trempealeau Group can be subdivided only where the Van Oser Member is present.

Ordovician System

The Prairie du Chien Group consists of the Oneota and Shakopee Formations. The Shakopee Formation is further subdivided into the lower New Richmond Member and upper Willow River Member. The Oneota Formation and the Willow River Member are very similar, consisting of massive dolomite with minor limestone and oolitic chert. The New Richmond Member consists of sandstone, shaly sandstone, or dolomitic sandstone. The Prairie du Chien Group can be subdivided only in wells where the New Richmond is present. Erosion that occurred prior to deposition of the overlying Ancell Group has removed the Prairie du Chien Group rocks in some areas of Brown County.

The Ancell Group consists of the St. Peter and Glenwood Formation. The St. Peter Formation is composed of two members—the lower Readstown Member, which consists of sandy shale with chert layers, and the overlying Tonti Member, which consists of poorly cemented fine to medium-grained sandstone. The overlying Glenwood Formation is a silty sandstone.

The St. Peter Formation varies areally in thickness because of erosion of the Prairie du Chien strata in pre-St. Peter time. The St. Peter reaches a maximum thickness of up to 300 ft under the Fox River Valley in the area of De Pere, but thins rapidly to as little as 40 ft several miles to the east and west.

The Ancell Group is overlain by the Sinnipee Group, which includes the Platteville, Decorah, and Galena Formations. The Platteville and Galena Formations consist of dolomite that contains fossil fragments and shaly layers. The Galena is distinguished from the Platteville by its chert content. The Decorah Formation is predominantly shale. The Sinnipee Group can be subdivided with certainty only in wells where shale of the Decorah Formation is present between the underlying Platteville and overlying Galena Formations.

The Maquoketa Formation overlies the Sinnipee Group in the area to the east of the Fox River. This formation consists of the Scales Member (a dolomitic shale), which is overlain by the Fort Atkinson Member (a fossiliferous dolomite), which is overlain by the Brainerd Member (another dolomitic shale). The Maquoketa Formation can be subdivided only in northeastern Brown County, where the Fort Atkinson Member is present.

Silurian System

The rocks of the Silurian System are not subdivided in the subsurface of Brown County. These rocks underlie the area east of the Fox River lowland, and consist of massive

¹ Wisconsin Geological and Natural History Survey.

dolomite containing variable amounts of fossil fragments, calcite and gypsum crystals, pyrite, and minor limestone.

Pleistocene

Pleistocene deposits overlie the Paleozoic rock in Brown County and are more than 50 ft thick in most places

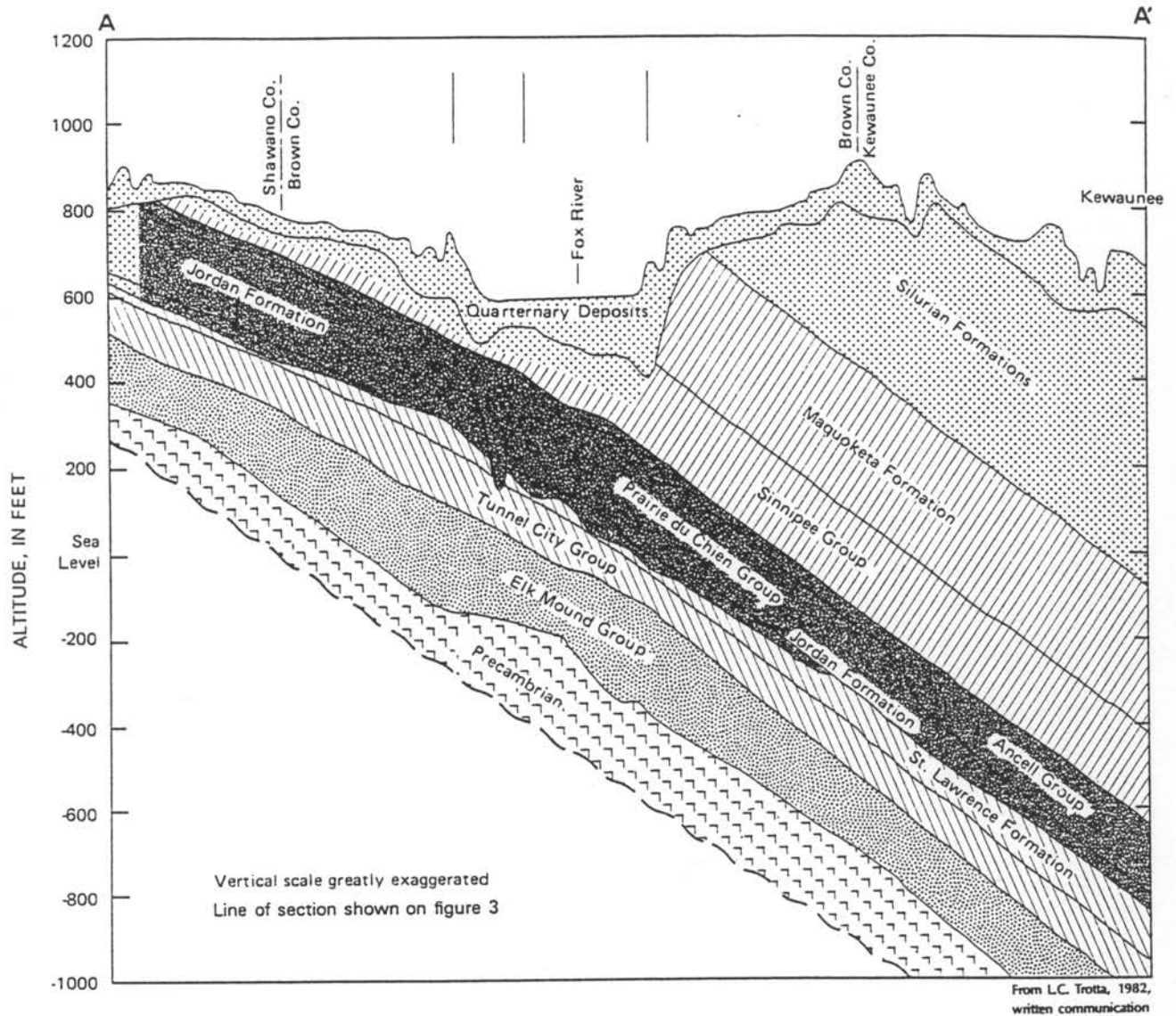
and more than 200 ft thick in the southwestern part of the county (fig. 4). These unconsolidated deposits were mapped by Need (1983) and the following description is based on that work.

Several glacial episodes are recorded in Brown County Pleistocene deposits. Seven tills and their associated fluvial







Table 1. Stratigraphy of Brown County

| Age | Rock unit | Lithology | Areal extent |
|---------------------------|--|--|--|
| Quaternary Pleistocene | Kewaunee Formation | Fluvial, lacustrine, wind blown, and peat deposits, and till | Predominantly fine-grained till except for Fox River valley and area adjacent to west side of Green Bay where lacustrine silt and clay are common. Sand and gravel deposits of small areal extent are present throughout the county. |
| | Horicon Formation | | |
| Silurian | Undifferentiated | Dolomite with varying amounts of fossil fragments, gypsum crystals, pyrite, and limestone. | Subcrops east of the Silurian escarpment. |
| | Maquoketa Formation Brainerd Member Fort Atkinson Member Scales Member | Predominantly dolomitic shale. The Fort Atkinson Member is fossiliferous dolomite. | Subcrops in a band generally less than 3 mi wide west of the Silurian escarpment. Present directly beneath the Silurian dolomite. |
| | Sinnipee Group Galena Formation Decorah Formation Platteville Formation | Galena and Platteville Formations are dolomite. The Decorah Formation is shale. | Subcrops just east of the Fox River and throughout the county west of the river. |
| Ordovician | Ancell Group Glenwood Formation St. Peter Formation Tonti Member Readstown Member | The Glenwood Formation is a silty sandstone, the Tonti Member is a fine- to medium-grained sandstone and the Readstown Member is a sandy shale. | Commonly present in the Fox River valley but thins rapidly east and west of the valley. |
| | Prairie du Chien Group Shakopee Formation Willow River Member New Richmond Member Oneota Formation | The Prairie du Chien Group is generally dolomite with varying amounts of oolitic chert. The group can be subdivided only when the New Richmond Member, a sandstone, shaly sandstone, or dolomitic sandstone, is present. | Thin or absent where the St. Peter Sandstone is thick (Fox River valley). |
| | Trempealeau Group Jordan Formation St. Lawrence Formation | The Jordan Formation is a fine- to medium-grained sandstone. The St. Lawrence Formation is a silty glauconitic dolomite. | Present throughout the county. |
| Cambrian | Tunnel City Group Mazomanie Formation Lone Rock Formation | The Mazomanie Formation is a fine- to medium-grained sandstone. The Lone Rock Formation is a silty sandstone to a sandy dolomite. | Present throughout the county. |
| | Elk Mound Group Wonowoc Formation Eau Claire Formation Mount Simon Formation | The members of the Elk Mound Group are usually not differentiated. Where distinguishable the units generally present are a very fine to fine-grained sandstone and a medium- to coarse-grained sandstone. | Present throughout the county. |
| | Precambrian | Red granite | Basement rock throughout the county. |

1/ The stratigraphic nomenclature used in this report is that of the Wisconsin Geological and Natural History Survey and does not necessarily follow usage of the U.S. Geological Survey.



EXPLANATION

-  UPPER AQUIFER
-  MAQUOKETA-SINNIPEE CONFINING UNIT
-  ST. PETER AQUIFER
-  ST. LAWRENCE CONFINING UNIT
-  ELK MOUND AQUIFER
-  PRECAMBRIAN CONFINING UNIT

 TURNING POINT OF SECTION

Scale 1:500,000

0 5 10 MILES
0 5 10 KILOMETERS

Figure 2. Hydrogeologic section through study area.

and lacustrine deposits are present in the Brown County area. Tills were deposited by the Green Bay and Lake Michigan Lobes of the ice sheet. Fluvial sand and gravel were deposited by glacial meltwater from the lobes. Lacustrine sediment, generally fine grained (silt or clay), was deposited in two ice dammed lakes—Nipissing Lake and Lake Oshkosh.

Modern sediments deposited by wind, water, and the accumulation of organic matter are also present in Brown County. Figure 5 shows the areal distribution of groupings of Pleistocene surface deposits in Brown County. The groupings are till, silt and clay, and sand and gravel. Figure 6 is an east-west geologic section through northern Brown

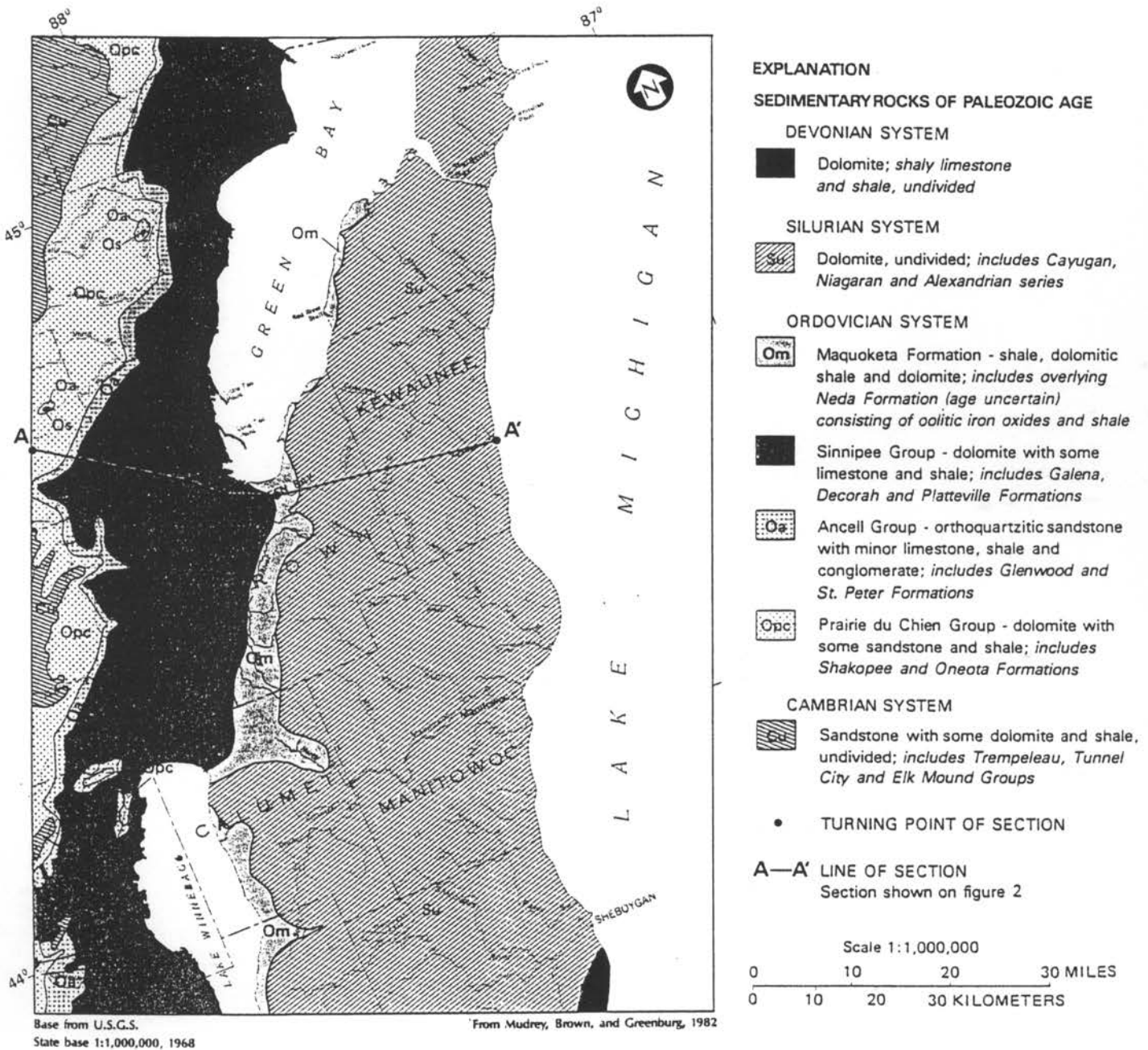


Figure 3. Bedrock geology.

County showing the vertical distribution of Pleistocene deposits.

The following is a brief description of the Pleistocene deposits in order of their relative age from youngest to oldest.

Modern Deposits

1. *Modern stream sediment* is silt loam and silt channel-fill and flood-plain deposits. It is present adjacent to most county streams.
2. *Windblown sand* is well-sorted fine sand in transverse dunes present in northwestern Brown County.
3. *Organic and hillslope sediment in topographic depressions* is loam to silty clay slopewash sediment overlain by peat and muck. It is present in small areas throughout the county.

Kewaunee Formation

1. *Nipissing Lake Plain and Lake Oshkosh Plain sediment* vary from clay to silt loam. This sediment is present at the surface in the Fox River lowland.
2. *Stream sediment in spillways* is gravelly sand, sand, and sandy gravel point-bar and channel-lag deposits in steep-walled channels that drained proglacial Lake Oshkosh. It is present in two locations in eastern Brown County.
3. *Till of the Middle Inlet Member* is reddish brown, calcareous, loam till and is the surface unit in northwestern Brown County. It is present discontinuously in the subsurface in the Fox River lowland near the west side of Green Bay.
4. *Till of the Glenmore Member* is reddish brown, calcareous, silty clay loam till that is the surface unit throughout most of eastern Brown County. It has been identified in the subsurface in the Fox River lowland west of the Fox River.
5. *The Duck Creek Ridge Complex* is sediment of the Middle Inlet and Kirby Lake Members, stream sediment, and clayey lake sediment. It is present in a glacially eroded, elongated ridge near the east side of Duck Creek.
6. *Meltwater-stream sediment exposed by glacial and postglacial erosion* is well-sorted sand exposed along elongated ridges and steep slopes. It is present at the surface and in the subsurface in western and northwestern Brown County.
7. *Till of the Kirby Lake Member* is reddish brown, calcareous, clay loam to silty clay loam till. It is not exposed at the surface but is present in the subsurface throughout northwestern and west-central Brown County.
8. *Till of the Chilton Member* is reddish brown, calcareous, silty clay loam till. It is exposed at the surface in southern Brown County and present in the subsurface in the Fox River lowland south of Green Bay.

9. *Till of the Valders Member* is reddish brown, calcareous, silt loam till and is exposed in southeastern Brown County but is not present to any significant extent in the subsurface.

10. *Clayey offshore sediment exposed by glacial and stream erosion* is silty clay loam, silty clay, and clay that was deposited in proglacial lakes predating the Chilton and Kirby Lake Members. This unit is exposed at the surface in northwestern Brown County and in the subsurface throughout most of the Fox River lowland.
11. *Meltwater stream sediment* is gravelly sand, sand, and sandy gravel with minor amounts of silt loam. It is present at the surface in southern Brown County near the Branch River and is discontinuous in the subsurface in the Fox River lowland.
12. *Till of the Branch River Member* is light reddish brown, calcareous, loam till. It is exposed at the surface in southern Brown County and around the margins of an erosional window of the Wayside till in northeastern Brown County. The Branch River Member is also thought to be present in the subsurface throughout the eastern part of the county.

Horicon Formation

1. *Till of the Wayside Member* is light-grayish brown, calcareous, stony loam till and is exposed at the surface in southern Brown County.
2. *Meltwater-stream sediment* is sand and gravel, discontinuous in the subsurface in eastern Brown County.

AQUIFERS AND CONFINING UNITS

The complex hydrogeologic system in the Brown County area consists of aquifers and confining units. The hydrogeologic system includes an upper aquifer and deep aquifers separated by confining units. Previous studies have defined the "sandstone aquifer" in the Brown County area to include Cambrian and Ordovician Formations older than the Maquoketa Formation (Donohue, 1976; Drescher, 1953; Knowles, 1964). Although it was recognized in previous studies that the "sandstone aquifer" did not have uniform hydraulic properties and was not a single aquifer, it was considered a single aquifer because hydraulic data on individual formations were not available. Most high-capacity wells in Brown County are drilled through and open to most of the formations of the "sandstone aquifer".

The division of aquifers and confining units in this report is based on the composition and hydraulic information of the rock groups or formations present in the Brown County area. Figure 2 shows rock groups and formations present in the Brown County area and the aquifers and confining units defined in this report. The general range in thickness of the aquifers and confining units can be seen in figures 7 and 15a. Table 2 lists hydraulic parameters for the aquifers and confining units. The locations of pump tests and

APPENDIX E

**BATHYMETRY INFORMATION AVAILABLE FROM THE NOAA
RECREATIONAL CHARTS FOR LAKE WINNEBAGO AND THE
LOWER FOX RIVER**

LAKE WINNEBAGO AND LOWER FOX RIVER WISCONSIN



RADAR REFLECTORS
Radar reflectors have been placed on many floating aids to navigation. Individual radar reflector identification on these aids has been omitted from this chart.

RACING BUOYS
Racing buoys within the limits of this chart are not shown here. Information may be obtained from the U.S. Coast Guard District Offices as racing and other privately maintained buoys are not all listed in the U.S. Coast Guard Light List.

NOTE B
The channel legend reflects the Corps of Engineers project depth. The Corps of Engineers publishes the controlling depth periodically in the U.S. Coast Guard Local Notice to Mariners. For further information on channel depths, direct inquiries to Office of the District Engineer, Corps of Engineers, Detroit, Michigan.

PLANE OF REFERENCE OF THIS CHART (Low Water Datum)
LAKE WINNEBAGO 745.8 ft.
FOX RIVER (Between Locks) See table below
LAKE MICHIGAN 577.5 ft.
Referred to mean water level at Rimouski, Quebec, International Great Lakes Datum (1885).

AIDS TO NAVIGATION Consult U.S. Coast Guard Light List for supplemental information concerning aids to navigation.

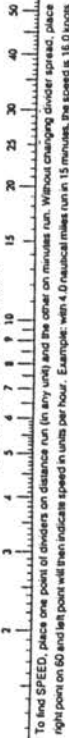
SYMBOLS AND ABBREVIATIONS For complete list of symbols and abbreviations see Chart No. 1.

BRIDGE AND OVERHEAD CABLE CLEARANCES When the water surface is above Low Water Datum, bridge and overhead clearances are reduced correspondingly. For clearances see U.S. Coast Pilot 6.

AUTHORITIES Hydrography and topography by the National Ocean Service, Coast and Geodetic Survey, with additional data from the Corps of Engineers, Geological Survey, and U.S. Coast Guard.

| | | | | | | | | | | | | |
|---------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|---|---|
| FATHOMS | 11 | 10 | 9 | 8 | 7 | 6 | 5 | 4 | 3 | 2 | 1 | 0 |
| FEET | 20 | 18 | 16 | 14 | 12 | 10 | 8 | 6 | 4 | 2 | 0 | 0 |
| METERS | 3.7 | 3.3 | 2.9 | 2.5 | 2.1 | 1.7 | 1.3 | 0.9 | 0.5 | 0.1 | 0 | 0 |

LOGARITHMIC SPEED SCALE



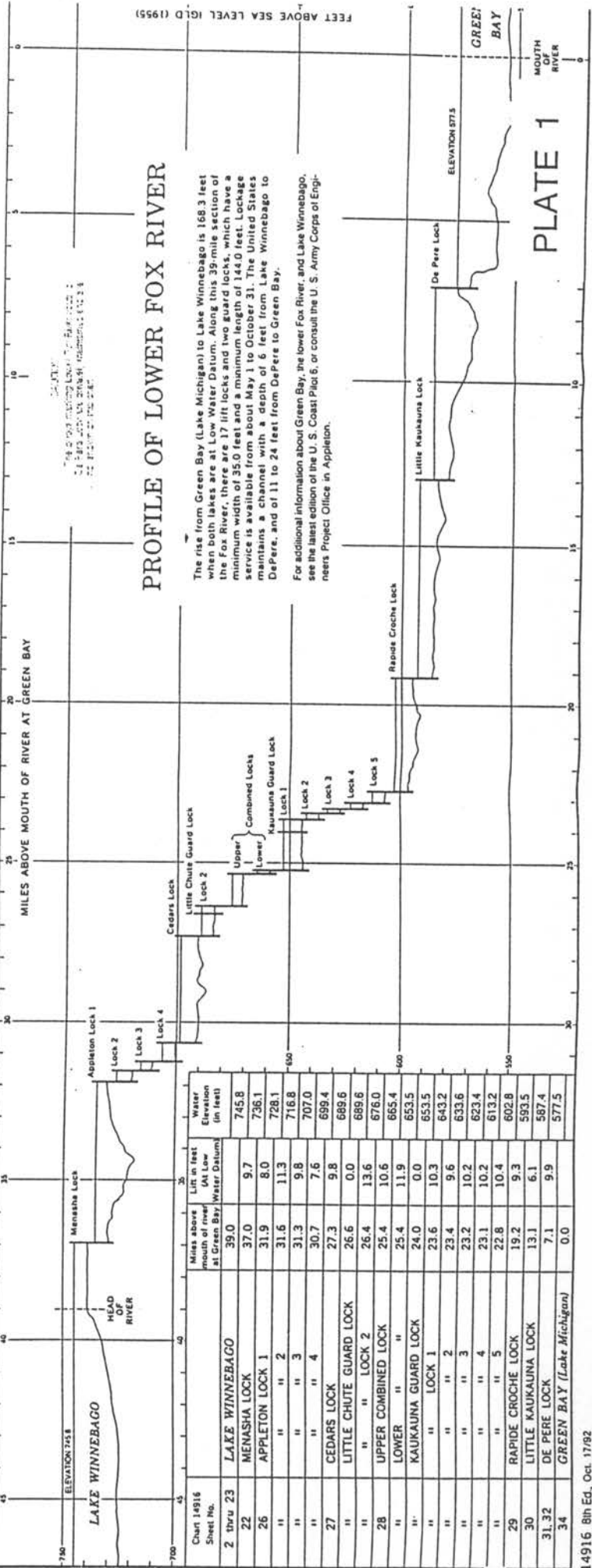
SUPPLEMENTAL INFORMATION
Consult U.S. Coast Pilot 6 for important supplemental information.

POLLUTION REPORTS
Report all spills of oil and hazardous substances to the National Response Center at 800-424-8802 (toll free), or to the nearest U.S. Coast Guard facility if telephone communication is impossible (33 CFR 153).

CAUTION
POTABLE WATER INTAKE (PWI)
Vessels operating in fresh water lakes or rivers shall not discharge sewage or ballast or brackish water into such areas adjacent to domestic water intakes as are designated by the Surgeon General (21 CFR 1250.93). Consult U.S. Coast Pilot 6 for important supplemental information.

NOTE A
Navigation regulations are published in Chapter 1 of the U.S. Coast Pilot. For more information on regulations, consult the U.S. Coast Pilot, or the Office of the Commander, U.S. Coast Guard, 1655 Broadway, New York, N.Y. 10019. For information on regulations, consult the Office of the District Engineer, Corps of Engineers, Detroit, Michigan.

© Firth-Ross Facilities



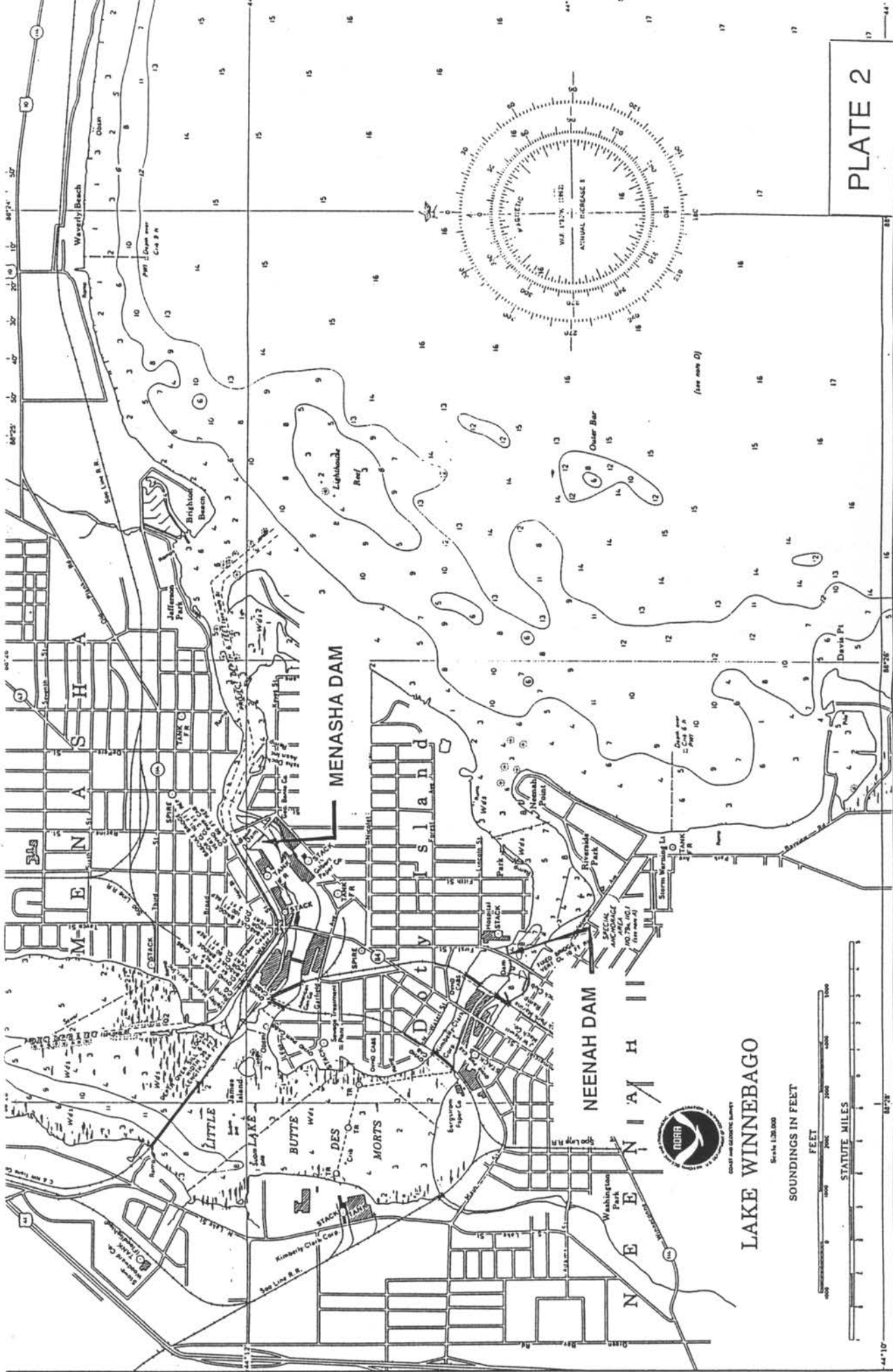


PLATE 2

LAKE WINNEBAGO

SOUNDINGS IN FEET

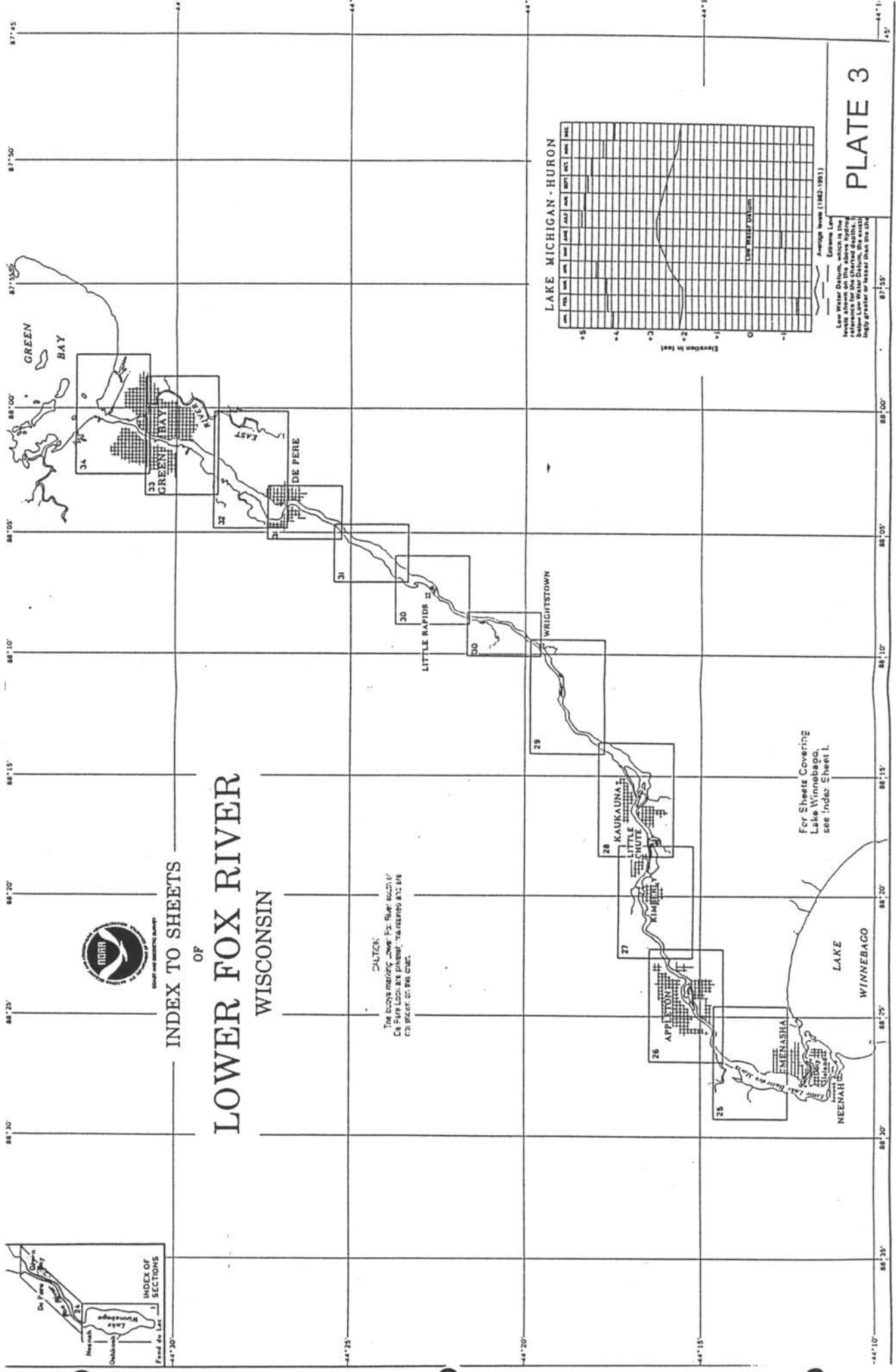
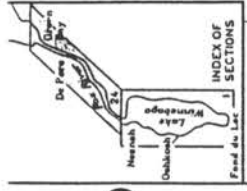
FEET

STATUTE MILES



U.S. COAST AND GEODETIC SURVEY

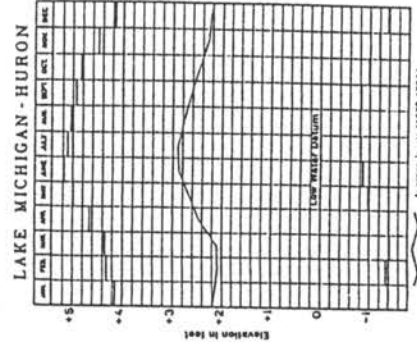
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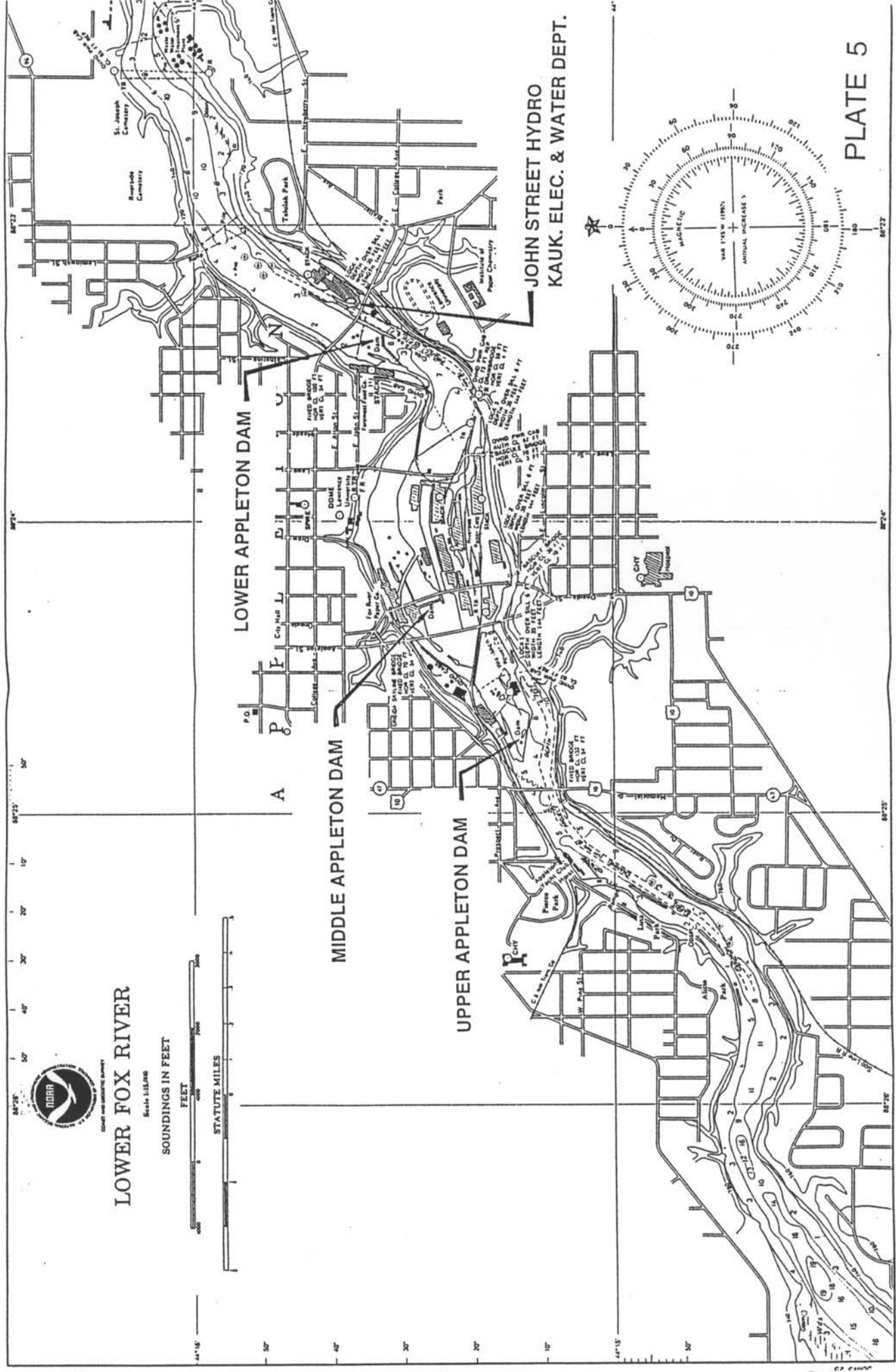
INDEX TO SHEETS
OF
LOWER FOX RIVER
WISCONSIN

CAUTION
The bridge marking above Fox River south of De Pere Lock and Dam project, the channel is 12 feet to 20 feet to the east.

For Sheets Covering Lake Winnebago, see Index Sheet 1.



Low Water Datum is the elevation of the water surface above the datum for the charted depths. It may be higher or lower than the charted datum.



LOWER FOX RIVER

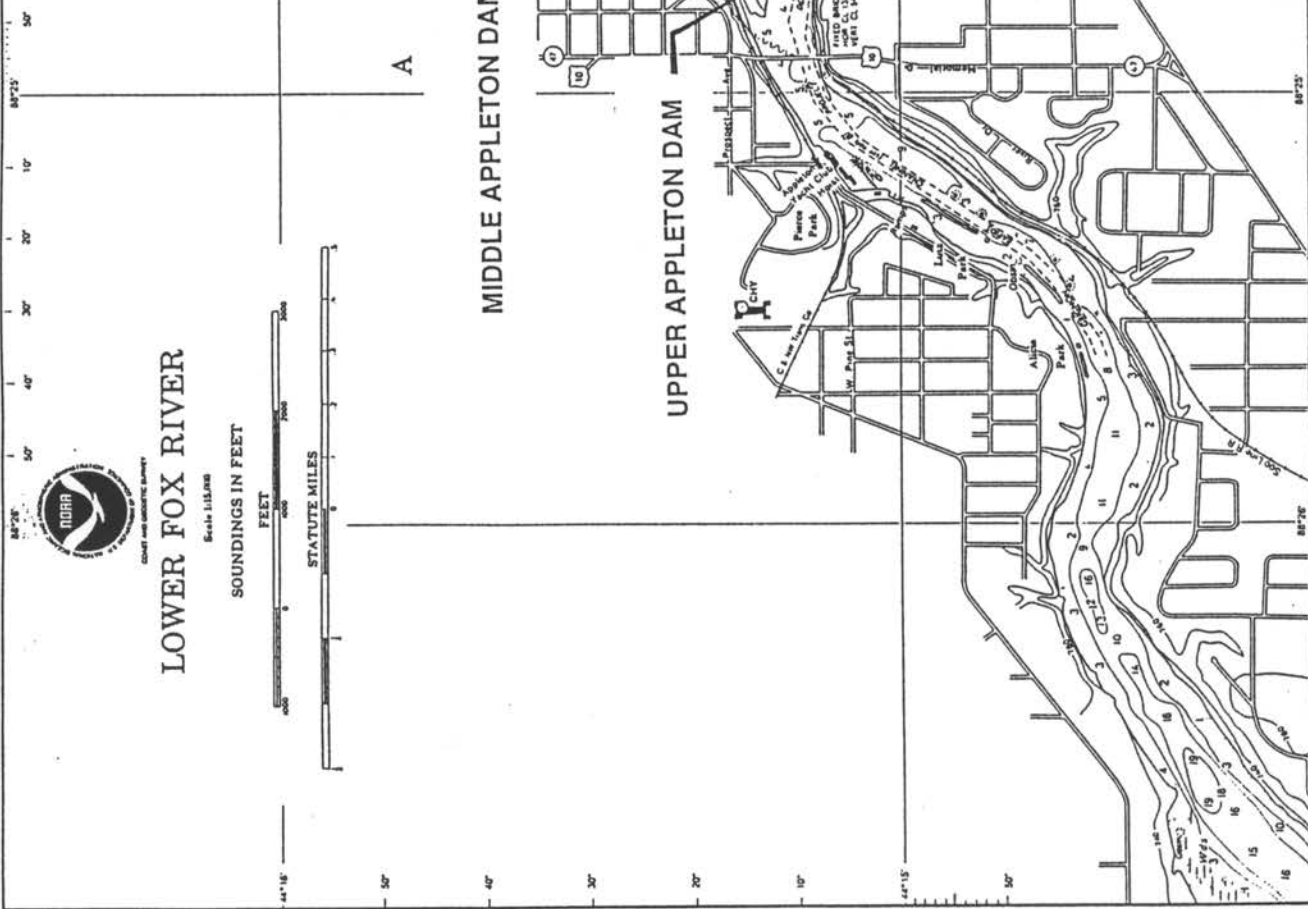
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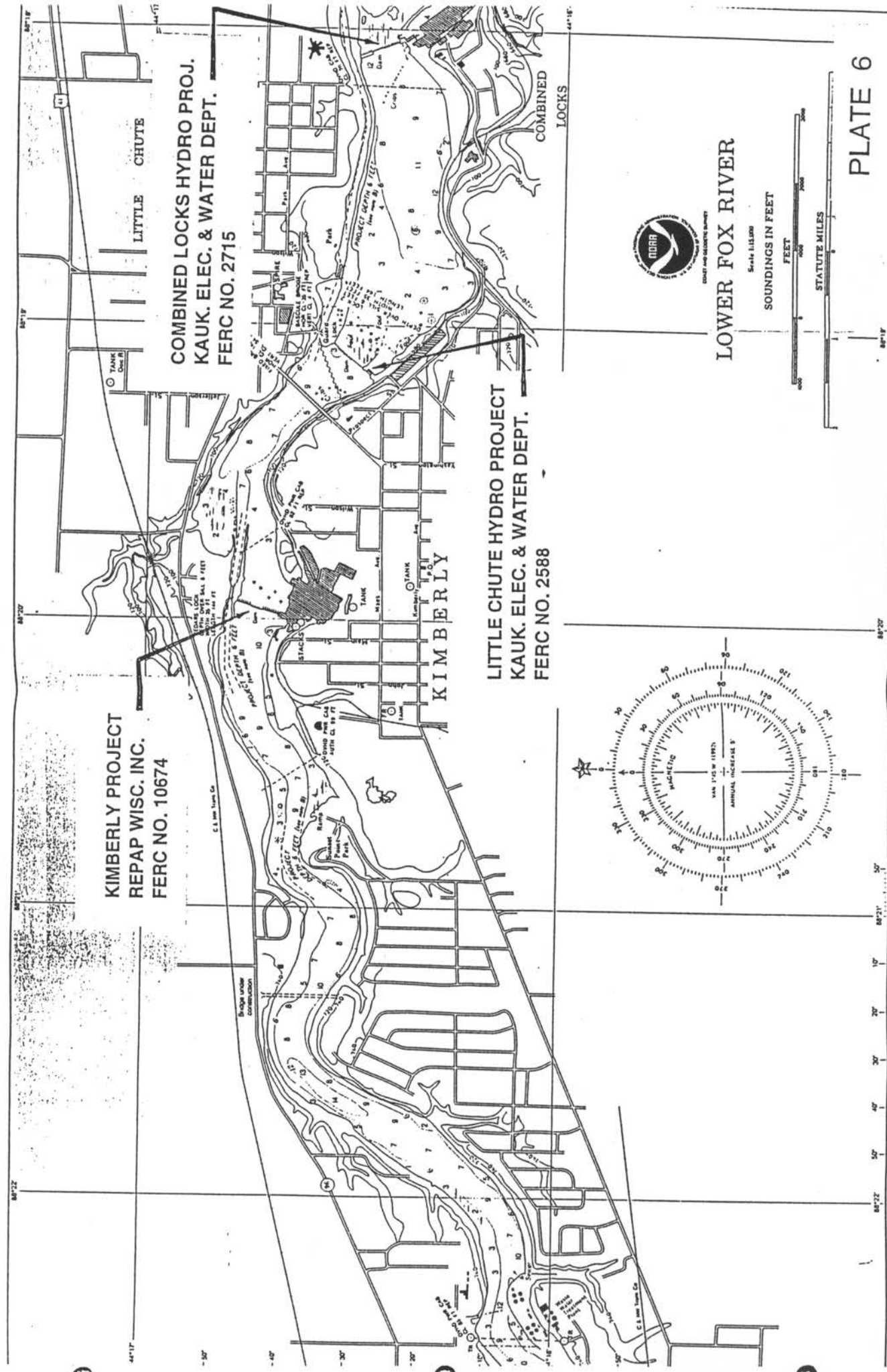
SOUNDINGS IN FEET

FEET



STATUTE MILES





KIMBERLY PROJECT
REPAP WISC. INC.
FERC NO. 10674

COMBINED LOCKS HYDRO PROJ.
KAUK. ELEC. & WATER DEPT.
FERC NO. 2715

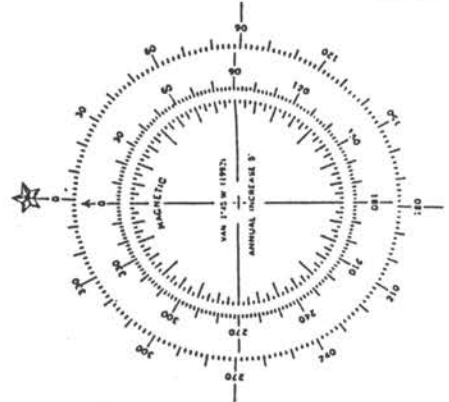
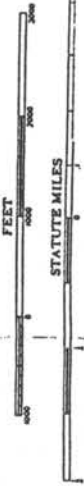
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KAUK. ELEC. & WATER DEPT.
FERC NO. 2588

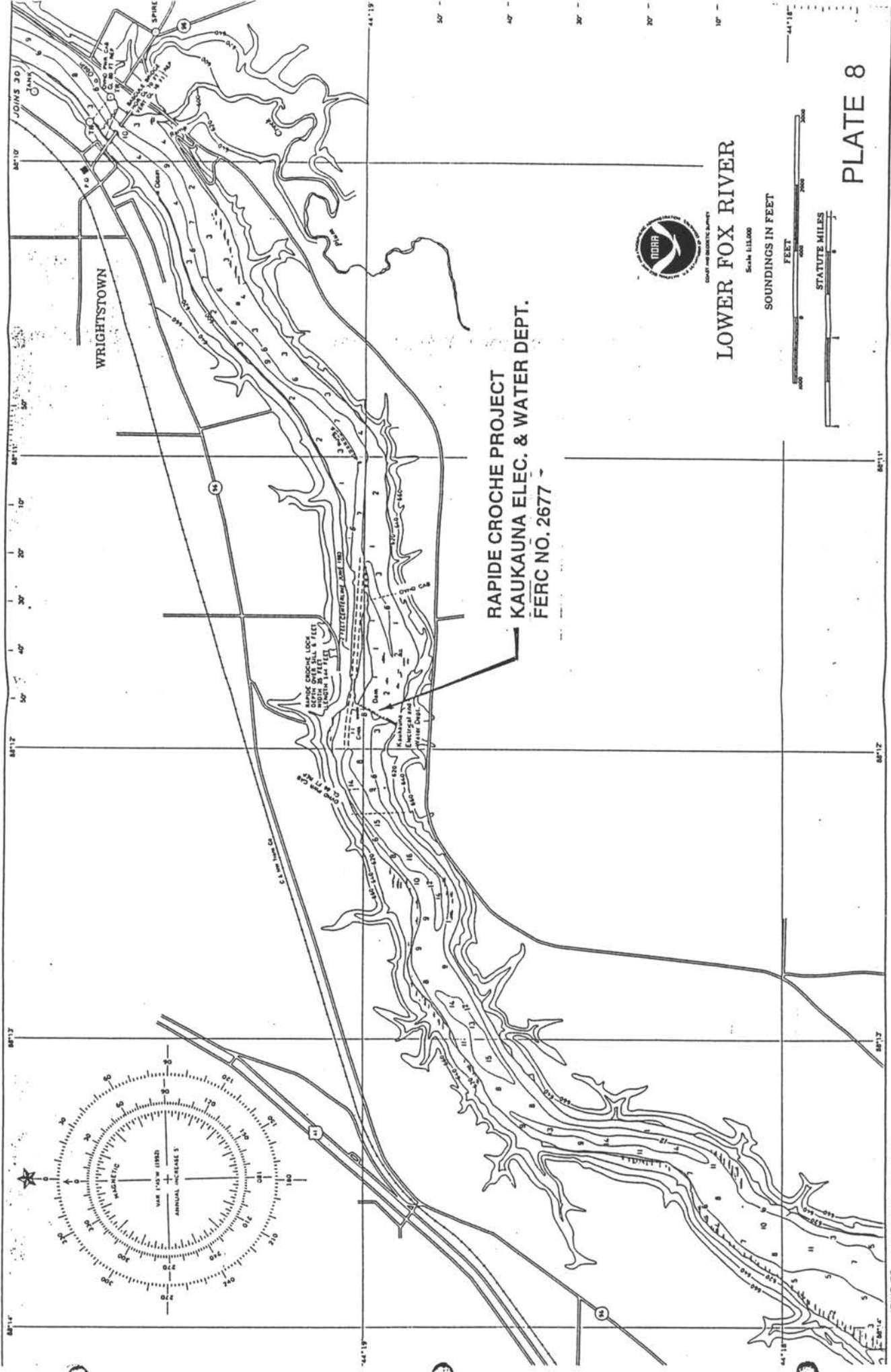
PLATE 6



LOWER FOX RIVER

Scale 1:15,000
 SOUNDINGS IN FEET





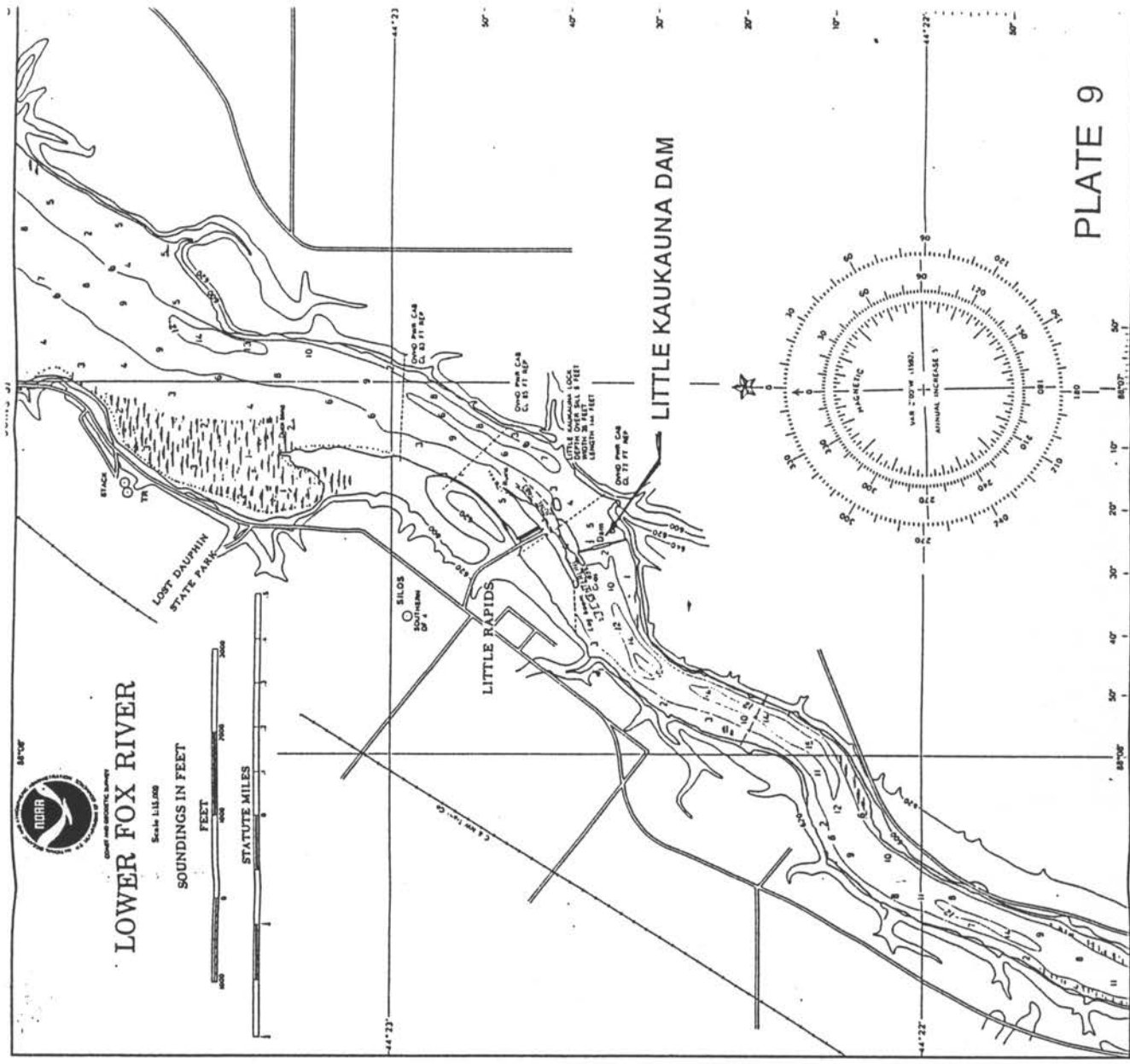
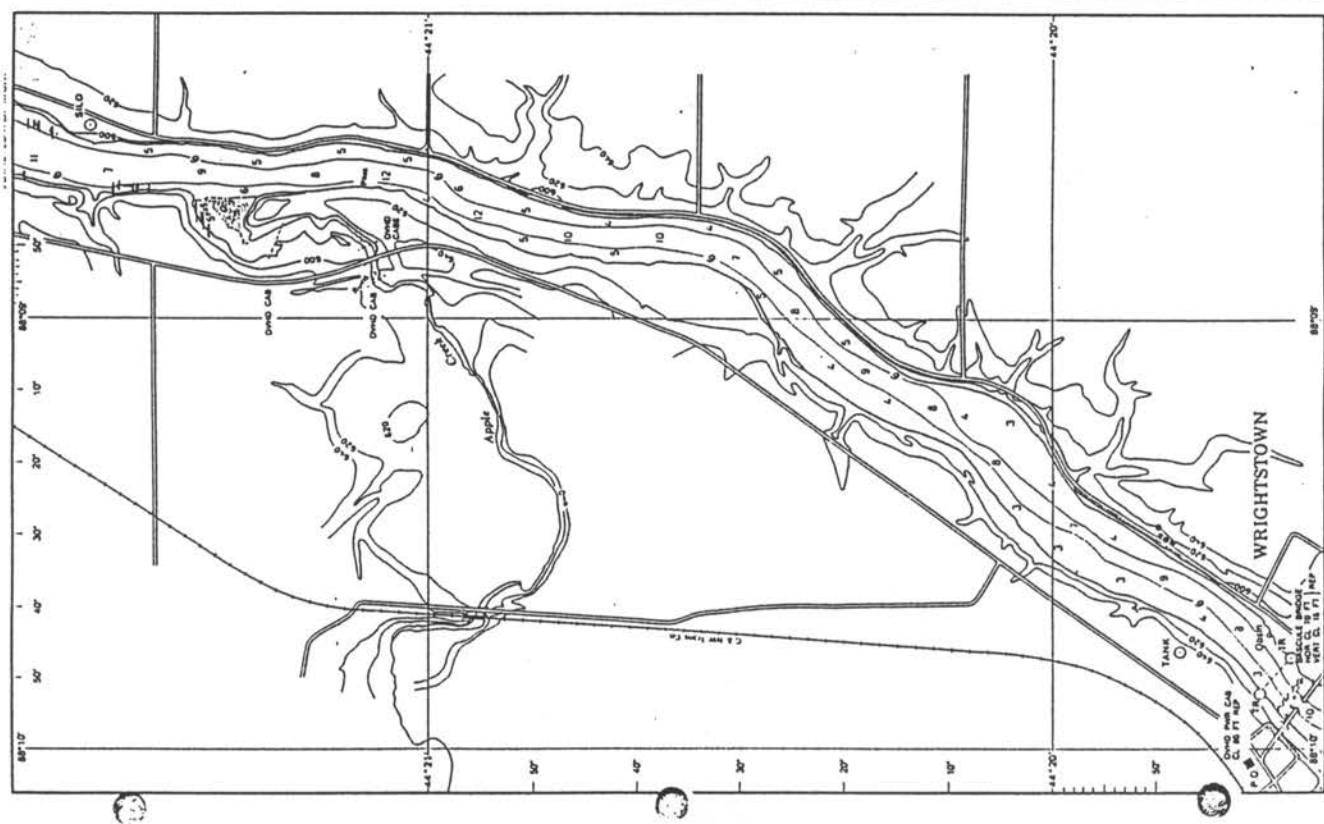
WRIGHTSTOWN

RAPIDS CROCHE PROJECT
KAUKAUNA ELEC. & WATER DEPT.
FERC NO. 2677

LOWER FOX RIVER

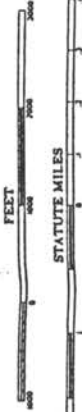
PLATE 8

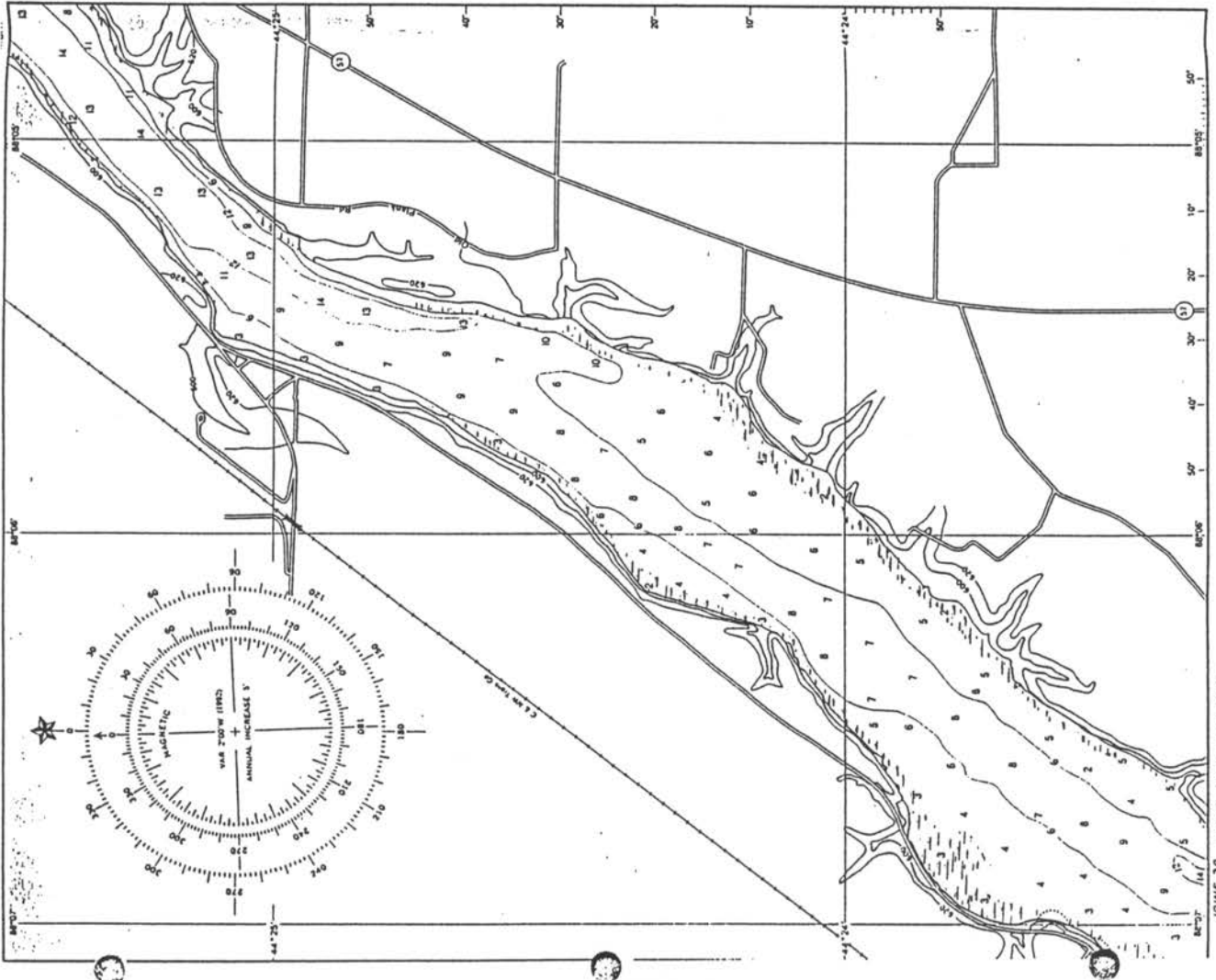
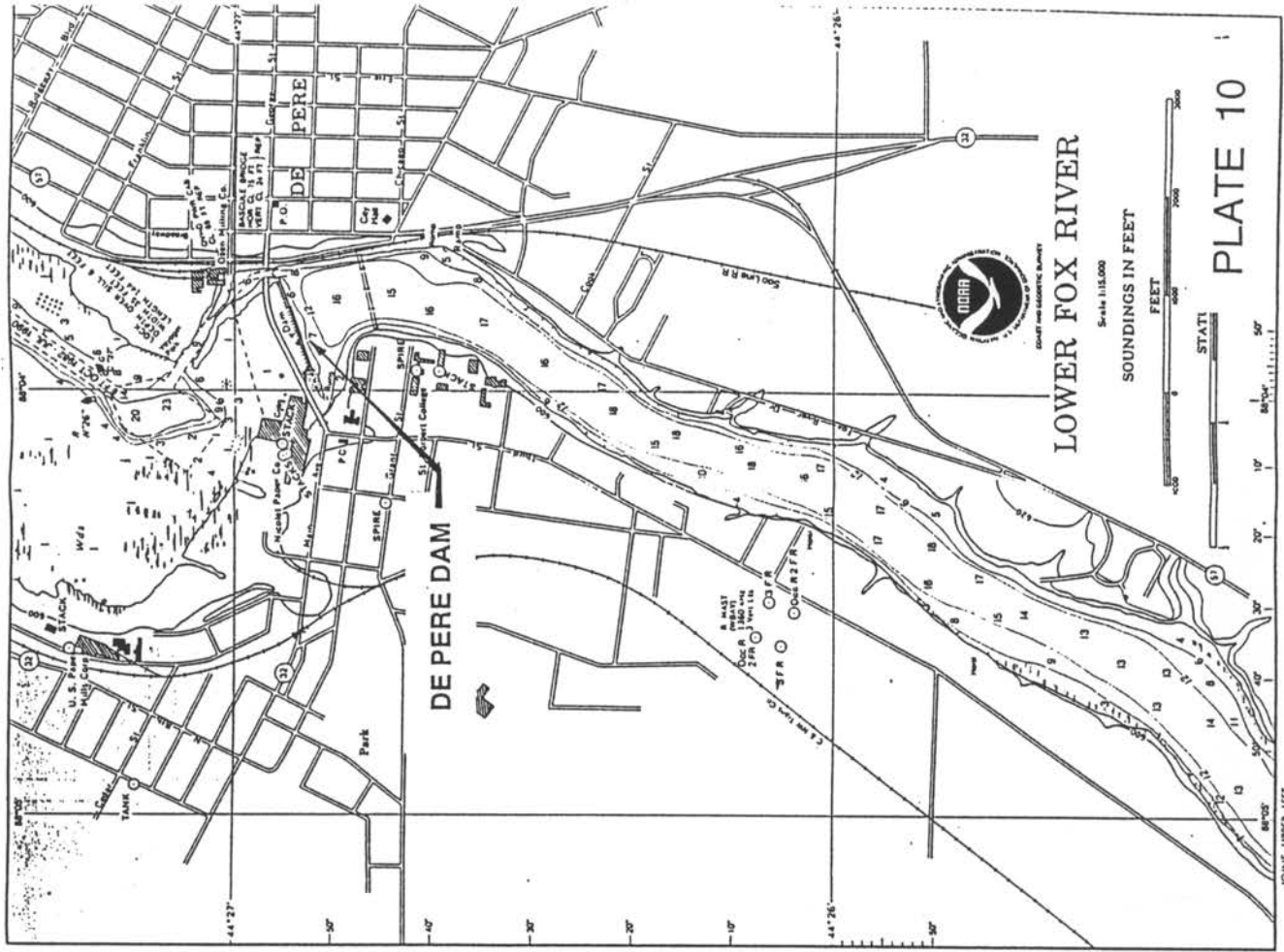
JOINS #8 14916 8th Ed. Oct. 1792



LOWER FOX RIVER

SOUNDINGS IN FEET
Scale 1:15,000







LOWER FOX RIVER
 Scale 1:11,000
 SOUNDINGS IN FEET
 STATUTE MILES



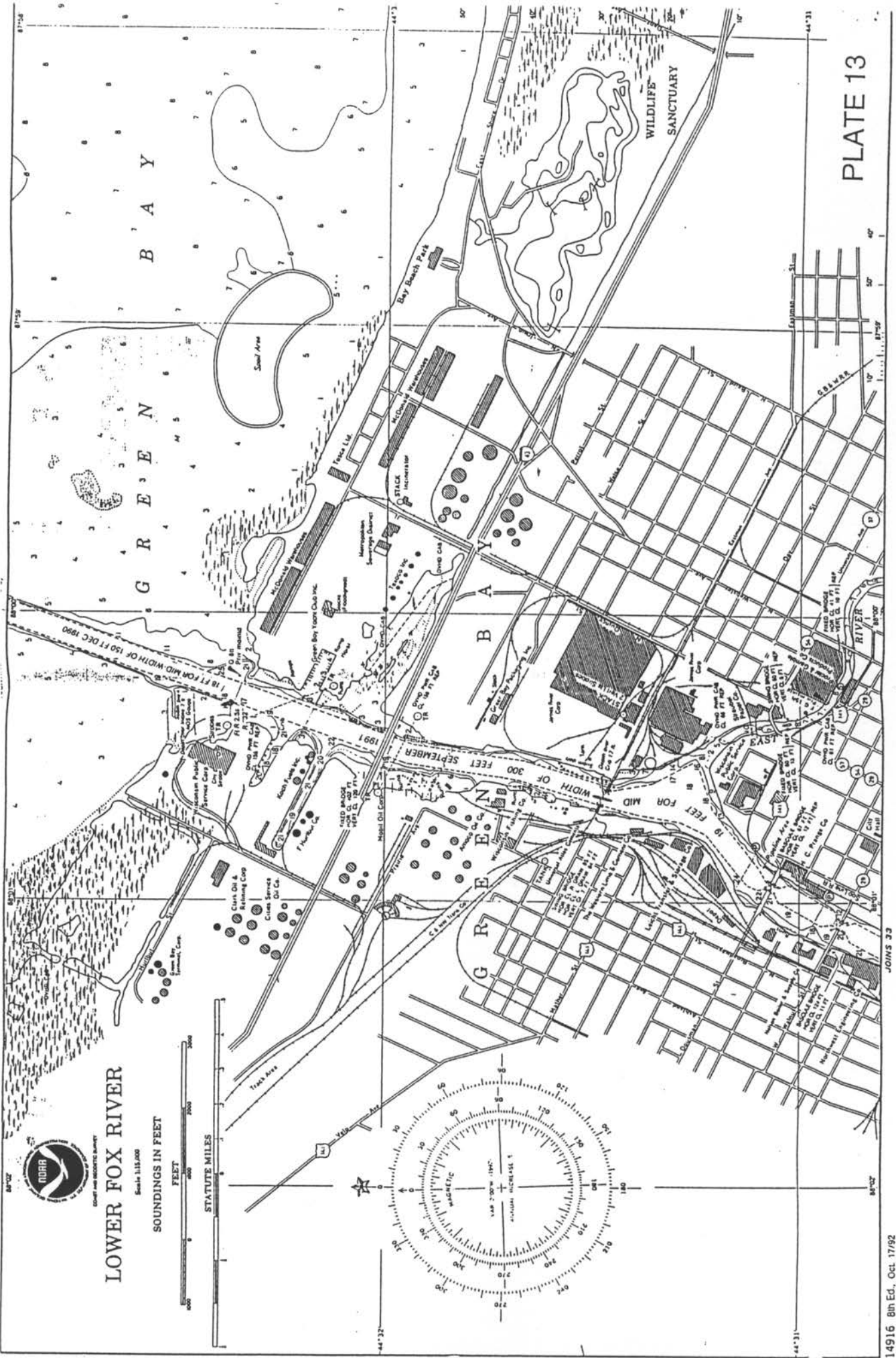
LOWER FOX RIVER

Scale 1:12,500

SOUNDINGS IN FEET



PLATE 12



LOWER FOX RIVER



Scale 1:15,000

SOUNDINGS IN FEET

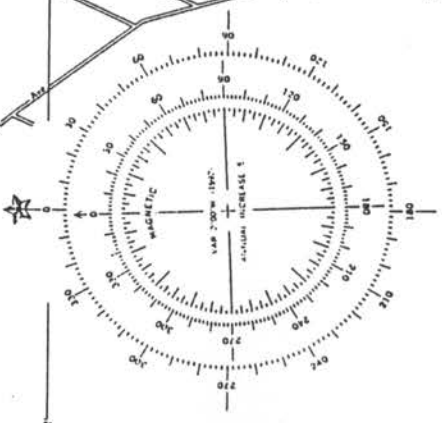


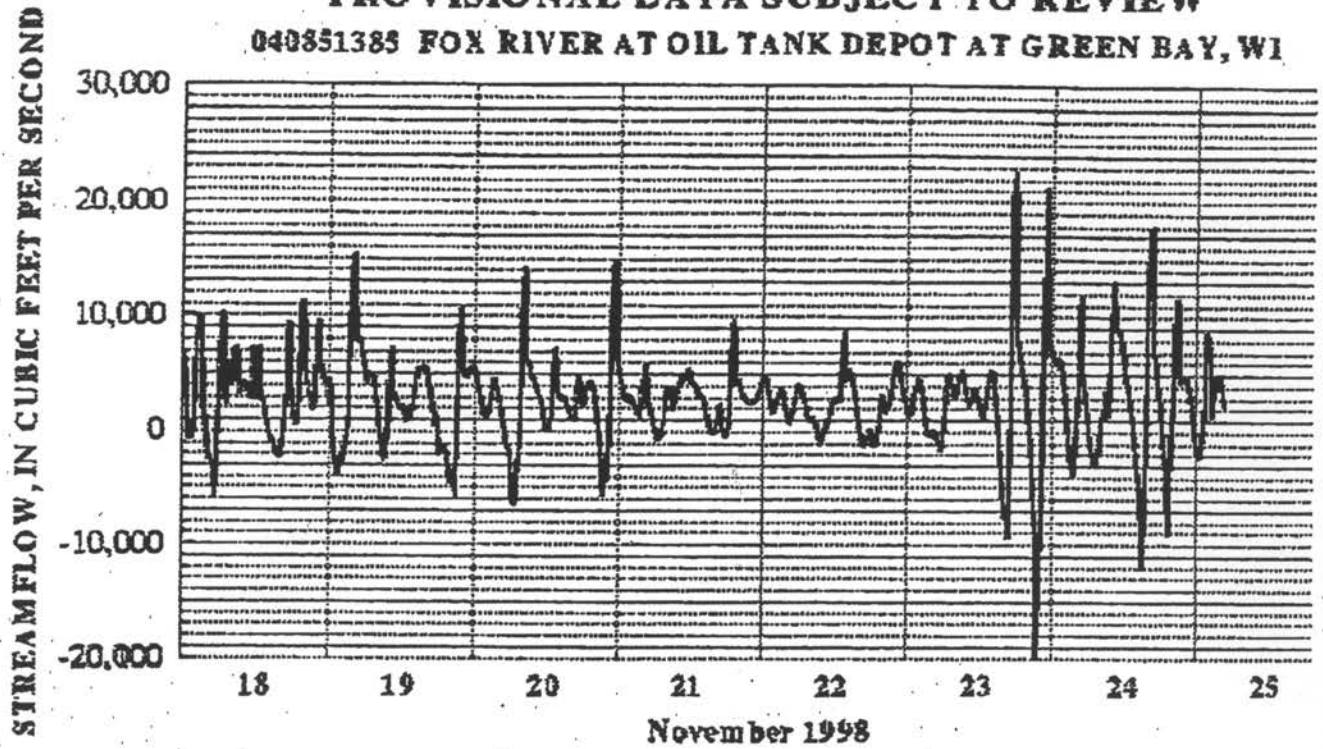
PLATE 13

APPENDIX F

**THE USGS HYDROGRAPHS FOR TWO STORM EVENTS IN
NOVEMBER 1998**

U.S. GEOLOGICAL SURVEY PROVISIONAL DATA SUBJECT TO REVIEW

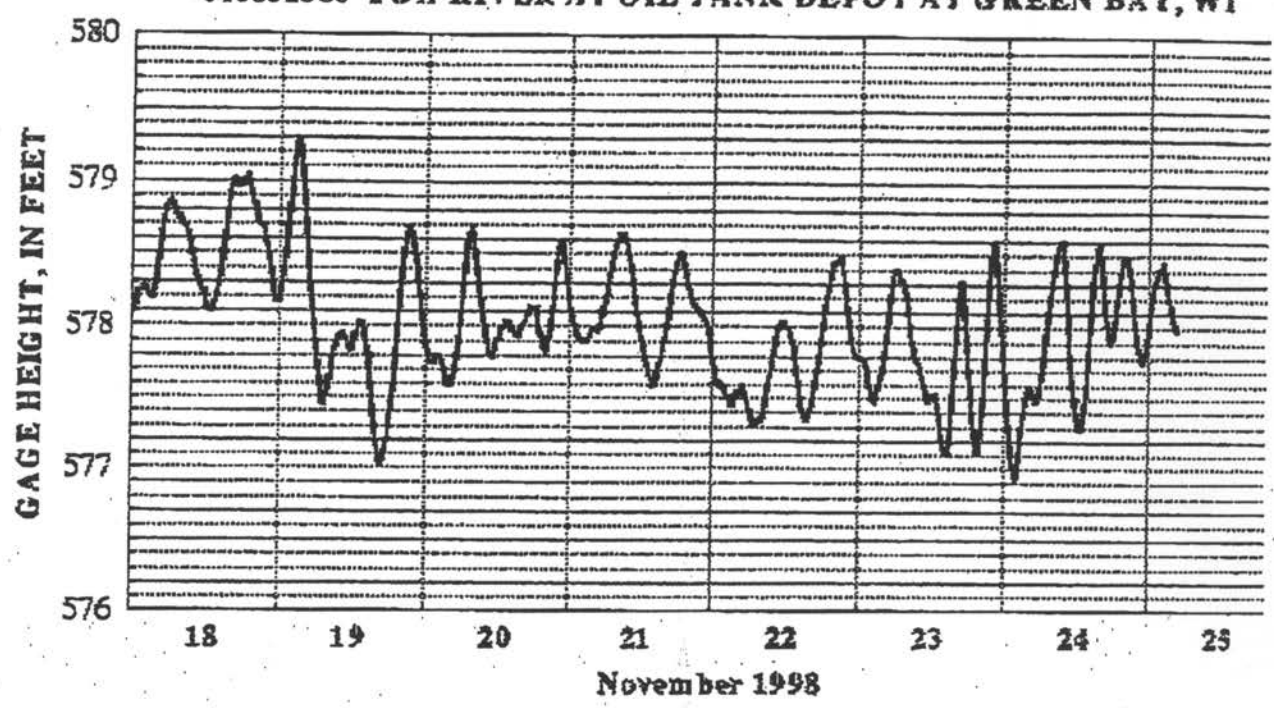
040851385 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI



Updated: 11-25-98 08:02

↑
High flow associated
with rapid seiche activity

U.S. GEOLOGICAL SURVEY
PROVISIONAL DATA SUBJECT TO REVIEW
040851385 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI



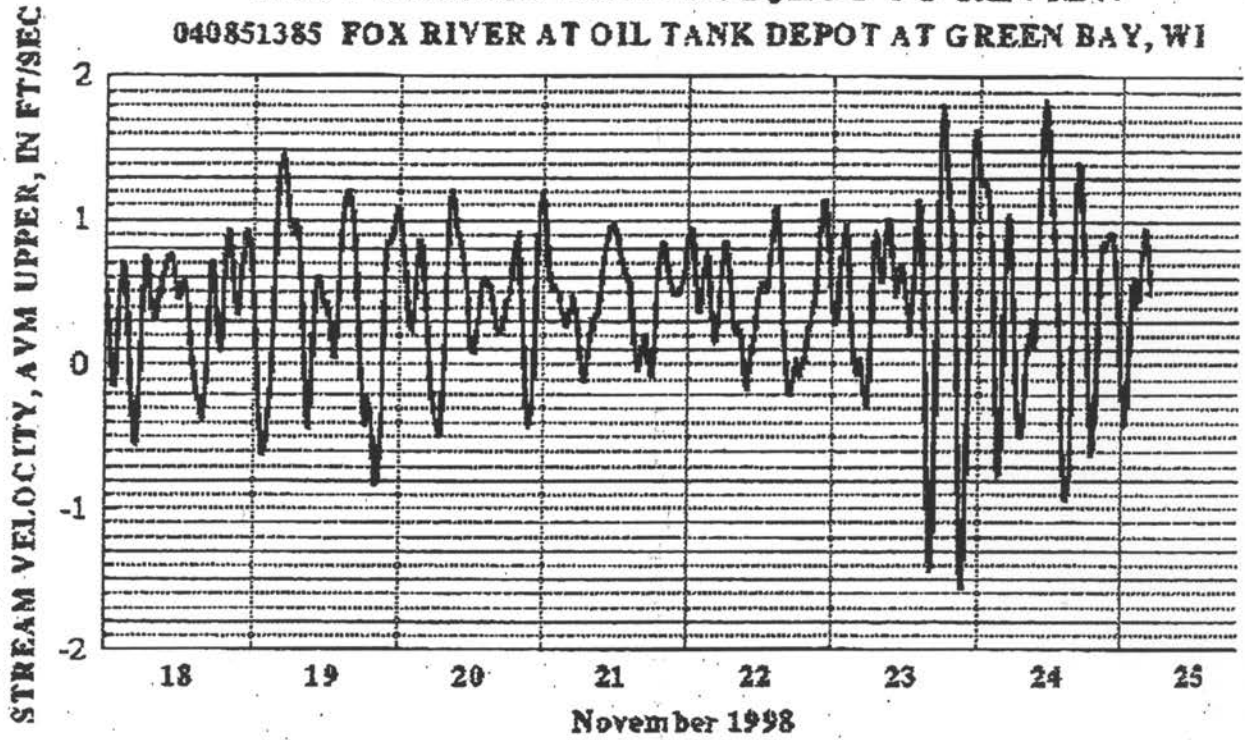
Updated: 11-25-98 07:17

↑
Rapid Seiche activity

218870

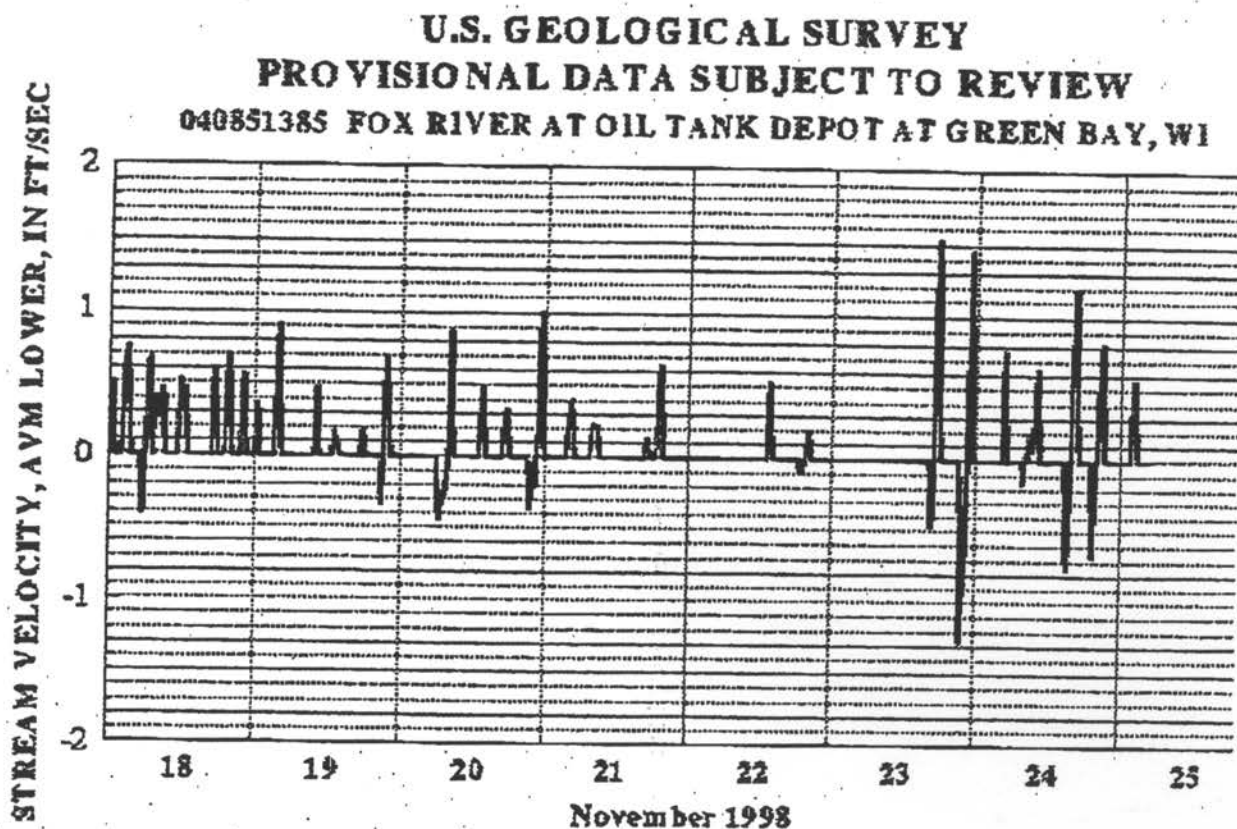
**U.S. GEOLOGICAL SURVEY
PROVISIONAL DATA SUBJECT TO REVIEW**

040851385 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI



Updated: 11-25-98 08:03

218869



Updated: 11-25-98 08:04

Map of region surrounding station

STATION.-- 040851385 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI

LOCATION.--Lat 44°31'43", long 88°01'12" in section 25, T.24 N., R.20 E.,
Brown County, Hydrologic Unit 04030204, about 0.5 mi upstream of Interstate
Highway 43 bridge in Green Bay, and 0.8 mi upstream from mouth.

DRAINAGE AREA.--6,330 square miles.

PERIOD OF RECORD.--October 1988 to current year.

GAGE.--Acoustical Velocity Meter (AVM) system. Two-path transducer installation.

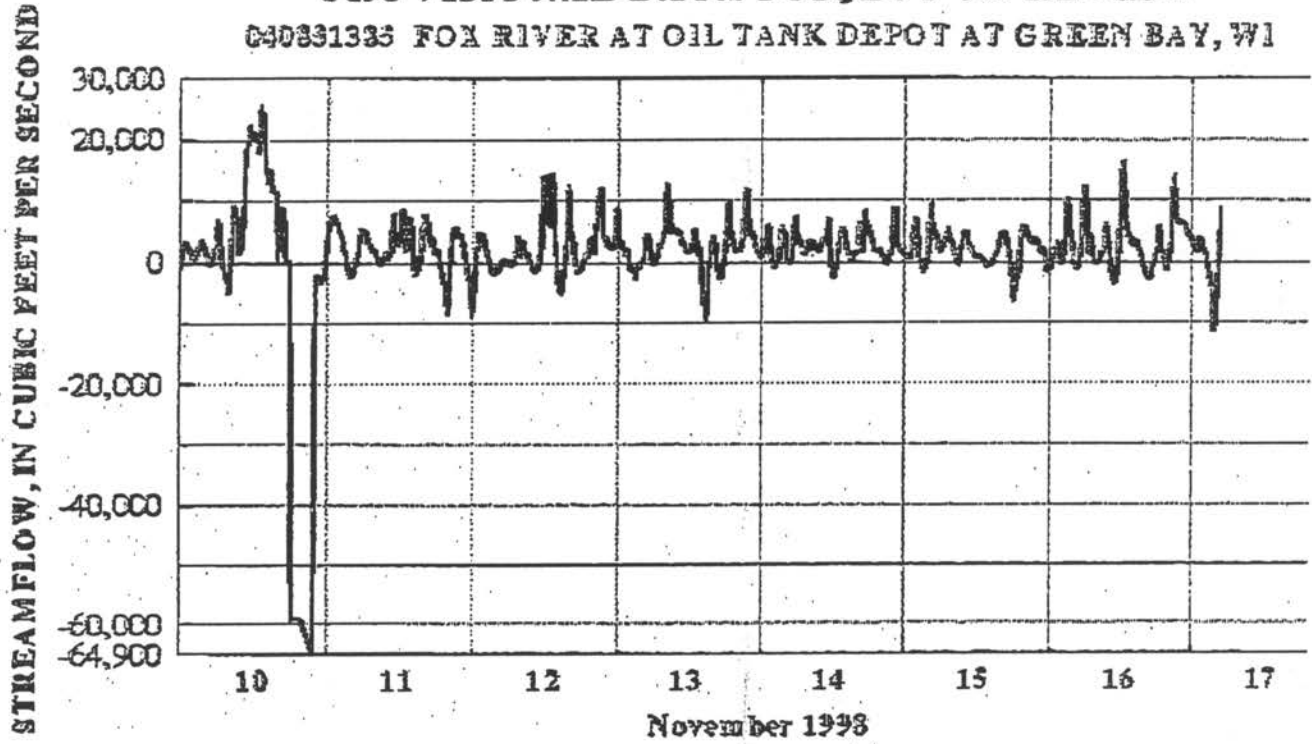
REMARKS.--Gage-height telemeter at station.

Retrieve postscript of discharge hydrograph or retrieve postscript of gage height hydrograph or
retrieve postscript of velocity (upper avm) hydrograph or retrieve postscript of velocity (lower avm)
hydrograph or complete station data from the 1997 Water Resources Data Report

218868

U.S. GEOLOGICAL SURVEY
PROVISIONAL DATA SUBJECT TO REVIEW

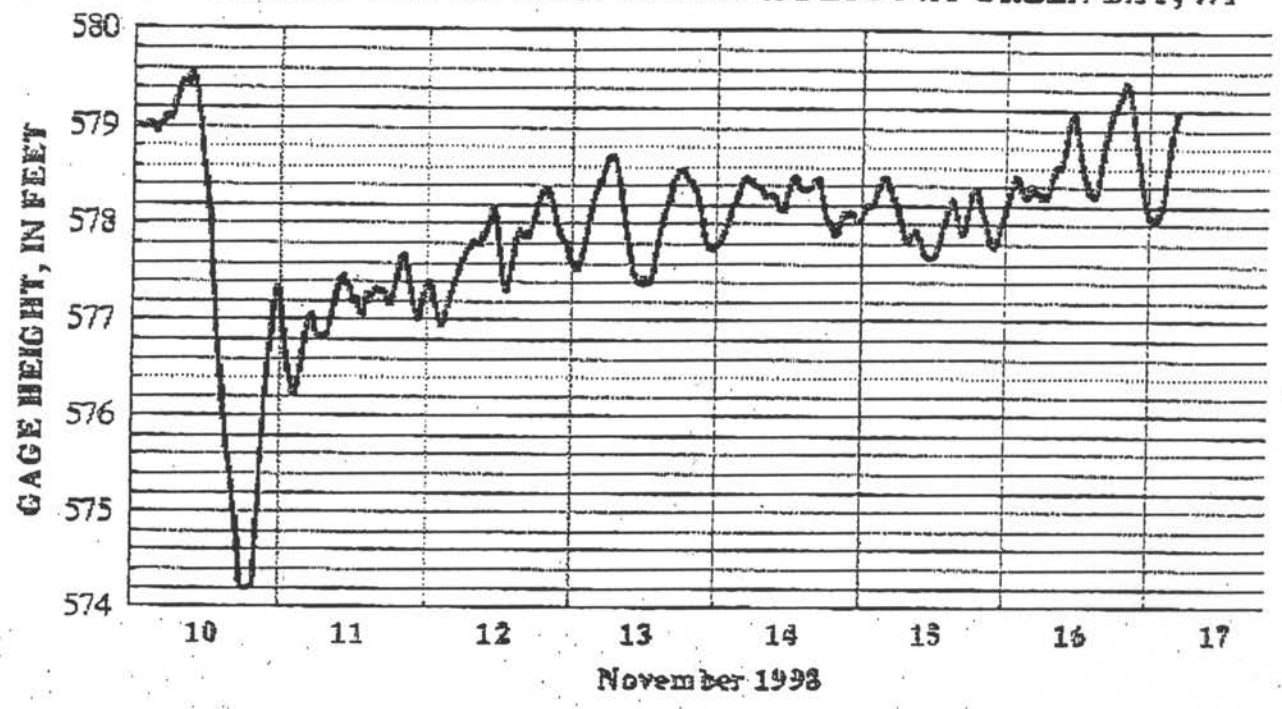
040831335 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI



Updated: 11-17-98 07:52

↑
11/10/98
Storm force winds, southerly;
with strong Low barometric pressure.

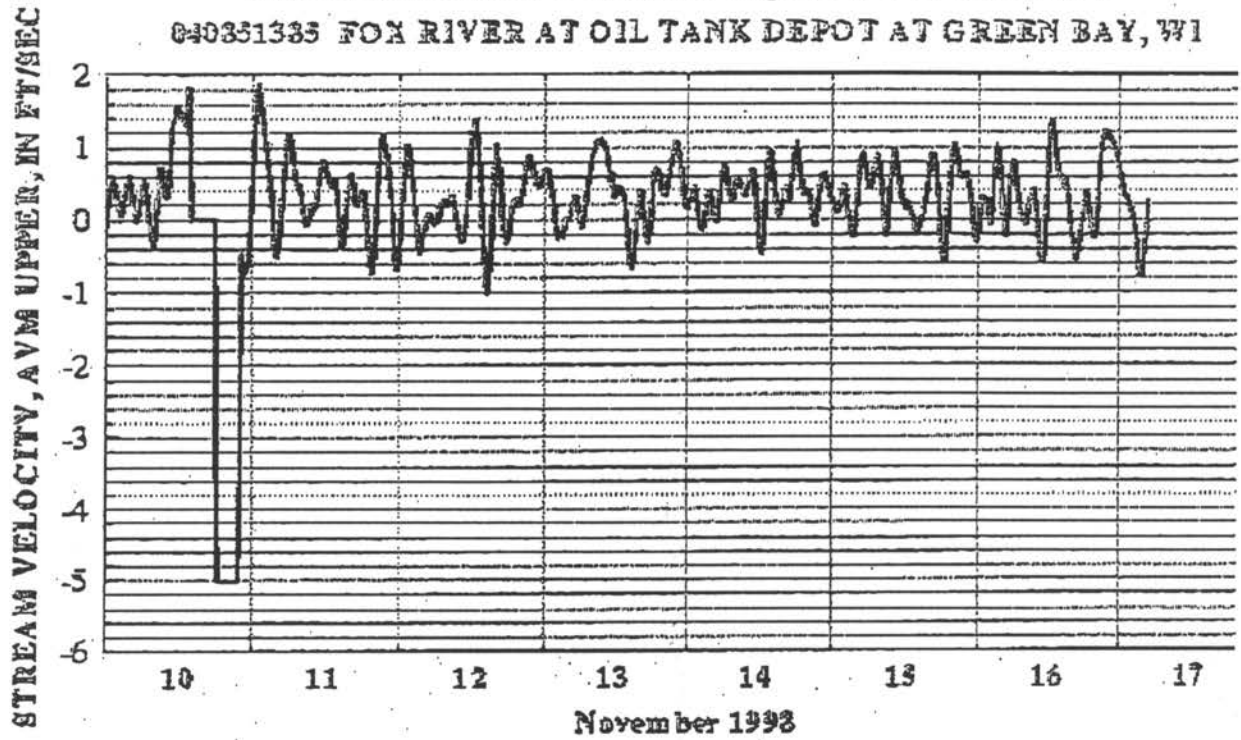
U.S. GEOLOGICAL SURVEY
PROVISIONAL DATA SUBJECT TO REVIEW
040331335 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI



Updated: 11-17-98 07:11

U.S. GEOLOGICAL SURVEY
PROVISIONAL DATA SUBJECT TO REVIEW

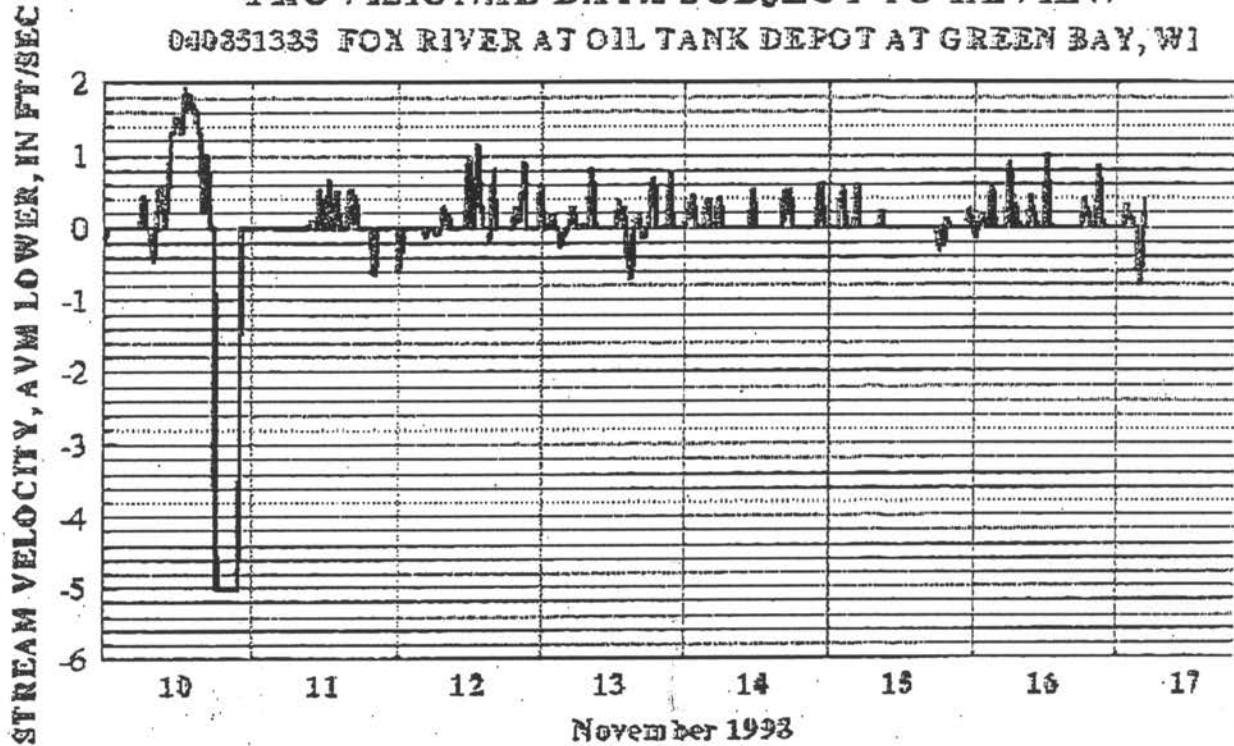
040351385 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI



Updated: 11-17-98 07:53

**U.S. GEOLOGICAL SURVEY
PROVISIONAL DATA SUBJECT TO REVIEW**

040851335 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI



Updated: 11-17-98 07:53

Map of region surrounding station

STATION.-- 040851385 FOX RIVER AT OIL TANK DEPOT AT GREEN BAY, WI

LOCATION.--Lat 44°31'43", long 88°01'12" in section 25, T.24 N., R.20
Brown County, Hydrologic Unit 04030204, about 0.5 mi upstream of I
Highway 43 bridge in Green Bay, and 0.8 mi upstream from mouth.

DRAINAGE AREA.--6,330 square miles.

PERIOD OF RECORD.--October 1988 to current year.

218709

APPENDIX G

PHYSICAL PARAMETERS TABLES

**Appendix G - Table 1
Lower Fox River Grain Size Results**

| Deposit or SMU Group | Grain Size Averages | | | |
|--|---------------------|-------------|-------------|-------------|
| | Gravel (%) | Sand (%) | Silt (%) | Clay (%) |
| Lake Winnebago | 0.0 | 14.0 | 48.7 | 37.3 |
| Little Lake Butte Des Morts Reach | | | | |
| Deposit A | 0.0 | 37.5 | 45.2 | 17.3 |
| Deposit B | 0.0 | 64.7 | 25.1 | 10.1 |
| Deposit C | 0.0 | 26.1 | 53.8 | 20.1 |
| Deposit POG | 2.2 | 57.4 | 34.4 | 6.0 |
| Deposit D | 0.3 | 43.8 | 44.1 | 11.9 |
| Deposit E | 0.1 | 27.7 | 50.5 | 21.8 |
| Deposit F | 0.0 | 27.1 | 50.8 | 22.1 |
| Deposit G | 0.0 | 55.7 | 31.0 | 13.3 |
| Deposit H | 0.0 | 67.7 | 20.3 | 12.0 |
| Interdeposit Areas | 3.2 | 49.3 | 35.6 | 12.0 |
| Reach Average | 0.6 | 45.7 | 39.1 | 14.7 |
| Appleton to Little Rapids Reach | | | | |
| Deposit I | 0.0 | 35.0 | 45.3 | 19.8 |
| Deposit J | 0.0 | 15.0 | 65.7 | 19.3 |
| Deposit K | 0.0 | 62.7 | 22.3 | 15.0 |
| Deposit L | 0.0 | 45.3 | 34.0 | 20.8 |
| Deposit M | 0.0 | 7.3 | 63.3 | 29.3 |
| Deposit N | 0.5 | 41.1 | 46.9 | 11.6 |
| Deposit O | 0.0 | 39.4 | 43.6 | 17.0 |
| Deposit P | 0.0 | 36.0 | 49.6 | 14.4 |
| Deposit Q | 0.0 | 49.0 | 39.7 | 11.3 |
| Deposit R | 0.0 | 12.0 | 56.0 | 32.0 |
| Deposit S | 0.0 | 46.5 | 36.0 | 17.5 |
| Deposit T | 0.0 | 87.7 | 7.3 | 5.0 |
| Deposit U | 0.0 | 51.8 | 35.8 | 12.5 |
| Deposit V | 0.0 | 32.2 | 52.0 | 15.8 |
| Deposit W | 0.0 | 50.1 | 32.5 | 17.4 |
| Deposit X | 0.0 | 33.2 | 52.8 | 14.0 |
| Deposit Y | 0.0 | 45.0 | 39.7 | 15.3 |
| Deposit Z | 0.0 | 34.7 | 42.7 | 22.7 |
| Deposit AA | 0.0 | 54.7 | 20.7 | 24.7 |
| Deposit BB | 0.0 | 47.7 | 33.0 | 19.3 |
| Deposit CC | 0.0 | 31.3 | 26.0 | 42.7 |
| Deposit DD | 0.0 | 32.6 | 42.1 | 25.3 |
| Interdeposit Areas | NA | NA | NA | NA |
| Reach Average | 0.0 | 40.5 | 40.3 | 19.2 |

**Appendix G - Table 1
Lower Fox River Grain Size Results**

| Deposit or SMU Group | Grain Size Averages | | | |
|---|---------------------|-------------|-------------|-------------|
| | Gravel (%) | Sand (%) | Silt (%) | Clay (%) |
| Little Rapids to DePere Reach | | | | |
| Deposit EE | 0.5 | 26.8 | 49.7 | 23.0 |
| Deposit FF | 0.0 | 27.2 | 51.6 | 21.1 |
| Deposit GG | 1.3 | 18.0 | 57.6 | 23.1 |
| Deposit HH | 2.9 | 21.7 | 57.1 | 18.4 |
| Interdeposit Areas | 3.7 | 31.9 | 24.3 | 40.1 |
| Reach Average | 1.2 | 23.4 | 54.0 | 21.4 |
| DePere to Green Bay Reach | | | | |
| SMU 20 to 25 | 0.0 | 42.3 | 42.5 | 15.2 |
| SMU 26 to 31 | 0.0 | 50.8 | 34.5 | 14.7 |
| SMU 32 to 37 | 0.0 | 31.8 | 49.9 | 18.3 |
| SMU 38 to 43 | 0.0 | 34.5 | 47.4 | 18.1 |
| SMU 44 to 49 | 0.0 | 37.8 | 44.6 | 17.6 |
| SMU 50 to 55 | 0.0 | 40.5 | 44.2 | 15.3 |
| SMU 56 to 61 | 0.0 | 32.1 | 51.9 | 16.0 |
| SMU 62 to 67 | 0.0 | 29.8 | 51.7 | 18.6 |
| SMU 68 to 73 | 0.5 | 34.8 | 41.6 | 23.1 |
| SMU 74 to 79 | 0.0 | 34.8 | 42.2 | 23.0 |
| SMU 80 to 85 | 0.0 | 45.4 | 36.8 | 17.8 |
| SMU 86 to 91 | 0.0 | 45.5 | 37.6 | 17.0 |
| SMU 92 to 97 | 0.0 | 60.3 | 27.9 | 11.8 |
| SMU 98 to 103 | 0.0 | 73.2 | 17.8 | 9.0 |
| SMU 104 to 109 | 0.0 | 41.7 | 40.5 | 17.8 |
| SMU 110 to 115 | 0.0 | 44.2 | 38.9 | 16.9 |
| Reach Average | 0.0 | 42.5 | 40.6 | 16.9 |
| Lower Fox River Average Values | 0.5 | 38.0 | 43.5 | 18.0 |

**Appendix G - Table 2
Green Bay Grain Size Results**

| Green Bay Zone | Sample Label | Gravel | Sand | Silt | Clay |
|--------------------------|---------------------|---------------|--------------|--------------|--------------|
| Zone 2 (2A & 2B) | S00030 | 0.0% | 64.8% | 30.7% | 4.5% |
| Zone 2 (2A & 2B) | S00031 | 0.1% | 93.1% | 2.3% | 4.5% |
| Zone 2 (2A & 2B) | S00032 | 0.0% | 98.6% | 0.0% | 1.4% |
| Zone 2 (2A & 2B) | S00037 | 0.0% | 70.4% | 17.6% | 12.0% |
| Zone 2 (2A & 2B) | S00038 | 0.7% | 69.1% | 23.7% | 6.5% |
| Zone 2 (2A & 2B) | S00039 | 0.0% | 65.8% | 20.2% | 14.0% |
| Zone 2 (2A & 2B) | S00040 | 0.0% | 51.6% | 29.4% | 19.0% |
| Zone 2 (2A & 2B) | S00056 | 0.1% | 90.9% | 4.5% | 4.5% |
| Zone 2 (2A & 2B) | S00057 | 0.0% | 61.8% | 32.2% | 6.0% |
| Zone 2 (2A & 2B) | S00058 | 0.1% | 67.8% | 18.1% | 14.0% |
| Zone 2 (2A & 2B) | S00063 | 0.2% | 72.5% | 19.3% | 8.0% |
| Zone 2 Averages | | 0.1% | 73.3% | 18.0% | 8.6% |
| Zone 3A | S00042 | 0.0% | 99.2% | 0.8% | 0.0% |
| Zone 3A | S00043 | 0.0% | 97.6% | 0.7% | 1.7% |
| Zone 3A Averages | | 0.0% | 98.4% | 0.8% | 0.9% |
| Zone 3B | S00041 | 0.0% | 83.3% | 10.2% | 6.5% |
| Zone 3B | S00047 | 0.0% | 73.1% | 20.9% | 6.0% |
| Zone 3B | S00048 | 0.0% | 66.3% | 21.7% | 12.0% |
| Zone 3B | S00054 | 0.2% | 27.9% | 46.9% | 25.0% |
| Zone 3B Averages | | 0.1% | 62.7% | 24.9% | 12.4% |
| Zone 4 | S00044 | 3.0% | 96.1% | 0.9% | 0.0% |
| Zone 4 | S00045 | 0.0% | 92.9% | 5.3% | 1.8% |
| Zone 4 | S00046 | 0.8% | 97.7% | 1.5% | 0.0% |
| Zone 4 | S00055 | 1.6% | 98.4% | 0.0% | 0.0% |
| Zone 4 Averages | | 1.4% | 96.3% | 1.9% | 0.5% |
| Green Bay Average | | 0.3% | 78.0% | 14.6% | 7.0% |

Notes: 1) All samples collected from 0 to 10 cm.

Appendix G - Table 3
Lower Fox River - Atterberg Limits

| Deposit or SMU | Sample Label | Sample Depth (cm) | Liquid Limit | Plastic Limit | Plasticity Index | Plasticity Index |
|--|--------------|-------------------|--------------|---------------|------------------|------------------|
| Little Lake Butte des Morts Reach | | | | | | |
| Deposit A | BA-SD01e | 61 - 79 | 74 | N/A | 49 | na |
| Deposit A | BA-SD02d | 48 - 58 | 50 | N/A | 26 | na |
| Deposit A | BA-SD03comp | 0 - 30 | 148 | N/A | 84 | na |
| Deposit A | BA-SD04c | 30 - 43 | 178 | N/A | 65 | na |
| Deposit A | BA-SD04d | 43 - 53 | 46 | N/A | 26 | na |
| Deposit A | BA-SD08d | 45 - 61 | 35 | N/A | 18 | na |
| Deposit C | SDC-C-1-G | 5 - 35 | 113.4 | 51.4 | 62 | MH |
| Deposit C | SDC-C-4-G | 5 - 35 | 186.3 | 61.3 | 125 | MH |
| Deposit E | SDC-E-2-G | 5 - 35 | 104.7 | 56.8 | 47.9 | MH |
| Deposit E | SDC-E-5b-G | 5 - 35 | N/A | Non-Plastic | N/A | na |
| Deposit F | GT0110 | 0 - 182.9 | 114 | 64.3 | 49.7 | na |
| Appleton to Little Rapids Reach | | | | | | |
| Deposit W | SDC-W-2-G | 5 - 35 | 71.1 | 37.2 | 33.9 | MH |
| Deposit W | SDC-W-5-G | 5 - 35 | 106.9 | 58.9 | 48 | MH |
| Deposit X | GT0143 | 0 - 195.1 | 92.6 | 52.9 | 39.7 | na |
| Deposit X | GT0144 | 0 - 195.1 | 83.3 | 44.9 | 38.4 | na |
| Deposit X | SDC-X-1-G | 5 - 35 | N/A | Non-Plastic | N/A | MH |
| Deposit X | SDC-X-2-G | 5 - 35 | 73.4 | 52.6 | 20.8 | CH |
| Little Rapids to De Pere Reach | | | | | | |
| Deposit EE | GT0125 | 0 - 182.9 | 73.2 | 36.6 | 36.6 | na |
| Deposit EE | SDC-EE22-3-G | 5 - 35 | 61.3 | 30.3 | 31 | CH |
| Deposit EE | SDC-EE22-4-G | 5 - 35 | 85 | 45.1 | 39.9 | MH |
| Deposit EE | SDC-EE23-1-G | 5 - 35 | N/A | Non-Plastic | N/A | na |
| Deposit EE | SDC-EE23-4-G | 5 - 35 | 144 | 45.9 | 98.1 | MH |
| Deposit EE | SDC-EE24-1-G | 5 - 35 | 92.5 | 45.2 | 47.3 | MH |
| Deposit EE | SDC-EE24-3-G | 5 - 32 | 76.6 | 39.7 | 36.9 | MH |
| Deposit EE | SDC-EE25-2-G | 5 - 35 | 93.4 | 50 | 43.4 | MH |
| Deposit EE | SDC-EE25-3-G | 5 - 35 | 176.7 | 113.4 | 63.3 | MH |
| Deposit EE | SDC-EE26-2-G | 5 - 35 | 88.8 | 48.5 | 40.3 | MH |
| Deposit EE | SDC-EE26-5-G | 5 - 35 | 89.5 | 44 | 45.5 | MH |
| Deposit GG | GT0068 | 0 - 182.9 | 89.4 | 45.1 | 44.3 | na |
| Deposit HH | GT0079 | 0 - 182.9 | 85.4 | 44.5 | 40.9 | na |
| De Pere to Green Bay Reach | | | | | | |
| SMU 20 | GT0005 | 0 - 173.7 | 94.3 | 47 | 47.3 | na |
| SMU 20 | SDC-DPD-2-G | 5 - 35 | 95 | 49.5 | 45.5 | MH |
| SMU 24 | GT0013 | 0 - 185.9 | 97.3 | 53 | 44.3 | na |
| SMU 41 | GT0036 | 0 - 182.9 | 37.7 | 21.5 | 16.2 | na |
| SMU 45 | GT0048 | 0 - 195.1 | 68.9 | 33.5 | 35.4 | na |
| SMU 45 | SDC-DPD-3-G | 5 - 35 | 156.9 | 109.6 | 47.3 | MH |
| SMU 48 | GT0040 | 0 - 182.9 | 44.6 | 23.6 | 21 | na |
| SMU 62 | GT0052 | 0 - 213.4 | 89 | 47.6 | 41.4 | na |

Notes:

- Atterberg Limits testing performed according to ASTM D-4318.
- Samples listed as non-plastic could not be cut with the grooving tool without tearing or slipping in the cup. Every effort was made to test these samples, but a liquid limit could not be determined.
- Classifications are based on ASTM D-2487. The samples were visually determined to be organic. Samples classified as "na" were not determined by the laboratory.

Appendix G - Table 4
Lower Fox River & Green Bay Maximum PCB Sampling
Depth and Deposit/SMU Area

| Deposit/SMU Group | Maximum PCB Sampling Depth | | Sediment Area | |
|-------------------------------|----------------------------|-----------|---------------|---------------|
| | (m) | (ft) | (hectares) | (acres) |
| LLBdM | | | | |
| Deposit A | 1.80 | 5.90 | 15.26 | 37.71 |
| Deposit B | 0.43 | 1.41 | 14.74 | 36.42 |
| Deposit C | 0.91 | 2.98 | 12.36 | 30.54 |
| Deposit POG | 1.89 | 6.20 | 21.32 | 52.68 |
| Deposit D | 1.22 | 4.00 | 25.24 | 62.37 |
| Deposit E | 1.74 | 5.71 | 202.51 | 500.41 |
| Deposit F | 1.83 | 6.00 | 16.91 | 41.79 |
| Deposit G | 0.30 | 0.98 | 4.11 | 10.16 |
| Deposit H | 0.38 | 1.25 | 1.08 | 2.67 |
| Reach Totals | na | na | 313.53 | 774.75 |
| Appleton-Little Rapids | | | | |
| Deposit I | 0.54 | 1.77 | 2.98 | 7.36 |
| Deposit J | 0.42 | 1.38 | 2.51 | 6.20 |
| Deposit K | 0.21 | 0.69 | 0.53 | 1.31 |
| Deposit L | 0.30 | 0.98 | 1.06 | 2.62 |
| Deposit M | 0.36 | 1.18 | 1.33 | 3.29 |
| Deposit N | 0.89 | 2.92 | 2.25 | 5.56 |
| Deposit O | 0.35 | 1.15 | 1.85 | 4.57 |
| Deposit P | 0.94 | 3.08 | 3.14 | 7.76 |
| Deposit Q | 0.55 | 1.80 | 0.42 | 1.04 |
| Deposit R | 0.13 | 0.43 | 0.77 | 1.90 |
| Deposit S | 0.34 | 1.12 | 16.64 | 41.12 |
| Deposit T | 0.52 | 1.71 | 2.08 | 5.14 |
| Deposit U | 0.26 | 0.85 | 1.74 | 4.30 |
| Deposit V | 0.63 | 2.07 | 2.41 | 5.96 |
| Deposit W | 1.52 | 4.99 | 56.41 | 139.39 |
| Deposit X | 1.83 | 6.00 | 25.60 | 63.26 |
| Deposit Y | 0.34 | 1.12 | 3.19 | 7.88 |
| Deposit Z | 0.83 | 2.72 | 2.44 | 6.03 |
| Deposit AA | 0.35 | 1.15 | 0.81 | 2.00 |
| Deposit BB | 0.39 | 1.28 | 1.58 | 3.90 |
| Deposit CC | 0.43 | 1.41 | 8.47 | 20.93 |
| Deposit DD | 0.53 | 1.74 | 14.92 | 36.87 |
| Reach Totals | NA | NA | 153.13 | 378.39 |

**Appendix G - Table 4
Lower Fox River & Green Bay Maximum PCB Sampling
Depth and Deposit/SMU Area**

| Deposit/SMU Group | Maximum PCB Sampling Depth | | Sediment Area | |
|---------------------------------|----------------------------|-----------|----------------|------------------|
| | (m) | (ft) | (hectares) | (acres) |
| Little Rapids to De Pere | | | | |
| Deposit EE | 2.30 | 7.54 | 258.81 | 639.53 |
| Deposit FF | 0.46 | 1.51 | 0.49 | 1.21 |
| Deposit GG | 2.30 | 7.54 | 2.40 | 5.93 |
| Deposit HH | 2.30 | 7.54 | 4.46 | 11.02 |
| Reach Totals | NA | NA | 266.16 | 657.69 |
| De Pere to Green Bay | | | | |
| SMU 20-25 | 2.13 | 6.99 | 113.39 | 280.19 |
| SMU 26-31 | 2.13 | 6.99 | 22.04 | 54.46 |
| SMU 32-37 | 2.74 | 8.99 | 26.78 | 66.17 |
| SMU 38-43 | 2.74 | 8.99 | 46.46 | 114.80 |
| SMU 44-49 | 3.35 | 10.99 | 107.15 | 264.77 |
| SMU 50-55 | 1.52 | 4.99 | 32.91 | 81.32 |
| SMU 56-61 | 3.96 | 12.99 | 29.66 | 73.29 |
| SMU 62-67 | 2.13 | 6.99 | 18.22 | 45.02 |
| SMU 68-73 | 2.74 | 8.99 | 21.58 | 53.33 |
| SMU 74-79 | 1.52 | 4.99 | 11.81 | 29.18 |
| SMU 80-85 | 2.13 | 6.99 | 10.62 | 26.24 |
| SMU 86-91 | 2.13 | 6.99 | 11.27 | 27.85 |
| SMU 92-97 | 0.91 | 2.98 | 19.76 | 48.83 |
| SMU 98-103 | 0.91 | 2.98 | 14.00 | 34.59 |
| SMU 104-109 | 0.30 | 0.98 | 17.02 | 42.06 |
| SMU 110-115 | 1.52 | 4.99 | 20.82 | 51.45 |
| Reach Totals | NA | NA | 523.49 | 1293.57 |
| Green Bay Zones | | | | |
| Zone 2 (2A&2B) | 0.91 | 2.98 | 11,081 | 27,382 |
| Zone 3A | 0.30 | 0.98 | 85,891 | 212,240 |
| Zone 3B | 0.62 | 2.03 | 69,339 | 171,340 |
| Zone 4 | 0.30 | 0.98 | 254,977 | 630,059 |
| Bay Totals | NA | NA | 421,288 | 1,041,021 |

na - Total value result not applicable.

Appendix G - Table 5
Lower Fox River and Green Bay TOC and Bulk Density Results

| Sampling Location | Average TOC Values | | Sampling Location | Average TOC Values | |
|--|--------------------|--------------|--------------------------------------|--------------------|--------------|
| | (mg/kg) | Percent | | (mg/kg) | Percent |
| Lake Winnebago | 78,000 | 7.80% | Little Rapids to DePere Reach | | |
| Little Lake Butte des Morts Reach | | | Creek Trib. | 20,300 | 2.03% |
| Creek Trib. | 31,000 | 3.10% | Deposit EE | 55,957 | 5.60% |
| Deposit A | 90,359 | 9.04% | Deposit FF | 49,183 | 4.92% |
| Deposit B | 26,064 | 2.61% | Deposit GG | 59,318 | 5.93% |
| Deposit C | 70,577 | 7.06% | Deposit HH | 64,196 | 6.42% |
| Deposit POG | 79,129 | 7.91% | Interdeposit | 29,333 | 2.93% |
| Deposit D | 54,863 | 5.49% | Reach Average | 49,791 | 4.98% |
| Deposit E | 61,210 | 6.12% | DePere to Green Bay Reach | | |
| Deposit F | 122,917 | 12.29% | Past Mouth | 45,826 | 4.58% |
| Deposit G | 39,633 | 3.96% | SMU 20-25 | 50,855 | 5.09% |
| Deposit H | 37,100 | 3.71% | SMU 26-31 | 36,761 | 3.68% |
| Reach Average | 64,650 | 6.47% | SMU 32-37 | 56,387 | 5.64% |
| Appleton to Little Rapids Reach | | | SMU 38-43 | 45,921 | 4.59% |
| Creek Trib. | 12,600 | 1.26% | SMU 44-49 | 47,306 | 4.73% |
| Deposit I | 43,555 | 4.36% | SMU 50-55 | 37,107 | 3.71% |
| Deposit J | 35,300 | 3.53% | SMU 56-61 | 56,616 | 5.66% |
| Deposit K | 31,567 | 3.16% | SMU 62-67 | 66,420 | 6.64% |
| Deposit L | 24,920 | 2.49% | SMU 68-73 | 50,735 | 5.07% |
| Deposit M | 54,900 | 5.49% | SMU 74-79 | 50,979 | 5.10% |
| Deposit N* | --- | --- | SMU 80-85 | 53,088 | 5.31% |
| Deposit O | 54,917 | 5.49% | SMU 86-91 | 47,022 | 4.70% |
| Deposit P | 43,109 | 4.31% | SMU 92-97 | 27,769 | 2.78% |
| Deposit Q | 73,360 | 7.34% | SMU 98-103 | 20,543 | 2.05% |
| Deposit R | 3,300 | 0.33% | SMU 104-109 | 29,033 | 2.90% |
| Deposit S | 80,300 | 8.03% | SMU 110-115 | 46,474 | 4.65% |
| Deposit T | 86,000 | 8.60% | Reach Average | 45,188 | 4.52% |
| Deposit U | 45,033 | 4.50% | Entire River Average | 49,378 | 4.94% |
| Deposit V | 52,767 | 5.28% | Green Bay | | |
| Deposit W | 38,005 | 3.80% | Zone 2 (2A & 2B) | 14,845 | 1.48% |
| Deposit X | 51,962 | 5.20% | Zone 3A | 1,900 | 0.19% |
| Deposit Y | 0 | 0.00% | Zone 3B | 23,325 | 2.33% |
| Deposit Z | 0 | 0.00% | Zone 4 | 1,400 | 0.14% |
| Deposit AA | 0 | 0.00% | Lake Michigan | 3,461 | 0.35% |
| Deposit BB | 16,100 | 1.61% | Other | 83,600 | 8.36% |
| Deposit CC | 21,486 | 2.15% | USGS Reference | 56,800 | 5.68% |
| Deposit DD | 38,924 | 3.89% | | | |
| Interdeposit | 25,000 | 2.50% | | | |
| Reach Average | 37,881 | 3.79% | | | |

1) Reach and entire river averages do not include tributary results.

na - Parameter result not available.

* Data for Depsoit N is not included due to completion of the remediation demonstration project.

Appendix G - Table 6
Lower Fox River - Total Solids

| Deposit/Interval | Average | Minimum | Maximum |
|--|----------------|----------------|----------------|
| Lake Winnebago | 14.07% | 12.80% | 15.70% |
| Little Lake Butte Des Morts Reach | | | |
| Deposit A | 36.80% | 36.80% | 82.50% |
| Deposit B | 57.21% | 26.80% | 82.40% |
| Deposit C | 31.60% | 15.80% | 76.30% |
| Deposit POG | 32.09% | 18.10% | 67.00% |
| Deposit D | 41.95% | 21.50% | 73.80% |
| Deposit E | 37.78% | 12.60% | 71.30% |
| Deposit F | 26.11% | 17.10% | 38.50% |
| Deposit G | 47.40% | 40.40% | 56.80% |
| Deposit H | 57.93% | 54.70% | 61.00% |
| Interdeposit | 42.60% | 19.20% | 66.00% |
| Entire Reach | 41.15% | 12.60% | 82.50% |
| Appleton to Little Rapids Reach | | | |
| Deposit I | 50.18% | 35.60% | 81.40% |
| Deposit J | 46.13% | 43.20% | 51.20% |
| Deposit K | 51.00% | 39.00% | 61.70% |
| Deposit L | 59.18% | 47.70% | 87.20% |
| Deposit M | 35.73% | 33.70% | 37.60% |
| Deposit O | 41.34% | 33.30% | 56.80% |
| Deposit P | 42.21% | 20.80% | 72.70% |
| Deposit Q | 37.49% | 31.40% | 42.10% |
| Deposit R | 61.10% | 61.10% | 61.10% |
| Deposit S | 47.60% | 31.80% | 63.40% |
| Deposit T | 34.93% | 22.90% | 49.60% |
| Deposit U | 47.78% | 35.50% | 79.80% |
| Deposit V | 32.53% | 26.50% | 43.10% |
| Deposit W | 44.05% | 19.70% | 75.10% |
| Deposit X | 37.49% | 21.00% | 70.30% |
| Deposit Y | 46.63% | 36.80% | 52.50% |
| Deposit Z | 51.50% | 46.30% | 54.80% |
| Deposit AA | 67.70% | 62.60% | 72.20% |
| Deposit BB | 57.80% | 43.50% | 67.20% |
| Deposit CC | 57.08% | 35.70% | 69.60% |
| Deposit DD | 44.61% | 19.90% | 75.50% |
| Interdeposit | 26.00% | 26.00% | 26.00% |
| Entire Reach | 46.37% | 19.70% | 87.20% |

Appendix G - Table 6
Lower Fox River - Total Solids, continued

| Deposit/Interval | Average | Minimum | Maximum |
|--------------------------------------|----------------|----------------|----------------|
| Little Rapids to DePere Reach | | | |
| Deposit EE | 37.07% | 16.70% | 88.20% |
| Deposit EG | 42.60% | 26.00% | 69.70% |
| Deposit FF | 45.14% | 21.90% | 86.10% |
| Deposit GG | 36.36% | 25.70% | 85.90% |
| Deposit HH | 36.70% | 21.10% | 85.80% |
| Interdeposit | 36.03% | 19.80% | 76.30% |
| Entire Reach | 38.98% | 16.70% | 88.20% |
| DePere to Green Bay Reach | | | |
| Less than 61 cm deep | 35.70% | 12.70% | 80.20% |
| 62 cm - 240 cm | 45.60% | 22.10% | 81.70% |
| Entire Reach (all depths) | 41.20% | 12.70% | 83.90% |
| Entire River | | | |
| Entire River | 44.40% | 12.70% | 88.20% |
| Green Bay | | | |
| Zones 2A/2B | 49.52% | 30.10% | 73.80% |
| Zone 3A | 28.45% | 2.60% | 72.00% |
| Zone 3B | 28.37% | 15.10% | 59.20% |
| Zone 4 | 72.58% | 68.60% | 77.60% |
| Entire Bay | 44.73% | 2.60% | 77.60% |

Notes: 1) All samples collected above 61 cm (2 ft) except in the De Pere to Green Bay Reach. These sample results were delineated to evaluate the solids content in the upper 61 cm of sediment.

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-----------------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| Lake Winnebago | | | | | |
| | SDC-LW-1-P-S | 0 - 5 | NA | 0.17 | NA |
| | SDC-LW-2-P-S | 0 - 5 | NA | 0.14 | NA |
| | SDC-LW-3-P-S | 0 - 5 | NA | 0.15 | NA |
| Reach Average | | | --- | 0.15 | --- |
| LLBdM Reach | | | | | |
| A | 2A1.1 | 0 - 5 | NA | 0.49 | NA |
| A | 2A1.2 | 5 - 15 | NA | 0.47 | NA |
| A | 2A1.3 | 15 - 25 | NA | 0.40 | NA |
| A | 2A1.4 | 25 - 37 | NA | 0.52 | NA |
| A | 2A10.1 | 0 - 5 | NA | 0.35 | NA |
| A | 2A10.2 | 5 - 15 | NA | 0.37 | NA |
| A | 2A10.3 | 15 - 25 | NA | 0.80 | NA |
| A | 2A2.2 | 5 - 15 | NA | 0.55 | NA |
| A | 2A2.3 | 15 - 25 | NA | 0.54 | NA |
| A | 2A3.3 | 15 - 25 | NA | 0.42 | NA |
| A | 2A3.4 | 25 - 35 | NA | 0.54 | NA |
| A | 2A4.1 | 0 - 5 | NA | 0.26 | NA |
| A | 2A4.2 | 5 - 15 | NA | 0.42 | NA |
| A | 2A5.1 | 0 - 5 | NA | 0.49 | NA |
| A | 2A5.2 | 5 - 15 | NA | 0.45 | NA |
| A | 2A5.3 | 15 - 25 | NA | 0.39 | NA |
| A | 2A5.4 | 25 - 35 | NA | 0.33 | NA |
| A | 2A6.1 | 0 - 5 | NA | 0.31 | NA |
| A | 2A6.2 | 5 - 15 | NA | 0.44 | NA |
| A | 2A6.3 | 15 - 25 | NA | 0.42 | NA |
| A | 2A6.4 | 25 - 35 | NA | 0.44 | NA |
| A | 2A6.5 | 35 - 45 | NA | 0.34 | NA |
| A | 2A6.6 | 45 - 55 | NA | 0.35 | NA |
| A | 2A6.7 | 55 - 65 | NA | 0.37 | NA |
| A | 2A7.1 | 0 - 5 | NA | 0.51 | NA |
| A | 2A7.2 | 5 - 15 | NA | 0.54 | NA |
| A | 2A7.4 | 25 - 35 | NA | 0.57 | NA |
| A | 2A7.5 | 35 - 45 | NA | 1.50 | NA |
| A | 2A7.6 | 45 - 51 | NA | 1.71 | NA |
| A | 2A8.1 | 0 - 5 | NA | 0.54 | NA |
| A | 2A8.2 | 5 - 15 | NA | 0.57 | NA |
| A | 2A8.3 | 15 - 25 | NA | 0.60 | NA |
| A | 2A8.4 | 25 - 35 | NA | 0.99 | NA |
| A | 2A9.1 | 0 - 5 | NA | 0.40 | NA |
| A | 2A9.2 | 5 - 15 | NA | 0.38 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| A | 2A9.3 | 15 - 25 | NA | 0.44 | NA |
| A | 2A9.5 | 35 - 45 | NA | 0.42 | NA |
| A | 2A9.6 | 45 - 61 | NA | 0.41 | NA |
| A | 2NS1.2 OF 2 | - | NA | 0.98 | NA |
| A | 2NS3.1 OF 2 | - | NA | 0.82 | NA |
| A | 3A1.1 | 0 - 5 | NA | 0.35 | NA |
| A | 3A1.2 | 0 - 10 | NA | 0.39 | NA |
| A | 3A1.3 | 10 - 20 | NA | 0.44 | NA |
| A | 3A1.4 | 20 - 30 | NA | 0.40 | NA |
| A | 3A1.5 | 30 - 42 | NA | 0.33 | NA |
| A | 3A2.1 | 0 - 5 | NA | 0.40 | NA |
| A | 3A2.2 | 5 - 15 | NA | 0.55 | NA |
| A | 3A2.3 | 15 - 25 | NA | 0.42 | NA |
| A | 3A2.4 | 25 - 38 | NA | 0.41 | NA |
| A | 3A21.1 | 0 - 5 | NA | 0.41 | NA |
| A | 3A21.2 | 5 - 15 | NA | 0.48 | NA |
| A | 3A21.3 | 15 - 25 | NA | 1.18 | NA |
| A | 3NS1.1 | 0 - 5 | NA | 0.52 | NA |
| A | 3NS2.1 | 0 - 5 | NA | 1.20 | NA |
| A | 3NS2.2 | 5 - 15 | NA | 1.09 | NA |
| A | 3NS2.3 | 15 - 25 | NA | 0.51 | NA |
| A | 3NS2.4 | 25 - 35 | NA | 0.36 | NA |
| A | 3NS4.1 | 0 - 5 | NA | 0.61 | NA |
| A | 3NS4.2 | 5 - 15 | NA | 0.85 | NA |
| A | 3NS4.3 | 15 - 22 | NA | 0.69 | NA |
| A | 4A1.1 | 0 - 2 | NA | 0.40 | NA |
| A | 4A1.10 | 18 - 20 | NA | 0.25 | NA |
| A | 4A1.11 | 20 - 22 | NA | 0.29 | NA |
| A | 4A1.12 | 22 - 24 | NA | 0.25 | NA |
| A | 4A1.13 | 24 - 26 | NA | 0.24 | NA |
| A | 4A1.14 | 26 - 28 | NA | 0.27 | NA |
| A | 4A1.15 | 28 - 30 | NA | 0.21 | NA |
| A | 4A1.2 | 2 - 4 | NA | 0.44 | NA |
| A | 4A1.3 | 4 - 6 | NA | 0.43 | NA |
| A | 4A1.4 | 6 - 8 | NA | 0.38 | NA |
| A | 4A1.5 | 8 - 10 | NA | 0.35 | NA |
| A | 4A1.6 | 10 - 12 | NA | 0.37 | NA |
| A | 4A1.7 | 12 - 14 | NA | 0.32 | NA |
| A | 4A1.8 | 14 - 16 | NA | 0.31 | NA |
| A | 4A1.9 | 16 - 18 | NA | 0.27 | NA |
| A | A1.1 | 0 - 6 | NA | 0.44 | NA |
| A | A1.2 | 6 - 30 | NA | 0.38 | NA |
| A | A1.3 | 30 - 55 | NA | 0.74 | NA |
| A | A2.1 | 0 - 20 | NA | 0.46 | NA |
| A | A2.2 | 20 - 40 | NA | 0.80 | NA |
| A | A2.3 | 40 - 60 | NA | 0.35 | NA |
| A | A3.1 | 0 - 25 | NA | 0.48 | NA |
| A | A3.2 | 25 - 55 | NA | 0.49 | NA |
| A | A4.1 | 0 - 24 | NA | 0.55 | NA |
| A | A4.2 | 24 - 48 | NA | 0.94 | NA |
| A | A5.1 | 0 - 30 | NA | 0.42 | NA |
| A | A5.2 | 30 - 52 | NA | 0.38 | NA |
| A | A5.3 | 52 - 74 | NA | 0.49 | NA |
| A | AC1c1 | 0 - 10 | NA | 0.39 | NA |
| A | AC1c2 | 10 - 25 | NA | 0.36 | NA |
| A | AC1c3 | 25 - 36 | NA | 1.36 | NA |
| A | AC2c1 | 0 - 27 | NA | 0.45 | NA |
| A | AC2c2 | 27 - 54 | NA | 0.47 | NA |
| A | AC2c3 | 54 - 60 | NA | 0.32 | NA |
| A | BA-SD01e | 61 - 79 | NA | 1.82 | NA |
| A | BA-SD02d | 48 - 58 | NA | 1.61 | NA |
| A | BA-SD03comp | 0 - 30 | NA | 1.08 | NA |
| A | BA-SD04c | 30 - 43 | NA | 1.16 | NA |
| A | BA-SD04d | 43 - 53 | NA | 1.80 | NA |
| A | BA-SD07b | 15 - 30 | NA | 1.62 | NA |
| A | BA-SD08d | 45 - 61 | NA | 1.57 | NA |
| A | POG2 | 0 - 5 | NA | 1.46 | NA |
| B | 2B2.1 | 0 - 5 | NA | 0.90 | NA |
| B | 2B2.2 | 5 - 13 | NA | 1.61 | NA |
| B | B1.1 | 0 - 6 | NA | 0.32 | NA |
| B | B1.2 | 6 - 17 | NA | 0.40 | NA |
| B | B1.3 | 17 - 20 | NA | 0.71 | NA |
| B | B2.1 | 0 - 19 | NA | 1.34 | NA |
| B | POG1 | 0 - 5 | NA | 1.70 | NA |
| C | 2C1.1 | 0 - 5 | NA | 0.26 | NA |
| C | 2C1.2 | 5 - 15 | NA | 0.23 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| C | 2C1.4 | 25 - 41 | NA | 0.36 | NA |
| C | 2C2.1 | 0 - 5 | NA | 0.26 | NA |
| C | 2C2.2 | 5 - 15 | NA | 0.26 | NA |
| C | 2C2.4 | 25 - 36 | NA | 0.41 | NA |
| C | 2C3.1 | 0 - 5 | NA | 0.42 | NA |
| C | 2C3.2 | 5 - 15 | NA | 0.32 | NA |
| C | 2C3.3 | 15 - 26 | NA | 1.46 | NA |
| C | 3C1.1 | 0 - 5 | NA | 0.50 | NA |
| C | 3C1.2 | 5 - 15 | NA | 0.52 | NA |
| C | 3C1.3 | 15 - 25 | NA | 0.65 | NA |
| C | 3C1.4 | 25 - 35 | NA | 0.39 | NA |
| C | 3C1.5 | 35 - 45 | NA | 0.42 | NA |
| C | 3C2.1 | 0 - 5 | NA | 0.34 | NA |
| C | 3C2.2 | 5 - 15 | NA | 0.31 | NA |
| C | 3C2.3 | 15 - 25 | NA | 0.26 | NA |
| C | 3C2.4 | 25 - 35 | NA | 0.37 | NA |
| C | 3C2.5 | 35 - 45 | NA | 0.32 | NA |
| C | 3C3.1 | 0 - 5 | NA | 0.51 | NA |
| C | 3C3.2 | 5 - 15 | NA | 0.44 | NA |
| C | 3C3.3 | 15 - 25 | NA | 0.35 | NA |
| C | 3C3.5 | 35 - 45 | NA | 0.63 | NA |
| C | 3C3.6 | 45 - 53 | NA | 0.52 | NA |
| C | 3C4.1 | 0 - 5 | NA | 0.43 | NA |
| C | 3C4.2 | 5 - 15 | NA | 0.38 | NA |
| C | 3C4.3 | 15 - 25 | NA | 0.37 | NA |
| C | 3C4.4 | 25 - 35 | NA | 0.44 | NA |
| C | C1.1 | 0 - 10 | NA | 0.42 | NA |
| C | C1.2 | 10 - 22 | NA | 0.55 | NA |
| C | C1.3 | 22 - 38 | NA | 0.94 | NA |
| C | SDC-C-1-P-S | 0 - 5 | NA | 0.21 | NA |
| C | SDC-C-2-P-S | 0 - 5 | NA | 0.25 | NA |
| C | SDC-C-3-P-S | 0 - 5 | NA | 0.33 | NA |
| C | SDC-C-4-P-S | 0 - 5 | NA | 0.22 | NA |
| C | SDC-C-5-P-S | 0 - 5 | NA | 0.24 | NA |
| C | SDC-C-1-G | 5-35 | 1.17 | 0.34 | 2.69 |
| C | SDC-C-4-G | 5-35 | 1.13 | 0.34 | 2.48 |
| POG | 2POG1 | 0 - 10 | NA | 0.63 | NA |
| POG | POG3 | 0 - 5 | NA | 0.65 | NA |
| POG | P-RI-1(0-2) | 0 - 61 | NA | 0.67 | NA |
| POG | P-RI-10(0-0.5) | 0 - 15 | NA | 0.52 | NA |
| POG | P-RI-11(0-2) | 0 - 61 | NA | 0.28 | NA |
| POG | P-RI-11(2-4) | 61 - 122 | NA | 0.32 | NA |
| POG | P-RI-11(4-6.2) | 122 - 189 | NA | 0.36 | NA |
| POG | P-RI-12(0-1.4) | 0 - 43 | NA | 0.33 | NA |
| POG | P-RI-13(0-1.1) | 0 - 34 | NA | 0.43 | NA |
| POG | P-RI-14(0-1.2) | 0 - 37 | NA | 0.40 | NA |
| POG | P-RI-15(0-2) | 0 - 61 | NA | 0.28 | NA |
| POG | P-RI-15(2-4) | 61 - 122 | NA | 0.32 | NA |
| POG | P-RI-15(2-4)-FD | 61 - 122 | NA | 0.33 | NA |
| POG | P-RI-15(4-6) | 122 - 183 | NA | 0.42 | NA |
| POG | P-RI-16(0-1.3) | 0 - 40 | NA | 0.28 | NA |
| POG | P-RI-17(0-1.2) | 0 - 37 | NA | 0.38 | NA |
| POG | P-RI-18(0-1.4) | 0 - 43 | NA | 0.27 | NA |
| POG | P-RI-19(0-0.5) | 0 - 15 | NA | 0.98 | NA |
| POG | P-RI-2(0-1) | 0 - 30 | NA | 0.72 | NA |
| POG | P-RI-20(0-2) | 0 - 61 | NA | 0.22 | NA |
| POG | P-RI-20(2-4.3) | 61 - 131 | NA | 0.37 | NA |
| POG | P-RI-21(0-1.8) | 0 - 55 | NA | 0.30 | NA |
| POG | P-RI-22(0-0.4) | 0 - 12 | NA | 0.62 | NA |
| POG | P-RI-3(0-1.0) | 0 - 30 | NA | 0.29 | NA |
| POG | P-RI-4(0-2) | 0 - 61 | NA | 0.24 | NA |
| POG | P-RI-4(2-3.4) | 61 - 104 | NA | 0.32 | NA |
| POG | P-RI-5(0-0.9) | 0 - 27 | NA | 0.44 | NA |
| POG | P-RI-6(0-2.2) | 0 - 67 | NA | 0.66 | NA |
| POG | P-RI-7(0-2) | 0 - 61 | NA | 0.20 | NA |
| POG | P-RI-7(2-2.7) | 61 - 82 | NA | 0.26 | NA |
| POG | P-RI-8(0-1.7) | 0 - 52 | NA | 0.24 | NA |
| POG | P-RI-Comp1(0-2) | 0 - 61 | NA | 0.31 | NA |
| POG | P-RI-Comp1(2-4) | 61 - 122 | NA | 0.34 | NA |
| POG | P-RI-Comp1(4-6) | 122 - 183 | NA | 0.40 | NA |
| D | 2D1.1 | 0 - 5 | NA | 0.71 | NA |
| D | 2D1.2 | 5 - 20 | NA | 0.90 | NA |
| D | 2D2.1 | 0 - 5 | NA | 0.42 | NA |
| D | 2D2.2 | 5 - 15 | NA | 0.59 | NA |
| D | 2D2.3 | 15 - 25 | NA | 0.62 | NA |
| D | 2D2.4 | 25 - 44 | NA | 1.16 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| D | 2D3.1 | 0 - 5 | NA | 0.50 | NA |
| D | 2D3.2 | 5 - 15 | NA | 0.51 | NA |
| D | 2D3.3 | 15 - 25 | NA | 0.52 | NA |
| D | 2D3.5 | 35 - 45 | NA | 0.36 | NA |
| D | 2D4.1 | 0 - 5 | NA | 0.42 | NA |
| D | 2D4.2 | 5 - 15 | NA | 0.54 | NA |
| D | 2D4.3 | 15 - 25 | NA | 0.34 | NA |
| D | 2D4.4 | 25 - 35 | NA | 0.70 | NA |
| D | 3D1.2 | 5 - 15 | NA | 0.42 | NA |
| D | 3D1.4 | 25 - 35 | NA | 0.35 | NA |
| D | 3D1.5 | 35 - 48 | NA | 0.41 | NA |
| D | 3D2.1 | 0 - 5 | NA | 0.37 | NA |
| D | 3D2.3 | 15 - 25 | NA | 0.29 | NA |
| D | 3D3.1 | 0 - 5 | NA | 0.37 | NA |
| D | 3D3.2 | 5 - 15 | NA | 0.70 | NA |
| D | D2.2 | 15 - 39 | NA | 0.40 | NA |
| D | D-RI-1(0-0.5) | 0 - 15 | NA | 0.53 | NA |
| D | D-RI-10(0-2.2) | 0 - 67 | NA | 0.31 | NA |
| D | D-RI-11(0-1.3) | 0 - 40 | NA | 1.07 | NA |
| D | D-RI-12(0-2) | 0 - 61 | NA | 0.42 | NA |
| D | D-RI-12(2-3.5) | 61 - 107 | NA | 0.98 | NA |
| D | D-RI-13(0-2) | 0 - 61 | NA | 0.40 | NA |
| D | D-RI-13(2-3.6) | 61 - 110 | NA | 1.20 | NA |
| D | D-RI-14(0-0.75) | 0 - 23 | NA | 0.54 | NA |
| D | D-RI-15(0-2) | 0 - 61 | NA | 0.77 | NA |
| D | D-RI-15(2-3.7) | 61 - 113 | NA | 0.99 | NA |
| D | D-RI-16(0-1.6) | 0 - 49 | NA | 0.66 | NA |
| D | D-RI-17(0-1.1) | 0 - 34 | NA | 0.29 | NA |
| D | D-RI-18(0-1.5) | 0 - 46 | NA | 0.39 | NA |
| D | D-RI-19(0-0.5) | 0 - 15 | NA | 0.52 | NA |
| D | D-RI-2(0-0.5) | 0 - 15 | NA | 0.83 | NA |
| D | D-RI-20(0-2) | 0 - 61 | NA | 0.87 | NA |
| D | D-RI-20(2-3) | 61 - 91 | NA | 1.37 | NA |
| D | D-RI-21(0-2) | 0 - 61 | NA | 0.88 | NA |
| D | D-RI-21(2-4) | 61 - 122 | NA | 1.33 | NA |
| D | D-RI-3(0-0.5) | 0 - 15 | NA | 0.52 | NA |
| D | D-RI-4(0-0.5) | 0 - 15 | NA | 0.47 | NA |
| D | D-RI-5(0-0.5) | 0 - 15 | NA | 0.40 | NA |
| D | D-RI-6(0-0.5) | 0 - 15 | NA | 0.51 | NA |
| D | D-RI-7(0-1.3) | 0 - 40 | NA | 0.56 | NA |
| D | D-RI-8(0-1.7) | 0 - 52 | NA | 0.59 | NA |
| D | D-RI-9(0-2) | 0 - 61 | NA | 0.49 | NA |
| D | D-RI-9(2-2.8) | 61 - 85 | NA | 0.98 | NA |
| D | D-RI-Comp1(0-2) | 0 - 61 | NA | 0.37 | NA |
| D | D-RI-Comp1(2-4) | 61 - 122 | NA | 1.27 | NA |
| D | D-RI-Comp2(0-2) | 0 - 61 | NA | 0.34 | NA |
| E | 2E1.1 | 0 - 5 | NA | 0.52 | NA |
| E | 2E1.2 | 5 - 23 | NA | 0.96 | NA |
| E | 2E10.1 | 0 - 5 | NA | 0.33 | NA |
| E | 2E10.2 | 5 - 15 | NA | 0.36 | NA |
| E | 2E11.1 | 0 - 5 | NA | 0.37 | NA |
| E | 2E11.2 | 5 - 15 | NA | 0.27 | NA |
| E | 2E11.3 | 15 - 25 | NA | 0.37 | NA |
| E | 2E12.1 | 0 - 5 | NA | 0.53 | NA |
| E | 2E12.2 | 5 - 15 | NA | 0.69 | NA |
| E | 2E12.3 | 15 - 30 | NA | 0.76 | NA |
| E | 2E13.2 | 5 - 19 | NA | 1.21 | NA |
| E | 2E14.1 | 0 - 5 | NA | 0.40 | NA |
| E | 2E14.2 | 5 - 15 | NA | 0.43 | NA |
| E | 2E14.3 | 15 - 25 | NA | 0.38 | NA |
| E | 2E15.1 | 0 - 5 | NA | 0.18 | NA |
| E | 2E15.2 | 5 - 15 | NA | 0.26 | NA |
| E | 2E15.3 | 15 - 30 | NA | 0.32 | NA |
| E | 2E16.1 | 0 - 5 | NA | 0.45 | NA |
| E | 2E16.2 | 5 - 15 | NA | 0.38 | NA |
| E | 2E16.4 | 25 - 32 | NA | 1.05 | NA |
| E | 2E17.1 | 0 - 5 | NA | 0.37 | NA |
| E | 2E17.3 | 15 - 29 | NA | 0.34 | NA |
| E | 2E18.1 | 0 - 5 | NA | 0.28 | NA |
| E | 2E18.3 | 15 - 29 | NA | 0.27 | NA |
| E | 2E19.3 | 15 - 25 | NA | 0.48 | NA |
| E | 2E19.4 | 25 - 37 | NA | 1.08 | NA |
| E | 2E2.1 | 0 - 5 | NA | 0.32 | NA |
| E | 2E2.2 | 5 - 15 | NA | 0.30 | NA |
| E | 2E2.3 | 15 - 25 | NA | 0.37 | NA |
| E | 2E20.2 | 5 - 15 | NA | 0.63 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| E | 2E20.3 | 15 - 25 | NA | 0.50 | NA |
| E | 2E20.4 | 25 - 35 | NA | 0.39 | NA |
| E | 2E21.3 | 15 - 25 | NA | 0.46 | NA |
| E | 2E24.1 | 0 - 5 | NA | 0.22 | NA |
| E | 2E24.2 | 5 - 15 | NA | 0.30 | NA |
| E | 2E24.4 | 25 - 35 | NA | 0.32 | NA |
| E | 2E3.1 | 0 - 5 | NA | 0.33 | NA |
| E | 2E3.2 | 5 - 15 | NA | 0.35 | NA |
| E | 2E3.3 | 15 - 25 | NA | 0.37 | NA |
| E | 2E4.2 | 5 - 15 | NA | 0.43 | NA |
| E | 2E4.3 | 15 - 25 | NA | 0.26 | NA |
| E | 2E5.1 | 0 - 5 | NA | 0.38 | NA |
| E | 2E5.2 | 5 - 15 | NA | 0.34 | NA |
| E | 2E6.1 | 0 - 5 | NA | 0.34 | NA |
| E | 2E6.2 | 5 - 15 | NA | 0.31 | NA |
| E | 2E6.3 | 15 - 24 | NA | 0.32 | NA |
| E | 2E7.1 | 0 - 5 | NA | 0.33 | NA |
| E | 2E7.2 | 5 - 15 | NA | 0.27 | NA |
| E | 2E7.3 | 15 - 25 | NA | 0.30 | NA |
| E | 2E8.1 | 0 - 5 | NA | 0.26 | NA |
| E | 2E8.2 | 5 - 15 | NA | 0.37 | NA |
| E | 2E8.3 | 15 - 25 | NA | 0.38 | NA |
| E | 2E8.4 | 25 - 35 | NA | 0.49 | NA |
| E | 2E9.2 | 5 - 15 | NA | 0.39 | NA |
| E | 2E9.3 | 15 - 25 | NA | 0.28 | NA |
| E | 2POG3 | 0 - 10 | NA | 0.54 | NA |
| E | E1-1.1 | 0 - 18 | NA | 0.41 | NA |
| E | E1-1.2 | 18 - 43 | NA | 0.44 | NA |
| E | E1-1.3 | 43 - 18 | NA | 0.43 | NA |
| E | E1C1C1 | 0 - 7 | NA | 0.19 | NA |
| E | E1C1C2 | 7 - 18 | NA | 0.28 | NA |
| E | E1C1C3 | 18 - 30 | NA | 0.47 | NA |
| E | E2-1.1 | 0 - 13 | NA | 0.29 | NA |
| E | E2-1.2 | 13 - 27 | NA | 0.49 | NA |
| E | E2-1.3 | 27 - 33 | NA | 0.43 | NA |
| E | E2-2.1 | 0 - 6 | NA | 0.69 | NA |
| E | E2-2.2 | 6 - 13 | NA | 0.67 | NA |
| E | E2-2.3 | 13 - 21 | NA | 0.73 | NA |
| E | E2-3.1 | 0 - 17 | NA | 0.46 | NA |
| E | E2-3.2 | 17 - 36 | NA | 0.44 | NA |
| E | E2-3.3 | 36 - 39 | NA | 0.44 | NA |
| E | E2-4.1 | 0 - 7 | NA | 0.91 | NA |
| E | E2-4.2 | 7 - 18 | NA | 0.71 | NA |
| E | E2-4.3 | 18 - 30 | NA | 0.60 | NA |
| E | E2C1C1 | - | NA | 0.41 | NA |
| E | E2C1C2 | - | NA | 0.62 | NA |
| E | E2C1C3 | - | NA | 0.53 | NA |
| E | E-RI-1(0-0.5) | 0 - 15 | NA | 0.94 | NA |
| E | E-RI-10(0-1.5) | 0 - 46 | NA | 1.28 | NA |
| E | E-RI-11(0-2) | 0 - 61 | NA | 0.94 | NA |
| E | E-RI-11(2-3.6) | 61 - 110 | NA | 1.14 | NA |
| E | E-RI-12(0-2) | 0 - 61 | NA | 0.26 | NA |
| E | E-RI-12(2-4.2) | 61 - 128 | NA | 0.81 | NA |
| E | E-RI-13(0-2) | 0 - 61 | NA | 0.24 | NA |
| E | E-RI-13(2-3.75) | 61 - 114 | NA | 0.94 | NA |
| E | E-RI-14(0-2) | 0 - 61 | NA | 0.55 | NA |
| E | E-RI-15(0-2) | 0 - 61 | NA | 0.53 | NA |
| E | E-RI-16(0-2) | 0 - 61 | NA | 0.25 | NA |
| E | E-RI-16(2-3) | 61 - 91 | NA | 0.33 | NA |
| E | E-RI-17(0-2) | 0 - 61 | NA | 0.27 | NA |
| E | E-RI-17(2-4) | 61 - 122 | NA | 0.31 | NA |
| E | E-RI-2(0-2) | 0 - 61 | NA | 0.48 | NA |
| E | E-RI-2(2-4) | 61 - 122 | NA | 1.26 | NA |
| E | E-RI-2(4-4.7) | 122 - 143 | NA | 1.29 | NA |
| E | E-RI-3(0-2) | 0 - 61 | NA | 0.75 | NA |
| E | E-RI-3(2-2.8) | 61 - 85 | NA | 0.93 | NA |
| E | E-RI-4(0-2) | 0 - 61 | NA | 0.75 | NA |
| E | E-RI-4(2-3) | 61 - 91 | NA | 0.82 | NA |
| E | E-RI-5(0-2) | 0 - 61 | NA | 0.66 | NA |
| E | E-RI-6(0-2) | 0 - 61 | NA | 0.91 | NA |
| E | E-RI-6(2-4) | 61 - 122 | NA | 0.82 | NA |
| E | E-RI-7(0-2) | 0 - 61 | NA | 0.89 | NA |
| E | E-RI-7(2-2.8) | 61 - 85 | NA | 1.15 | NA |
| E | E-RI-8(0-2) | 0 - 61 | NA | 0.93 | NA |
| E | E-RI-8(2-3.25) | 61 - 99 | NA | 1.01 | NA |
| E | E-RI-9(0-2) | 0 - 61 | NA | 0.98 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|--|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| E | E-RI-9(2-4) | 61 - 122 | NA | 1.17 | NA |
| E | E-RI-9(4-5.7) | 122 - 174 | NA | 1.17 | NA |
| E | E-RI-Comp1(0-2) | 0 - 61 | NA | 0.42 | NA |
| E | E-RI-Comp1(2-4) | 61 - 122 | NA | 0.75 | NA |
| E | E-RI-Comp2(0-2) | 0 - 61 | NA | 0.29 | NA |
| E | SDC-E-1-P-S | 0 - 5 | NA | 0.25 | NA |
| E | SDC-E-2-P-S | 0 - 5 | NA | 0.20 | NA |
| E | SDC-E-3-P-S | 0 - 5 | NA | 0.34 | NA |
| E | SDC-E-4-P-S | 0 - 5 | NA | 0.14 | NA |
| E | SDC-E-5-P-S | 0 - 5 | NA | 0.63 | NA |
| E | SDC-E-6-P-S | 0 - 5 | NA | 0.44 | NA |
| E | SDC-E-2-G | 5-35 | 1.18 | 0.43 | 2.62 |
| E | SDC-E-5b-G | 5-35 | 1.12 | 0.32 | 2.24 |
| F | 2F1.1 | 0 - 5 | NA | 0.51 | NA |
| F | 2F1.2 | 5 - 27 | NA | 0.50 | NA |
| F | 2F2.1 | 0 - 5 | NA | 0.36 | NA |
| F | 2F2.2 | 5 - 15 | NA | 0.29 | NA |
| F | 2F2.3 | 15 - 25 | NA | 0.20 | NA |
| F | 2F3.2 | 5 - 15 | NA | 0.36 | NA |
| F | F1.1 | 0 - 4 | NA | 0.28 | NA |
| F | F1.2 | 4 - 15 | NA | 0.22 | NA |
| F | F1.3 | 15 - 26 | NA | 0.25 | NA |
| F | F2.1 | 0 - 6 | NA | 0.33 | NA |
| F | F2.2 | 6 - 17 | NA | 0.23 | NA |
| F | F2.3 | 17 - 29 | NA | 0.20 | NA |
| G | G1.1 | 0 - 17 | NA | 0.63 | NA |
| G | G1.2 | 17 - 30 | NA | 0.54 | NA |
| G | G1.3 | 30 - 37 | NA | 0.88 | NA |
| H | H1.1 | 0 - 8 | NA | 0.83 | NA |
| H | H1.2 | 8 - 17 | NA | 0.99 | NA |
| H | H1.3 | 17 - 38 | NA | 0.91 | NA |
| Reach Average | | | 1.15 | 0.55 | 2.51 |
| Appleton to Little Rapids Reach | | | | | |
| I | I1.1 | 0 - 14 | NA | 0.46 | NA |
| I | I1.2 | 14 - 34 | NA | 0.56 | NA |
| I | I1.3 | 34 - 54 | NA | 0.58 | NA |
| I | POG9 | 0 - 2 | NA | 1.66 | NA |
| J | J1.1 | 0 - 20 | NA | 0.59 | NA |
| J | J1.2 | 20 - 42 | NA | 0.61 | NA |
| J | J1.3 | 42 - 50 | NA | 0.75 | NA |
| K | K1.1 | 0 - 11 | NA | 1.01 | NA |
| K | K1.2 | 11 - 16 | NA | 0.78 | NA |
| K | K1.3 | 16 - 21 | NA | 0.52 | NA |
| L | L1.1 | 0 - 15 | NA | 0.77 | NA |
| L | L1.2 | 15 - 30 | NA | 0.68 | NA |
| L | L1.3 | 30 - 41 | NA | 0.73 | NA |
| L | POG8 | 0 - 5 | NA | 1.92 | NA |
| M | M1.1 | 0 - 19 | NA | 0.49 | NA |
| M | M1.2 | 19 - 36 | NA | 0.43 | NA |
| M | M1.3 | 36 - 41 | NA | 0.46 | NA |
| O | 3O1.2 | 5 - 15 | NA | 0.61 | NA |
| O | 3O1.3 | 15 - 25 | NA | 0.49 | NA |
| O | 3O1.4 | 25 - 35 | NA | 0.42 | NA |
| O | 3O2.1 | 0 - 5 | NA | 0.88 | NA |
| O | O1.1 | 0 - 10 | NA | 0.61 | NA |
| O | O1.2 | 10 - 23 | NA | 0.47 | NA |
| O | O1.3 | 23 - 35 | NA | 0.48 | NA |
| P | 2P1.1 | 0 - 5 | NA | 0.96 | NA |
| P | 2P1.2 | 5 - 15 | NA | 1.33 | NA |
| P | 2P1.4 | 25 - 35 | NA | 1.07 | NA |
| P | 2P1.6 | 45 - 55 | NA | 1.02 | NA |
| P | 2P2.1 | 0 - 5 | NA | 0.24 | NA |
| P | 2P2.2 | 5 - 15 | NA | 0.33 | NA |
| P | 2P2.4 | 25 - 35 | NA | 0.54 | NA |
| P | 2P2.6 | 45 - 56 | NA | 0.57 | NA |
| P | 2P3.1 | 0 - 5 | NA | 0.43 | NA |
| P | 2P3.2 | 5 - 15 | NA | 0.67 | NA |
| P | 2P3.3 | 15 - 25 | NA | 0.42 | NA |
| P | 2P3.6 | 45 - 58 | NA | 0.43 | NA |
| Q | 2Q1.1 | 0 - 5 | NA | 0.45 | NA |
| Q | 2Q1.2 | 5 - 15 | NA | 0.48 | NA |
| Q | 2Q1.3 | 15 - 25 | NA | 0.56 | NA |
| Q | 2Q1.4 | 25 - 35 | NA | 0.50 | NA |
| Q | 2Q1.5 | 35 - 45 | NA | 0.57 | NA |
| Q | 3Q1.3 | 15 - 25 | NA | 0.44 | NA |
| Q | 3Q2.2 | 5 - 15 | NA | 0.56 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| Q | 3Q2.3 | 15 - 25 | NA | 0.39 | NA |
| Q | 3Q2.5 | 35 - 45 | NA | 0.48 | NA |
| R | 2R1.1 | 0 - 5 | NA | 0.99 | NA |
| S | 2S1.1 | 0 - 5 | NA | 1.05 | NA |
| S | 2S1.2 | 5 - 15 | NA | 0.40 | NA |
| T | 2T1.1 | 0 - 5 | NA | 0.27 | NA |
| T | 2T1.2 | 5 - 15 | NA | 0.41 | NA |
| T | T1.1 | 0 - 28 | NA | 0.72 | NA |
| U | POG10 | 0 - 5 | NA | 1.59 | NA |
| U | U1.1 | 0 - 12 | NA | 0.53 | NA |
| U | U1.2 | 12 - 26 | NA | 0.46 | NA |
| U | U1.3 | 26 - 40 | NA | 0.46 | NA |
| V | 2V1.2 | 5 - 15 | NA | 0.59 | NA |
| V | 2V2.1 | 0 - 5 | NA | 0.40 | NA |
| V | 2V2.2 | 5 - 15 | NA | 0.54 | NA |
| V | 2V2.3 | 15 - 25 | NA | 0.32 | NA |
| V | 2V2.4 | 25 - 35 | NA | 0.32 | NA |
| V | 2V2.5 | 35 - 45 | NA | 0.32 | NA |
| W | 2W2.1 | 0 - 5 | NA | 1.40 | NA |
| W | 2W2.2 | 5 - 15 | NA | 0.89 | NA |
| W | 2W4.1 | 0 - 5 | NA | 0.39 | NA |
| W | 2W4.2 | 5 - 15 | NA | 0.90 | NA |
| W | 2W5.2 | 5 - 15 | NA | 0.22 | NA |
| W | 2W6.1 | 0 - 5 | NA | 1.08 | NA |
| W | 2W6.2 | 5 - 22 | NA | 1.17 | NA |
| W | 2W7.1 | 0 - 5 | NA | 0.34 | NA |
| W | 2W7.4 | 25 - 35 | NA | 0.52 | NA |
| W | 2W8.1 | 0 - 5 | NA | 0.43 | NA |
| W | 2W8.3 | 15 - 25 | NA | 0.48 | NA |
| W | 2W9.1 | 0 - 5 | NA | 0.85 | NA |
| W | 2W9.2 | 5 - 15 | NA | 0.43 | NA |
| W | POG4 | 0 - 5 | NA | 1.42 | NA |
| W | SDC-W-1-P-S | 0 - 5 | NA | 0.67 | NA |
| W | SDC-W-2-P-S | 0 - 5 | NA | 0.54 | NA |
| W | SDC-W-3-P-S | 0 - 5 | NA | 1.12 | NA |
| W | SDC-W-4-P-S | 0 - 5 | NA | 0.57 | NA |
| W | SDC-W-5-P-S | 0 - 5 | NA | 0.34 | NA |
| W | W1 | 0 - 19 | NA | 0.64 | NA |
| W | W2.1 | 0 - 23 | NA | 0.47 | NA |
| W | W2.2 | 23 - 48 | NA | 0.51 | NA |
| W | W2.3 | 48 - 74 | NA | 0.50 | NA |
| W | W3.1 | 0 - 15 | NA | 0.59 | NA |
| W | W3.2 | 15 - 26 | NA | 0.48 | NA |
| W | WC1C1 | 0 - 17 | NA | 0.53 | NA |
| W | WC1C2 | 17 - 27 | NA | 0.66 | NA |
| W | WC1C3 | 27 - 41 | NA | 0.68 | NA |
| W | WC2C1 | 0 - 7 | NA | 0.97 | NA |
| W | SDC-W-2-G | 5-35 | 1.15 | 0.46 | 2.38 |
| W | SDC-W-5-G | 5-35 | 1.17 | 0.29 | 2.30 |
| X | 2X1.1 | 0 - 5 | NA | 0.33 | NA |
| X | 2X1.2 | 5 - 19 | NA | 0.39 | NA |
| X | 2X2.1 | 0 - 5 | NA | 0.47 | NA |
| X | 2X2.2 | 5 - 15 | NA | 0.58 | NA |
| X | 2X2.3 | 15 - 28 | NA | 0.42 | NA |
| X | 2X3.1 | 0 - 5 | NA | 0.48 | NA |
| X | 2X3.3 | 15 - 25 | NA | 0.39 | NA |
| X | 2X3.4 | 25 - 35 | NA | 0.44 | NA |
| X | 2X3.6 | 45 - 61 | NA | 0.52 | NA |
| X | 2X4.1 | 0 - 5 | NA | 0.88 | NA |
| X | 2X4.3 | 15 - 25 | NA | 0.83 | NA |
| X | 2X4.4 | 25 - 34 | NA | 0.62 | NA |
| X | 2X5.1 | 0 - 5 | NA | 0.59 | NA |
| X | 2X5.2 | 5 - 15 | NA | 0.60 | NA |
| X | 2X6.1 | 0 - 5 | NA | 0.24 | NA |
| X | 2X6.2 | 5 - 15 | NA | 0.49 | NA |
| X | 2X6.4 | 25 - 35 | NA | 0.53 | NA |
| X | 2X7.1 | 0 - 5 | NA | 1.12 | NA |
| X | 2X7.3 | 15 - 25 | NA | 0.48 | NA |
| X | SDC-X-1-P-S | 0 - 5 | NA | 0.37 | NA |
| X | SDC-X-2-P-S | 0 - 5 | NA | 1.26 | NA |
| X | SDC-X-3-P-S | 0 - 5 | NA | 0.52 | NA |
| X | SDC-X-4-P-S | 0 - 5 | NA | 0.41 | NA |
| X | SDC-X-5-P-S | 0 - 5 | NA | 0.34 | NA |
| X | X1.1 | 0 - 16 | NA | 0.39 | NA |
| X | X1.2 | 16 - 49 | NA | 0.35 | NA |
| X | X1.3 | 49 - 92 | NA | 0.41 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|---------------------------------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| X | XC1C1 | 0 - 8 | NA | 0.42 | NA |
| X | XC1C2 | 8 - 12.5 | NA | 0.44 | NA |
| X | XC1C3 | 12.5 - 17 | NA | 0.46 | NA |
| X | SDC-X-1-G | 5-35 | 1.26 | 0.55 | 2.49 |
| X | SDC-X-2-G | 5-35 | 1.19 | 0.40 | 2.58 |
| Y | Y1.1 | 0 - 14 | NA | 0.74 | NA |
| Y | Y1.2 | 14 - 24 | NA | 0.48 | NA |
| Y | Y1.3 | 24 - 34 | NA | 0.78 | NA |
| Z | Z1.1 | 0 - 11 | NA | 0.80 | NA |
| Z | Z1.2 | 11 - 39 | NA | 0.65 | NA |
| Z | Z1.3 | 39 - 44 | NA | 0.83 | NA |
| AA | AA1.1 | 0 - 6 | NA | 1.32 | NA |
| AA | AA1.2 | 6 - 12 | NA | 1.19 | NA |
| AA | AA1.3 | 12 - 17 | NA | 1.03 | NA |
| BB | BB1.1 | 0 - 14 | NA | 1.16 | NA |
| BB | BB1.2 | 14 - 31 | NA | 0.60 | NA |
| BB | BB1.3 | 31 - 39 | NA | 1.03 | NA |
| CC | 2CC1.1 | 0 - 5 | NA | 0.93 | NA |
| CC | 2CC1.2 | 5 - 15 | NA | 0.76 | NA |
| CC | 2CC1.3 | 15 - 25 | NA | 0.47 | NA |
| CC | 2CC2.1 | 0 - 5 | NA | 0.46 | NA |
| CC | CC1.1 | 0 - 6 | NA | 0.85 | NA |
| CC | CC1.2 | 6 - 12 | NA | 1.12 | NA |
| CC | CC1.3 | 12 - 17 | NA | 1.15 | NA |
| CC | CC2.1 | 0 - 5 | NA | 1.22 | NA |
| CC | CC2.2 | 5 - 9 | NA | 1.00 | NA |
| CC | CC2.3 | 9 - 13 | NA | 1.23 | NA |
| DD | 2DD1.1 | 0 - 5 | NA | 0.66 | NA |
| DD | 2DD1.3 | 15 - 25 | NA | 0.63 | NA |
| DD | 2DD2.1 | 0 - 5 | NA | 0.23 | NA |
| DD | 2DD2.3 | 15 - 26 | NA | 0.44 | NA |
| DD | 2DD3.1 | 0 - 5 | NA | 0.82 | NA |
| DD | 2DD3.2 | 5 - 15 | NA | 1.43 | NA |
| DD | 2DD4.2 | 5 - 15 | NA | 0.68 | NA |
| DD | 2DD4.3 | 15 - 25 | NA | 0.51 | NA |
| DD | 2DD6.1 | 0 - 5 | NA | 0.80 | NA |
| DD | 3DD1.2 | 5 - 15 | NA | 0.69 | NA |
| DD | 3DD1.3 | 15 - 27 | NA | 0.69 | NA |
| DD | 3DD2.1 | 0 - 5 | NA | 1.12 | NA |
| DD | 3DD2.2 | 5 - 15 | NA | 0.45 | NA |
| DD | DD1.1 | 0 - 9 | NA | 0.43 | NA |
| DD | DD1.2 | 9 - 18 | NA | 0.52 | NA |
| DD | DD1.3 | 18 - 26 | NA | 0.64 | NA |
| DD | DD2.1 | 0 - 15 | NA | 0.56 | NA |
| DD | DD2.2 | 15 - 36 | NA | 0.39 | NA |
| DD | DD2.3 | 36 - 49 | NA | 0.59 | NA |
| Reach Average | | | 1.19 | 0.66 | 2.44 |
| Little Rapids to De Pere Reach | | | | | |
| EE | 2EE1.1 | 0 - 5 | NA | 1.43 | NA |
| EE | 2EE1.2 | 5 - 18 | NA | 0.81 | NA |
| EE | 2EE10.1 | 0 - 5 | NA | 0.27 | NA |
| EE | 2EE10.2 | 5 - 15 | NA | 0.34 | NA |
| EE | 2EE10.3 | 15 - 25 | NA | 0.38 | NA |
| EE | 2EE10.4 | 25 - 32 | NA | 0.35 | NA |
| EE | 2EE11.1 | 0 - 5 | NA | 0.25 | NA |
| EE | 2EE11.2 | 5 - 15 | NA | 0.31 | NA |
| EE | 2EE11.3 | 15 - 25 | NA | 0.44 | NA |
| EE | 2EE11.4 | 25 - 35 | NA | 0.45 | NA |
| EE | 2EE12.1 | 0 - 5 | NA | 0.68 | NA |
| EE | 2EE12.2 | 5 - 15 | NA | 0.54 | NA |
| EE | 2EE12.3 | 15 - 28 | NA | 0.57 | NA |
| EE | 2EE13.1 | 0 - 5 | NA | 0.46 | NA |
| EE | 2EE13.2 | 5 - 15 | NA | 0.45 | NA |
| EE | 2EE13.3 | 15 - 25 | NA | 0.66 | NA |
| EE | 2EE13.4 | 25 - 35 | NA | 0.62 | NA |
| EE | 2EE14.1 | 0 - 5 | NA | 0.45 | NA |
| EE | 2EE14.2 | 5 - 15 | NA | 0.28 | NA |
| EE | 2EE14.3 | 15 - 25 | NA | 0.24 | NA |
| EE | 2EE14.4 | 25 - 35 | NA | 0.36 | NA |
| EE | 2EE14.5 | 35 - 46 | NA | 0.33 | NA |
| EE | 2EE15.1 | 0 - 5 | NA | 0.38 | NA |
| EE | 2EE15.2 | 5 - 15 | NA | 0.45 | NA |
| EE | 2EE15.3 | 15 - 29 | NA | 0.48 | NA |
| EE | 2EE16.1 | 0 - 5 | NA | 0.82 | NA |
| EE | 2EE16.2 | 5 - 15 | NA | 0.50 | NA |
| EE | 2EE17.1 | 0 - 5 | NA | 0.37 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| EE | 2EE17.2 | 5 - 15 | NA | 0.37 | NA |
| EE | 2EE17.3 | 15 - 25 | NA | 0.39 | NA |
| EE | 2EE18.1 | 0 - 5 | NA | 1.11 | NA |
| EE | 2EE19.2 | 5 - 15 | NA | 0.41 | NA |
| EE | 2EE19.4 | 25 - 34 | NA | 0.51 | NA |
| EE | 2EE2.1 | 0 - 5 | NA | 0.41 | NA |
| EE | 2EE2.2 | 5 - 15 | NA | 0.47 | NA |
| EE | 2EE2.3 | 15 - 25 | NA | 0.39 | NA |
| EE | 2EE2.4 | 25 - 35 | NA | 0.26 | NA |
| EE | 2EE2.5 | 35 - 45 | NA | 0.24 | NA |
| EE | 2EE2.6 | 45 - 55 | NA | 0.30 | NA |
| EE | 2EE2.7 | 55 - 66 | NA | 0.29 | NA |
| EE | 2EE20.1 | 0 - 5 | NA | 0.53 | NA |
| EE | 2EE21.2 | 5 - 15 | NA | 0.32 | NA |
| EE | 2EE21.3 | 15 - 25 | NA | 0.84 | NA |
| EE | 2EE22.1 | 0 - 5 | NA | 0.30 | NA |
| EE | 2EE22.2 | 5 - 15 | NA | 0.47 | NA |
| EE | 2EE22.3 | 15 - 25 | NA | 0.37 | NA |
| EE | 2EE22.4 | 25 - 35 | NA | 0.34 | NA |
| EE | 2EE22.5 | 35 - 45 | NA | 0.40 | NA |
| EE | 2EE23.1 | 0 - 5 | NA | 0.23 | NA |
| EE | 2EE23.2 | 5 - 15 | NA | 0.33 | NA |
| EE | 2EE23.3 | 15 - 23 | NA | 0.65 | NA |
| EE | 2EE24.1 | 0 - 5 | NA | 0.40 | NA |
| EE | 2EE24.2 | 5 - 15 | NA | 0.55 | NA |
| EE | 2EE24.3 | 15 - 29 | NA | 0.38 | NA |
| EE | 2EE25.1 | 0 - 5 | NA | 0.35 | NA |
| EE | 2EE25.2 | 5 - 15 | NA | 0.33 | NA |
| EE | 2EE25.3 | 15 - 25 | NA | 0.34 | NA |
| EE | 2EE26.1 | 0 - 5 | NA | 1.10 | NA |
| EE | 2EE27.1 | 0 - 5 | NA | 0.38 | NA |
| EE | 2EE27.2 | 5 - 15 | NA | 0.35 | NA |
| EE | 2EE28.1 | 0 - 5 | NA | 0.22 | NA |
| EE | 2EE28.2 | 5 - 15 | NA | 0.47 | NA |
| EE | 2EE28.3 | 15 - 25 | NA | 0.45 | NA |
| EE | 2EE29.1 | 0 - 5 | NA | 0.37 | NA |
| EE | 2EE29.2 | 5 - 15 | NA | 0.43 | NA |
| EE | 2EE3.1 | 0 - 5 | NA | 0.95 | NA |
| EE | 2EE3.2 | 5 - 15 | NA | 1.16 | NA |
| EE | 2EE3.3 | 15 - 28 | NA | 0.78 | NA |
| EE | 2EE30.1 | 0 - 5 | NA | 1.33 | NA |
| EE | 2EE30.2 | 5 - 18 | NA | 0.82 | NA |
| EE | 2EE31.1 | 0 - 5 | NA | 0.41 | NA |
| EE | 2EE31.2 | 5 - 15 | NA | 0.40 | NA |
| EE | 2EE31.3 | 15 - 25 | NA | 0.51 | NA |
| EE | 2EE32.1 | 0 - 5 | NA | 0.31 | NA |
| EE | 2EE32.2 | 5 - 15 | NA | 0.31 | NA |
| EE | 2EE32.3 | 15 - 29 | NA | 0.41 | NA |
| EE | 2EE33.1 | 0 - 5 | NA | 0.24 | NA |
| EE | 2EE33.2 | 5 - 15 | NA | 0.47 | NA |
| EE | 2EE33.3 | 15 - 25 | NA | 0.40 | NA |
| EE | 2EE34.1 | 0 - 5 | NA | 0.62 | NA |
| EE | 2EE34.2 | 5 - 15 | NA | 0.40 | NA |
| EE | 2EE34.3 | 15 - 25 | NA | 0.49 | NA |
| EE | 2EE34.4 | 25 - 35 | NA | 0.49 | NA |
| EE | 2EE34.5 | 35 - 45 | NA | 0.63 | NA |
| EE | 2EE35.1 | 0 - 5 | NA | 0.33 | NA |
| EE | 2EE35.2 | 5 - 15 | NA | 0.36 | NA |
| EE | 2EE35.3 | 15 - 25 | NA | 0.45 | NA |
| EE | 2EE35.4 | 25 - 38 | NA | 0.53 | NA |
| EE | 2EE36.1 | 0 - 5 | NA | 0.32 | NA |
| EE | 2EE36.2 | 5 - 15 | NA | 0.43 | NA |
| EE | 2EE36.3 | 15 - 30 | NA | 0.53 | NA |
| EE | 2EE37.1 | 0 - 5 | NA | 0.95 | NA |
| EE | 2EE37.2 | 5 - 15 | NA | 1.10 | NA |
| EE | 2EE38.1 | 0 - 5 | NA | 0.59 | NA |
| EE | 2EE38.2 | 5 - 15 | NA | 0.34 | NA |
| EE | 2EE39.1 | 0 - 5 | NA | 0.51 | NA |
| EE | 2EE39.2 | 5 - 15 | NA | 0.51 | NA |
| EE | 2EE39.3 | 15 - 25 | NA | 0.48 | NA |
| EE | 2EE4.1 | 0 - 5 | NA | 0.47 | NA |
| EE | 2EE4.2 | 5 - 15 | NA | 0.88 | NA |
| EE | 2EE4.3 | 15 - 25 | NA | 0.49 | NA |
| EE | 2EE4.4 | 25 - 35 | NA | 0.38 | NA |
| EE | 2EE4.5 | 35 - 48 | NA | 0.43 | NA |
| EE | 2EE40.1 | 0 - 5 | NA | 0.46 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| EE | 2EE40.2 | 5 - 15 | NA | 0.51 | NA |
| EE | 2EE40.3 | 15 - 29 | NA | 0.59 | NA |
| EE | 2EE41.1 | 0 - 5 | NA | 0.49 | NA |
| EE | 2EE41.2 | 5 - 15 | NA | 0.22 | NA |
| EE | 2EE41.3 | 15 - 25 | NA | 0.38 | NA |
| EE | 2EE42.1 | 0 - 5 | NA | 0.45 | NA |
| EE | 2EE42.2 | 5 - 15 | NA | 0.33 | NA |
| EE | 2EE42.3 | 15 - 23 | NA | 0.42 | NA |
| EE | 2EE43.1 | 0 - 5 | NA | 0.25 | NA |
| EE | 2EE43.2 | 5 - 15 | NA | 0.29 | NA |
| EE | 2EE43.3 | 15 - 25 | NA | 0.34 | NA |
| EE | 2EE43.4 | 25 - 34 | NA | 0.52 | NA |
| EE | 2EE44.1 | 0 - 5 | NA | 0.32 | NA |
| EE | 2EE44.3 | 15 - 25 | NA | 0.33 | NA |
| EE | 2EE44.4 | 25 - 33 | NA | 0.30 | NA |
| EE | 2EE45.1 | 0 - 5 | NA | 0.31 | NA |
| EE | 2EE45.3 | 15 - 25 | NA | 0.33 | NA |
| EE | 2EE45.4 | 25 - 35 | NA | 0.40 | NA |
| EE | 2EE46.1 | 0 - 5 | NA | 0.49 | NA |
| EE | 2EE46.3 | 15 - 25 | NA | 0.46 | NA |
| EE | 2EE46.4 | 25 - 35 | NA | 0.45 | NA |
| EE | 2EE47.1 | 0 - 5 | NA | 1.17 | NA |
| EE | 2EE48.1 | 0 - 5 | NA | 0.75 | NA |
| EE | 2EE48.2 | 5 - 15 | NA | 0.52 | NA |
| EE | 2EE48.3 | 15 - 25 | NA | 0.65 | NA |
| EE | 2EE49.3 | 15 - 25 | NA | 0.51 | NA |
| EE | 2EE49.4 | 25 - 38 | NA | 0.44 | NA |
| EE | 2EE50.1 | 0 - 5 | NA | 0.45 | NA |
| EE | 2EE50.3 | 15 - 25 | NA | 0.39 | NA |
| EE | 2EE50.4 | 25 - 41 | NA | 0.38 | NA |
| EE | 2EE51.3 | 15 - 25 | NA | 0.37 | NA |
| EE | 2EE51.4 | 25 - 35 | NA | 0.70 | NA |
| EE | 2EE52.2 | 5 - 15 | NA | 0.53 | NA |
| EE | 2EE52.3 | 15 - 30 | NA | 0.62 | NA |
| EE | 2EE53.1 | 0 - 5 | NA | 0.37 | NA |
| EE | 2EE53.2 | 5 - 15 | NA | 0.41 | NA |
| EE | 2EE53.3 | 15 - 25 | NA | 0.56 | NA |
| EE | 2EE53.4 | 25 - 35 | NA | 0.48 | NA |
| EE | 2EE54.1 | 0 - 5 | NA | 0.54 | NA |
| EE | 2EE54.2 | 5 - 15 | NA | 0.54 | NA |
| EE | 2EE54.3 | 15 - 25 | NA | 0.45 | NA |
| EE | 2EE55.1 | 0 - 5 | NA | 0.61 | NA |
| EE | 2EE55.3 | 15 - 25 | NA | 0.50 | NA |
| EE | 2EE55.4 | 25 - 38 | NA | 0.43 | NA |
| EE | 2EE6.1 | 0 - 5 | NA | 0.87 | NA |
| EE | 2EE6.2 | 5 - 15 | NA | 1.40 | NA |
| EE | 2EE6.3 | 15 - 25 | NA | 1.29 | NA |
| EE | 2EE7.1 | 0 - 5 | NA | 0.56 | NA |
| EE | 2EE7.2 | 5 - 15 | NA | 0.37 | NA |
| EE | 2EE7.3 | 15 - 25 | NA | 0.49 | NA |
| EE | 2EE7.4 | 25 - 35 | NA | 0.48 | NA |
| EE | 2EE7.5 | 35 - 51 | NA | 0.43 | NA |
| EE | 2EE8.1 | 0 - 5 | NA | 0.24 | NA |
| EE | 2EE8.2 | 5 - 15 | NA | 0.33 | NA |
| EE | 2EE8.3 | 15 - 25 | NA | 0.23 | NA |
| EE | 2EE8.4 | 25 - 35 | NA | 0.22 | NA |
| EE | 2EE8.5 | 35 - 45 | NA | 0.28 | NA |
| EE | 2EE8.6 | 45 - 55 | NA | 0.29 | NA |
| EE | 2EE8.7 | 55 - 67 | NA | 0.49 | NA |
| EE | 2EE9.1 | 0 - 5 | NA | 0.19 | NA |
| EE | 2EE9.2 | 5 - 15 | NA | 0.35 | NA |
| EE | 2EE9.3 | 15 - 25 | NA | 0.25 | NA |
| EE | 2EE9.4 | 25 - 38 | NA | 0.28 | NA |
| EE | 3EE1.1 | 0 - 5 | NA | 1.22 | NA |
| EE | 3EE1.2 | 5 - 15 | NA | 0.71 | NA |
| EE | 3EE1.3 | 15 - 25 | NA | 0.37 | NA |
| EE | 3EE2.1 | 0 - 5 | NA | 0.51 | NA |
| EE | 3EE2.2 | 5 - 15 | NA | 0.41 | NA |
| EE | 3EE2.3 | 15 - 25 | NA | 0.49 | NA |
| EE | 4EE1.1 | 0 - 2 | NA | 0.35 | NA |
| EE | 4EE1.10 | 18 - 20 | NA | 0.33 | NA |
| EE | 4EE1.11 | 20 - 22 | NA | 0.41 | NA |
| EE | 4EE1.12 | 22 - 24 | NA | 0.41 | NA |
| EE | 4EE1.13 | 24 - 26 | NA | 0.42 | NA |
| EE | 4EE1.14 | 26 - 28 | NA | 0.43 | NA |
| EE | 4EE1.15 | 28 - 30 | NA | 0.39 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| EE | 4EE1.16 | 30 - 32 | NA | 0.37 | NA |
| EE | 4EE1.17 | 32 - 34 | NA | 0.66 | NA |
| EE | 4EE1.18 | 34 - 36 | NA | 0.44 | NA |
| EE | 4EE1.19 | 36 - 38 | NA | 0.49 | NA |
| EE | 4EE1.2 | 2 - 4 | NA | 0.35 | NA |
| EE | 4EE1.20 | 38 - 40 | NA | 0.49 | NA |
| EE | 4EE1.21 | 40 - 42 | NA | 0.53 | NA |
| EE | 4EE1.3 | 4 - 6 | NA | 0.22 | NA |
| EE | 4EE1.4 | 6 - 8 | NA | 0.35 | NA |
| EE | 4EE1.5 | 8 - 10 | NA | 0.52 | NA |
| EE | 4EE1.6 | 10 - 12 | NA | 0.47 | NA |
| EE | 4EE1.8 | 14 - 16 | NA | 0.29 | NA |
| EE | 4EE1.9 | 16 - 18 | NA | 0.38 | NA |
| EE | 4EE2.1 | 0 - 2 | NA | 0.38 | NA |
| EE | 4EE2.10 | 16 - 18 | NA | 0.41 | NA |
| EE | 4EE2.11 | 18 - 20 | NA | 0.38 | NA |
| EE | 4EE2.12 | 20 - 22 | NA | 0.40 | NA |
| EE | 4EE2.13 | 22 - 24 | NA | 0.45 | NA |
| EE | 4EE2.14 | 24 - 26 | NA | 0.52 | NA |
| EE | 4EE2.2 | 2 - 4 | NA | 0.40 | NA |
| EE | 4EE2.3 | 4 - 6 | NA | 0.29 | NA |
| EE | 4EE2.4 | 6 - 8 | NA | 0.30 | NA |
| EE | 4EE2.5 | 8 - 10 | NA | 0.28 | NA |
| EE | 4EE2.6 | 10 - 12 | NA | 0.30 | NA |
| EE | 4EE2.7 | 12 - 14 | NA | 0.30 | NA |
| EE | 4EE2.8 | 14 - 16 | NA | 0.32 | NA |
| EE | 4EE2.9 | 16 - 18 | NA | 0.35 | NA |
| EE | EE1.1 | 0 - 9 | NA | 0.78 | NA |
| EE | EE1.2 | 9 - 20 | NA | 0.64 | NA |
| EE | EE1.3 | 20 - 28 | NA | 0.75 | NA |
| EE | EE10.1 | 0 - 11 | NA | 0.31 | NA |
| EE | EE10.2 | 11 - 27 | NA | 0.45 | NA |
| EE | EE10.3 | 27 - 33 | NA | 0.47 | NA |
| EE | EE11.1 | 0 - 8 | NA | 0.25 | NA |
| EE | EE11.2 | 8 - 26 | NA | 0.34 | NA |
| EE | EE11.3 | 26 - 32 | NA | 0.44 | NA |
| EE | EE12.1 | 0 - 19 | NA | 0.87 | NA |
| EE | EE12.2 | 19 - 38 | NA | 0.67 | NA |
| EE | EE13.1 | 0 - 15 | NA | 0.48 | NA |
| EE | EE13.3 | 22 - 30 | NA | 0.90 | NA |
| EE | EE14.1 | 0 - 20 | NA | 0.89 | NA |
| EE | EE14.2 | 20 - 40 | NA | 0.62 | NA |
| EE | EE14.3 | 40 - 60 | NA | 0.80 | NA |
| EE | EE16.1 | 0 - 5 | NA | 0.42 | NA |
| EE | EE16.2 | 5 - 23 | NA | 0.39 | NA |
| EE | EE16.3 | 23 - 41 | NA | 0.58 | NA |
| EE | EE17.1 | 0 - 5 | NA | 0.33 | NA |
| EE | EE17.2 | 5 - 15 | NA | 0.41 | NA |
| EE | EE17.3 | 15 - 27 | NA | 0.50 | NA |
| EE | EE18.1 | 0 - 6 | NA | 0.23 | NA |
| EE | EE18.2 | 6 - 17 | NA | 0.28 | NA |
| EE | EE18.3 | 17 - 25 | NA | 0.47 | NA |
| EE | EE19.1 | 0 - 3 | NA | 0.33 | NA |
| EE | EE19.2 | 3 - 14 | NA | 0.37 | NA |
| EE | EE19.3 | 14 - 33 | NA | 0.42 | NA |
| EE | EE2.1 | 0 - 7 | NA | 0.48 | NA |
| EE | EE2.2 | 7 - 15 | NA | 0.58 | NA |
| EE | EE2.3 | 15 - 22 | NA | 0.67 | NA |
| EE | EE20.1 | 0 - 6 | NA | 0.37 | NA |
| EE | EE20.2 | 6 - 18 | NA | 0.56 | NA |
| EE | EE20.3 | 18 - 27 | NA | 1.01 | NA |
| EE | EE21.1 | 0 - 4 | NA | 0.40 | NA |
| EE | EE21.2 | 4 - 14 | NA | 0.56 | NA |
| EE | EE21.3 | 14 - 29 | NA | 0.46 | NA |
| EE | EE22.1 | 0 - 4 | NA | 0.28 | NA |
| EE | EE22.2 | 4 - 17 | NA | 0.33 | NA |
| EE | EE22.3 | 17 - 34 | NA | 0.50 | NA |
| EE | EE23.1 | 0 - 17 | NA | 0.28 | NA |
| EE | EE23.2 | 17 - 23 | NA | 0.32 | NA |
| EE | EE23.3 | 23 - 38 | NA | 0.39 | NA |
| EE | EE24.1 | 0 - 10 | NA | 0.38 | NA |
| EE | EE24.2 | 10 - 16 | NA | 0.65 | NA |
| EE | EE24.3 | 16 - 21 | NA | 0.58 | NA |
| EE | EE25.1 | 0 - 10 | NA | 0.29 | NA |
| EE | EE25.2 | 10 - 29 | NA | 0.42 | NA |
| EE | EE25.3 | 29 - 38 | NA | 0.48 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| EE | EE26.1 | 0 - 4 | NA | 0.55 | NA |
| EE | EE26.2 | 4 - 8 | NA | 0.73 | NA |
| EE | EE26.3 | 8 - 42 | NA | 0.42 | NA |
| EE | EE27.1 | 0 - 6 | NA | 0.25 | NA |
| EE | EE27.2 | 6 - 19.5 | NA | 0.34 | NA |
| EE | EE27.3 | 19.5 - 33 | NA | 0.40 | NA |
| EE | EE28.1 | 0 - 22 | NA | 0.40 | NA |
| EE | EE28.2 | 22 - 40 | NA | 0.40 | NA |
| EE | EE28.3 | 40 - 65 | NA | 0.48 | NA |
| EE | EE29.1 | 0 - 26 | NA | 1.30 | NA |
| EE | EE3.1 | 0 - 5 | NA | 0.38 | NA |
| EE | EE3.2 | 5 - 17 | NA | 0.46 | NA |
| EE | EE3.3 | 17 - 31 | NA | 0.55 | NA |
| EE | EE4.1 | 0 - 10 | NA | 0.42 | NA |
| EE | EE4.2 | 10 - 21 | NA | 0.48 | NA |
| EE | EE4.3 | 21 - 28 | NA | 0.52 | NA |
| EE | EE5.1 | 0 - 10 | NA | 0.36 | NA |
| EE | EE5.2 | 10 - 23 | NA | 0.49 | NA |
| EE | EE5.3 | 23 - 28 | NA | 0.44 | NA |
| EE | EE6.1 | 0 - 7 | NA | 0.51 | NA |
| EE | EE6.2 | 7 - 19.5 | NA | 0.39 | NA |
| EE | EE6.3 | 19.5 - 31 | NA | 0.48 | NA |
| EE | EE7.1 | 0 - 10 | NA | 0.43 | NA |
| EE | EE7.2 | 10 - 20.5 | NA | 0.59 | NA |
| EE | EE7.3 | 20.5 - 30 | NA | 0.47 | NA |
| EE | EE8.1 | 0 - 5 | NA | 0.31 | NA |
| EE | EE8.2 | 5 - 19 | NA | 0.36 | NA |
| EE | EE8.3 | 19 - 34 | NA | 0.48 | NA |
| EE | EE9.1 | 0 - 30 | NA | 0.47 | NA |
| EE | EE9.2 | 30 - 59 | NA | 0.53 | NA |
| EE | EE9.3 | 59 - 88 | NA | 0.47 | NA |
| EE | EEC1C1 | 0 - 8 | NA | 0.76 | NA |
| EE | EEC1C2 | 8 - 14 | NA | 1.13 | NA |
| EE | EEC1C3 | 14 - 20 | NA | 0.99 | NA |
| EE | EEC2C1 | 0 - 15 | NA | 0.30 | NA |
| EE | EEC2C2 | 15 - 28 | NA | 0.47 | NA |
| EE | EEC2C3 | 28 - 41 | NA | 0.35 | NA |
| EE | EEC3C1 | 0 - 13 | NA | 0.35 | NA |
| EE | EEC3C2 | 13 - 30 | NA | 0.44 | NA |
| EE | EEC3C3 | 30 - 41 | NA | 0.45 | NA |
| EE | EE-RI-1(0-2) | 0 - 61 | NA | 0.34 | NA |
| EE | EE-RI-1(2-4) | 61 - 122 | NA | 0.50 | NA |
| EE | EE-RI-1(6-7.8) | 183 - 238 | NA | 1.26 | NA |
| EE | EE-RI-10(0-2) | 0 - 61 | NA | 0.29 | NA |
| EE | EE-RI-10(2-4) | 61 - 122 | NA | 0.58 | NA |
| EE | EE-RI-10(6-7.1) | 183 - 216 | NA | 0.57 | NA |
| EE | EE-RI-11(0-2) | 0 - 61 | NA | 0.35 | NA |
| EE | EE-RI-11(2-4) | 61 - 122 | NA | 0.57 | NA |
| EE | EE-RI-11(4-4.5) | 122 - 137 | NA | 1.53 | NA |
| EE | EE-RI-12(0-2) | 0 - 61 | NA | 0.38 | NA |
| EE | EE-RI-12(2-4) | 61 - 122 | NA | 0.51 | NA |
| EE | EE-RI-12(4-4.7) | 122 - 143 | NA | 1.47 | NA |
| EE | EE-RI-13(0-2) | 0 - 61 | NA | 0.33 | NA |
| EE | EE-RI-13(4-6) | 122 - 183 | NA | 0.72 | NA |
| EE | EE-RI-13(6-6.9) | 183 - 210 | NA | 1.73 | NA |
| EE | EE-RI-14(0-0.7) | 0 - 21 | NA | 0.33 | NA |
| EE | EE-RI-15(0-2) | 0 - 61 | NA | 0.27 | NA |
| EE | EE-RI-15(2-4) | 61 - 122 | NA | 0.43 | NA |
| EE | EE-RI-15(6-7.3) | 183 - 223 | NA | 1.25 | NA |
| EE | EE-RI-16(0-1.8) | 0 - 55 | NA | 0.28 | NA |
| EE | EE-RI-17(0-2) | 0 - 61 | NA | 0.29 | NA |
| EE | EE-RI-17(2-3.1) | 61 - 94 | NA | 0.46 | NA |
| EE | EE-RI-18(0-2) | 0 - 61 | NA | 0.35 | NA |
| EE | EE-RI-18(2-3.3) | 61 - 101 | NA | 1.13 | NA |
| EE | EE-RI-19(0-2) | 0 - 61 | NA | 0.38 | NA |
| EE | EE-RI-19(2-4.1) | 61 - 125 | NA | 0.53 | NA |
| EE | EE-RI-2(0-2) | 0 - 61 | NA | 0.30 | NA |
| EE | EE-RI-2(2-4) | 61 - 122 | NA | 0.42 | NA |
| EE | EE-RI-2(4-5) | 122 - 152 | NA | 1.04 | NA |
| EE | EE-RI-20(0-2) | 0 - 61 | NA | 0.30 | NA |
| EE | EE-RI-20(2-4.2) | 61 - 128 | NA | 0.51 | NA |
| EE | EE-RI-21(0-2) | 0 - 61 | NA | 0.27 | NA |
| EE | EE-RI-21(2-4) | 61 - 122 | NA | 0.50 | NA |
| EE | EE-RI-21(4-5.7) | 122 - 174 | NA | 1.33 | NA |
| EE | EE-RI-22(0-2) | 0 - 61 | NA | 0.40 | NA |
| EE | EE-RI-22(2-3.2) | 61 - 98 | NA | 0.68 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| EE | EE-RI-23(0-2) | 0 - 61 | NA | 0.31 | NA |
| EE | EE-RI-23(2-4.1) | 61 - 125 | NA | 0.66 | NA |
| EE | EE-RI-24(0-2) | 0 - 61 | NA | 0.27 | NA |
| EE | EE-RI-24(2-4) | 61 - 122 | NA | 0.43 | NA |
| EE | EE-RI-24(6-7.3) | 183 - 223 | NA | 1.77 | NA |
| EE | EE-RI-25(0-1.6) | 0 - 49 | NA | 0.74 | NA |
| EE | EE-RI-26(0-2) | 0 - 61 | NA | 0.30 | NA |
| EE | EE-RI-26(2-4) | 61 - 122 | NA | 0.46 | NA |
| EE | EE-RI-26(6-6.9) | 183 - 210 | NA | 1.97 | NA |
| EE | EE-RI-27(0-2) | 0 - 61 | NA | 0.32 | NA |
| EE | EE-RI-27(2-4) | 61 - 122 | NA | 0.43 | NA |
| EE | EE-RI-27(4-6.2) | 122 - 189 | NA | 0.78 | NA |
| EE | EE-RI-28(0-2) | 0 - 61 | NA | 0.26 | NA |
| EE | EE-RI-28(2-3.4) | 61 - 104 | NA | 0.44 | NA |
| EE | EE-RI-29(0-2) | 0 - 61 | NA | 0.37 | NA |
| EE | EE-RI-29(2-2.75) | 61 - 84 | NA | 0.60 | NA |
| EE | EE-RI-3(0-2) | 0 - 61 | NA | 0.30 | NA |
| EE | EE-RI-3(2-4) | 61 - 122 | NA | 0.42 | NA |
| EE | EE-RI-3(6-7) | 183 - 213 | NA | 0.58 | NA |
| EE | EE-RI-4(0-2) | 0 - 61 | NA | 0.29 | NA |
| EE | EE-RI-4(2-4) | 61 - 122 | NA | 0.78 | NA |
| EE | EE-RI-4(4-6.1) | 122 - 186 | NA | 1.27 | NA |
| EE | EE-RI-5(0-2) | 0 - 61 | NA | 0.31 | NA |
| EE | EE-RI-5(4-6) | 122 - 183 | NA | 0.54 | NA |
| EE | EE-RI-5(6-8) | 183 - 244 | NA | 0.56 | NA |
| EE | EE-RI-6(0-2) | 0 - 61 | NA | 0.31 | NA |
| EE | EE-RI-6(2-4) | 61 - 122 | NA | 0.44 | NA |
| EE | EE-RI-6(4-5.7) | 122 - 174 | NA | 0.97 | NA |
| EE | EE-RI-7(0-2) | 0 - 61 | NA | 0.34 | NA |
| EE | EE-RI-7(2-4) | 61 - 122 | NA | 0.42 | NA |
| EE | EE-RI-7(6-6.7) | 183 - 204 | NA | 0.70 | NA |
| EE | EE-RI-8(0-2) | 0 - 61 | NA | 0.31 | NA |
| EE | EE-RI-8(4-6) | 122 - 183 | NA | 0.51 | NA |
| EE | EE-RI-8(6-7.7) | 183 - 235 | NA | 0.56 | NA |
| EE | EE-RI-9(0-2) | 0 - 61 | NA | 0.29 | NA |
| EE | EE-RI-9(2-4) | 61 - 122 | NA | 0.41 | NA |
| EE | EE-RI-9(4-5.6) | 122 - 171 | NA | 0.81 | NA |
| EE | POG7 | 0 - 5 | NA | 1.49 | NA |
| EE | SDC-EE22-1-P-S | 0 - 5 | NA | 0.83 | NA |
| EE | SDC-EE22-2-P-S | 0 - 5 | NA | 0.43 | NA |
| EE | SDC-EE22-3-P-S | 0 - 5 | NA | 0.38 | NA |
| EE | SDC-EE22-4-P-S | 0 - 5 | NA | 1.31 | NA |
| EE | SDC-EE23-1-P-S | 0 - 5 | NA | 0.90 | NA |
| EE | SDC-EE23-2-P-S | 0 - 5 | NA | 0.62 | NA |
| EE | SDC-EE23-3-P-S | 0 - 5 | NA | 0.45 | NA |
| EE | SDC-EE23-4-P-S | 0 - 5 | NA | 0.48 | NA |
| EE | SDC-EE23-5-P-S | 0 - 5 | NA | 0.29 | NA |
| EE | SDC-EE24-1-P-S | 0 - 5 | NA | 0.75 | NA |
| EE | SDC-EE24-2-P-S | 0 - 5 | NA | 0.54 | NA |
| EE | SDC-EE24-3-P-S | 0 - 5 | NA | 0.49 | NA |
| EE | SDC-EE24-4-P-S | 0 - 5 | NA | 0.30 | NA |
| EE | SDC-EE24-5-P-S | 0 - 5 | NA | 0.40 | NA |
| EE | SDC-EE25-1-P-S | 0 - 5 | NA | 0.26 | NA |
| EE | SDC-EE25-2-P-S | 0 - 5 | NA | 0.34 | NA |
| EE | SDC-EE25-3-P-S | 0 - 5 | NA | 0.33 | NA |
| EE | SDC-EE25-4-P-S | 0 - 5 | NA | 0.39 | NA |
| EE | SDC-EE25-5-P-S | 0 - 5 | NA | 0.30 | NA |
| EE | SDC-EE26-1-P-S | 0 - 5 | NA | 0.39 | NA |
| EE | SDC-EE26-2-P-S | 0 - 5 | NA | 0.37 | NA |
| EE | SDC-EE26-3-P-S | 0 - 5 | NA | 0.34 | NA |
| EE | SDC-EE26-4-P-S | 0 - 5 | NA | 0.44 | NA |
| EE | SDC-EE26-5-P-S | 0 - 5 | NA | 0.22 | NA |
| EE | SDC-EE27-1-P-S | 0 - 5 | NA | 0.36 | NA |
| EE | SDC-EE27-2-P-S | 0 - 5 | NA | 0.37 | NA |
| EE | SDC-EE22-3-G | 5-35 | 1.17 | 0.42 | 2.61 |
| EE | SDC-EE22-4-G | 5-35 | 1.21 | 0.56 | 2.56 |
| EE | SDC-EE23-1-G | 5-32 | 1.48 | 0.93 | 2.36 |
| EE | SDC-EE23-4-G | 5-35 | 1.21 | 0.44 | 2.33 |
| EE | SDC-EE24-1-G | 5-35 | 0.93 | 0.29 | 2.44 |
| EE | SDC-EE24-3-G | 5-35 | 1.15 | 0.45 | 2.55 |
| EE | SDC-EE25-2-G | 5-35 | 1.14 | 0.38 | 2.43 |
| EE | SDC-EE25-3-G | 5-35 | 1.11 | 0.37 | 2.42 |
| EE | SDC-EE26-2-G | 5-35 | 1.13 | 0.36 | 2.52 |
| EE | SDC-EE26-5-G | 5-35 | 1.11 | 0.37 | 2.52 |
| EG | EGH-RI-Comp1(0-2) | 0 - 61 | NA | 0.31 | NA |
| EG | EGH-RI-Comp1(2-4) | 61 - 122 | NA | 0.43 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| EG | EGH-RI-Comp1(4-6) | 122 - 183 | NA | 0.55 | NA |
| EG | EGH-RI-Comp1(6-8) | 183 - 244 | NA | 1.24 | NA |
| FF | 3FF1.1 | 0 - 5 | NA | 1.87 | NA |
| FF | 3FF1.2 | 5 - 15 | NA | 1.12 | NA |
| FF | 3FF1.3 | 15 - 25 | NA | 1.18 | NA |
| FF | 3FF2.1 | 0 - 5 | NA | 0.40 | NA |
| FF | 3FF2.3 | 15 - 25 | NA | 0.39 | NA |
| FF | 3FF2.4 | 25 - 35 | NA | 0.55 | NA |
| FF | FF1.1 | 0 - 10 | NA | 0.25 | NA |
| FF | FF1.2 | 10 - 22 | NA | 0.35 | NA |
| FF | FF1.3 | 22 - 46 | NA | 0.41 | NA |
| GG | 2GG1.1 | 0 - 5 | NA | 0.37 | NA |
| GG | 2GG1.2 | 5 - 15 | NA | 0.48 | NA |
| GG | 2GG1.3 | 15 - 25 | NA | 0.88 | NA |
| GG | 2GG1.5 | 25 - 35 | NA | 0.58 | NA |
| GG | 2GG2.1 | 0 - 5 | NA | 0.40 | NA |
| GG | 2GG2.2 | 5 - 15 | NA | 0.41 | NA |
| GG | 2GG2.3 | 15 - 25 | NA | 0.36 | NA |
| GG | 2GG2.4 | 25 - 35 | NA | 0.46 | NA |
| GG | 2GG2.5 | 35 - 45 | NA | 0.68 | NA |
| GG | 2GG3.3 | 15 - 25 | NA | 0.35 | NA |
| GG | 2GG3.4 | 25 - 39 | NA | 0.45 | NA |
| GG | 3GG20.1 | 0 - 5 | NA | 0.46 | NA |
| GG | 3GG20.2 | 5 - 15 | NA | 0.37 | NA |
| GG | 3GG20.3 | 15 - 25 | NA | 0.38 | NA |
| GG | 3GG20.4 | 25 - 35 | NA | 0.36 | NA |
| GG | 3GG20.5 | 35 - 45 | NA | 0.40 | NA |
| GG | 3GG20.6 | 45 - 55 | NA | 0.48 | NA |
| GG | GG1.1 | 0 - 12 | NA | 0.31 | NA |
| GG | GG1.2 | 12 - 22 | NA | 0.31 | NA |
| GG | GG1.3 | 22 - 62 | NA | 0.41 | NA |
| GG | GG1.4 | 62 - 85 | NA | 0.47 | NA |
| GG | GG2.1 | 0 - 7 | NA | 0.40 | NA |
| GG | GG2.2 | 7 - 19 | NA | 0.35 | NA |
| GG | GG2.3 | 19 - 36 | NA | 0.43 | NA |
| GG | GG-RI-1(0-2) | 0 - 61 | NA | 0.33 | NA |
| GG | GG-RI-1(2-4.2) | 61 - 128 | NA | 0.49 | NA |
| GG | GG-RI-10(0-0.9) | 0 - 27 | NA | 0.44 | NA |
| GG | GG-RI-11(0-2) | 0 - 61 | NA | 0.37 | NA |
| GG | GG-RI-11(2-3.7) | 61 - 113 | NA | 0.51 | NA |
| GG | GG-RI-12(0-2) | 0 - 61 | NA | 0.58 | NA |
| GG | GG-RI-12(2-2.5) | 61 - 76 | NA | 1.86 | NA |
| GG | GG-RI-13(0-2) | 0 - 61 | NA | 0.37 | NA |
| GG | GG-RI-13(2-4.1) | 61 - 125 | NA | 0.90 | NA |
| GG | GG-RI-14(0-1.1) | 0 - 34 | NA | 0.37 | NA |
| GG | GG-RI-15(0-2) | 0 - 61 | NA | 0.37 | NA |
| GG | GG-RI-15(2-4.2) | 61 - 128 | NA | 0.61 | NA |
| GG | GG-RI-2(0-2) | 0 - 61 | NA | 0.37 | NA |
| GG | GG-RI-2(2-2.9) | 61 - 88 | NA | 0.73 | NA |
| GG | GG-RI-3(0-2) | 0 - 61 | NA | 0.33 | NA |
| GG | GG-RI-3(2-3.7) | 61 - 113 | NA | 0.44 | NA |
| GG | GG-RI-4(0-2) | 0 - 61 | NA | 0.31 | NA |
| GG | GG-RI-4(2-4) | 61 - 122 | NA | 0.38 | NA |
| GG | GG-RI-4(4-5.2) | 122 - 158 | NA | 0.55 | NA |
| GG | GG-RI-5(0-2.2) | 0 - 67 | NA | 0.39 | NA |
| GG | GG-RI-6(0-2) | 0 - 61 | NA | 0.33 | NA |
| GG | GG-RI-6(2-4) | 61 - 122 | NA | 0.45 | NA |
| GG | GG-RI-6(4-5.2) | 122 - 158 | NA | 0.58 | NA |
| GG | GG-RI-7(0-2) | 0 - 61 | NA | 0.39 | NA |
| GG | GG-RI-8(0-2) | 0 - 61 | NA | 0.32 | NA |
| GG | GG-RI-8(2-4) | 61 - 122 | NA | 0.42 | NA |
| GG | GG-RI-8(4-5.1) | 122 - 155 | NA | 0.50 | NA |
| GG | GG-RI-9(0-2) | 0 - 61 | NA | 0.34 | NA |
| GG | GG-RI-9(2-4.2) | 61 - 128 | NA | 0.77 | NA |
| HH | 2HH1.1 | 0 - 5 | NA | 0.30 | NA |
| HH | 2HH1.2 | 5 - 15 | NA | 0.32 | NA |
| HH | 2HH1.3 | 15 - 26 | NA | 0.33 | NA |
| HH | 2HH10.1 | 0 - 5 | NA | 0.25 | NA |
| HH | 2HH10.2 | 5 - 15 | NA | 0.43 | NA |
| HH | 2HH10.3 | 15 - 25 | NA | 0.46 | NA |
| HH | 2HH10.4 | 25 - 35 | NA | 0.38 | NA |
| HH | 2HH10.5 | 35 - 50 | NA | 0.34 | NA |
| HH | 2HH11.1 | 0 - 5 | NA | 0.35 | NA |
| HH | 2HH2.3 | 15 - 30 | NA | 0.32 | NA |
| HH | HH1.1 | 0 - 11 | NA | 0.24 | NA |
| HH | HH1.2 | 11 - 34 | NA | 0.32 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-----------------------------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| HH | HH1.3 | 34 - 46 | NA | 0.34 | NA |
| HH | HH-RI-1(0-2) | 0 - 61 | NA | 0.39 | NA |
| HH | HH-RI-1(2-3) | 61 - 91 | NA | 1.85 | NA |
| HH | HH-RI-10(0-0.7) | 0 - 21 | NA | 0.55 | NA |
| HH | HH-RI-2(0-2) | 0 - 61 | NA | 0.35 | NA |
| HH | HH-RI-2(2-3.25) | 61 - 99 | NA | 1.71 | NA |
| HH | HH-RI-3(0-2) | 0 - 61 | NA | 0.31 | NA |
| HH | HH-RI-3(2-4) | 61 - 122 | NA | 0.38 | NA |
| HH | HH-RI-3(4-6) | 122 - 183 | NA | 0.48 | NA |
| HH | HH-RI-3(6-6.7) | 183 - 204 | NA | 0.96 | NA |
| HH | HH-RI-4(0-1.2) | 0 - 37 | NA | 0.28 | NA |
| HH | HH-RI-5(0-2) | 0 - 61 | NA | 0.51 | NA |
| HH | HH-RI-5(2-4) | 61 - 122 | NA | 1.31 | NA |
| HH | HH-RI-5(4-5.1) | 122 - 155 | NA | 1.38 | NA |
| HH | HH-RI-6(0-2) | 0 - 61 | NA | 0.30 | NA |
| HH | HH-RI-6(2-4) | 61 - 122 | NA | 0.35 | NA |
| HH | HH-RI-6(4-5.2) | 122 - 158 | NA | 0.46 | NA |
| HH | HH-RI-7(0-0.5) | 0 - 15 | NA | 0.41 | NA |
| HH | HH-RI-8(0-2) | 0 - 61 | NA | 0.28 | NA |
| HH | HH-RI-8(2-2.9) | 61 - 88 | NA | 0.53 | NA |
| HH | HH-RI-9(0-2) | 0 - 61 | NA | 0.30 | NA |
| HH | HH-RI-9(2-3.7) | 61 - 113 | NA | 0.40 | NA |
| | 2FRA1.2 | 69 - 129 | NA | 0.98 | NA |
| | 2FRA2.2 | 34 - 64 | NA | 0.31 | NA |
| | 2FRA4.2 | 34 - 60 | NA | 0.31 | NA |
| | 2FRA4.3 | 69 - 96 | NA | 0.37 | NA |
| | 2FRA4.4 | 103 - 136 | NA | 0.44 | NA |
| | 2FRA4.5 | 137 - 162 | NA | 0.37 | NA |
| | 2FRA5.2 | 34 - 55 | NA | 0.29 | NA |
| | 2FRA5.3 | 69 - 98 | NA | 0.40 | NA |
| | 2FRA5.4 | 103 - 132 | NA | 0.41 | NA |
| | 2FRA5.5 | 137 - 167 | NA | 0.52 | NA |
| | POG6 | 0 - 5 | NA | 1.46 | NA |
| Reach Average | | | 1.16 | 0.51 | 2.47 |
| De Pere to Green Bay Reach | | | | | |
| 20 | 2FRBg27.1 | 0 - 5 | NA | 0.65 | NA |
| 20 | 2FRBg27.2 | 5 - 15 | NA | 0.62 | NA |
| 20 | 2FRBg27.3 | 15 - 25 | NA | 1.39 | NA |
| 20 | SDC-DPD-1-P-S | 0 - 5 | NA | 0.30 | NA |
| 20 | SDC-DPD-2-P-S | 0 - 5 | NA | 0.27 | NA |
| 20 | SDC-DPD-2-G | 5-35 | 1.15 | 0.39 | 2.32 |
| 21 | 2FRBg24.1 | 0 - 5 | NA | 0.35 | NA |
| 21 | 2FRBg24.2 | 5 - 15 | NA | 0.34 | NA |
| 21 | 2FRBg24.3 | 15 - 25 | NA | 0.31 | NA |
| 21 | 2FRBg24.4 | 25 - 44 | NA | 0.42 | NA |
| 21 | 2FRBg26.2 | 5 - 15 | NA | 0.82 | NA |
| 21 | 2FRBg26.3 | 15 - 25 | NA | 0.40 | NA |
| 23 | 2FRBg22.1 | 0 - 5 | NA | 0.52 | NA |
| 23 | 2FRBg22.2 | 5 - 15 | NA | 0.41 | NA |
| 25 | 2FRBg23.1 | 0 - 5 | NA | 0.95 | NA |
| 25 | 2FRBg23.3 | 15 - 25 | NA | 1.32 | NA |
| 25 | 2FRBg23.4 | 25 - 35 | NA | 0.74 | NA |
| 34 | 2FRBg20.1 | 0 - 5 | NA | 0.40 | NA |
| 34 | 2FRBg6.1 | 0 - 5 | NA | 0.31 | NA |
| 34 | 2FRBg6.2 | 5 - 15 | NA | 0.28 | NA |
| 34 | 2FRBg6.3 | 15 - 25 | NA | 0.35 | NA |
| 34 | 2FRBg6.5 | 35 - 45 | NA | 0.36 | NA |
| 41 | 2FRBg17.1 | 0 - 5 | NA | 0.31 | NA |
| 41 | 2FRBg17.2 | 5 - 15 | NA | 0.40 | NA |
| 41 | 2FRBg17.3 | 15 - 25 | NA | 0.41 | NA |
| 41 | 2FRBg17.4 | 25 - 31 | NA | 0.35 | NA |
| 43 | 2FRBg18.1 | 0 - 5 | NA | 0.56 | NA |
| 43 | 2FRBg18.2 | 5 - 15 | NA | 0.63 | NA |
| 43 | 2FRBg18.3 | 15 - 25 | NA | 0.83 | NA |
| 45 | 2FRBg13.1 | 0 - 5 | NA | 0.37 | NA |
| 45 | 2FRBg13.3 | 15 - 25 | NA | 0.42 | NA |
| 45 | 2FRBg13.4 | 25 - 35 | NA | 0.48 | NA |
| 45 | 2FRBg14.1 | 0 - 5 | NA | 0.43 | NA |
| 45 | 2FRBg14.3 | 15 - 25 | NA | 0.46 | NA |
| 45 | 2FRBg14.4 | 25 - 35 | NA | 0.59 | NA |
| 45 | SDC-DPD-3-P-S | 0 - 5 | NA | 0.28 | NA |
| 45 | SDC-DPD-3-G | 5-35 | 1.21 | 0.45 | 2.40 |
| 46 | 2FRBg16.1 | 0 - 5 | NA | 0.67 | NA |
| 46 | 2FRBg16.2 | 5 - 15 | NA | 0.49 | NA |
| 48 | 2FRBg15.1 | 0 - 5 | NA | 0.55 | NA |
| 48 | 2FRBg15.2 | 5 - 15 | NA | 1.55 | NA |

Appendix G - Table 7
Lower Fox River - Bulk Density Results

| Deposit/SMU | Sample Identification | Depth (cm) | Wet Density (g/cm ³) | Dry Density (g/cm ³) | Specific Gravity |
|-----------------------------|-----------------------|------------|----------------------------------|----------------------------------|------------------|
| 49 | 2FRBg12.1 | 0 - 5 | NA | 0.63 | NA |
| 49 | 2FRBg12.2 | 5 - 15 | NA | 0.94 | NA |
| 51 | 2FRBg3.1 | 0 - 5 | NA | 1.02 | NA |
| 51 | 2FRBg3.2 | 5 - 15 | NA | 1.27 | NA |
| 51 | 2FRBg3.3 | 15 - 26 | NA | 0.62 | NA |
| 52 | 2FRBg1.1 | 0 - 5 | NA | 0.37 | NA |
| 52 | 2FRBg1.2 | 5 - 15 | NA | 0.38 | NA |
| 52 | 2FRBg1.3 | 15 - 25 | NA | 0.37 | NA |
| 52 | 2FRBg1.4 | 25 - 33 | NA | 0.37 | NA |
| 53 | 2FRBg2.1 | 0 - 5 | NA | 0.32 | NA |
| 53 | 2FRBg2.2 | 5 - 15 | NA | 0.37 | NA |
| 53 | 2FRBg2.4 | 25 - 35 | NA | 0.44 | NA |
| 57 | 2FRBg5.1 | 0 - 5 | NA | 0.40 | NA |
| 57 | 2FRBg5.2 | 5 - 15 | NA | 0.46 | NA |
| 57 | 2FRBg5.3 | 15 - 25 | NA | 0.42 | NA |
| 61 | 2FRBg4.1 | 0 - 5 | NA | 0.98 | NA |
| 61 | 2FRBg4.2 | 5 - 15 | NA | 0.99 | NA |
| 70 | 2FRBg7.1 | 0 - 5 | NA | 0.30 | NA |
| 70 | 2FRBg7.2 | 5 - 15 | NA | 0.36 | NA |
| 70 | 2FRBg7.4 | 25 - 35 | NA | 0.41 | NA |
| 71 | 2FRBg11.1 | 0 - 5 | NA | 0.52 | NA |
| 71 | 2FRBg11.2 | 5 - 15 | NA | 0.42 | NA |
| 71 | 2FRBg11.3 | 15 - 25 | NA | 0.36 | NA |
| 77 | 2FRBg8.1 | 0 - 5 | NA | 0.67 | NA |
| 77 | 2FRBg8.2 | 5 - 15 | NA | 0.68 | NA |
| 77 | 2FRBg8.3 | 15 - 21 | NA | 0.78 | NA |
| 86 | 2FRBg9.1 | 0 - 5 | NA | 1.49 | NA |
| 86 | 2FRBg9.2 | 5 - 15 | NA | 0.52 | NA |
| 86 | 2FRBg9.3 | 15 - 27 | NA | 0.34 | NA |
| 94 | 2FRBg30.1 | 0 - 5 | NA | 0.43 | NA |
| 94 | 2FRBg30.2 | 5 - 15 | NA | 0.41 | NA |
| 94 | 2FRBg31.2 | 5 - 15 | NA | 0.30 | NA |
| 96 | SDC-DPD-4-P-S | 0 - 5 | NA | 1.32 | NA |
| 106 | 2FRBg28.1 | 0 - 5 | NA | 0.57 | NA |
| 106 | 2FRBg28.2 | 5 - 15 | NA | 0.68 | NA |
| 112 | 2FRBg10.1 | 0 - 5 | NA | 0.74 | NA |
| 112 | 2FRBg10.2 | 5 - 15 | NA | 0.46 | NA |
| 112 | 2FRBg10.4 | 25 - 35 | na | 0.51 | NA |
| 115 | SDC-DPD-5-P-S | 0 - 5 | NA | 0.28 | NA |
| | 2FRBg19.1 | 0 - 5 | NA | 0.63 | NA |
| | 2FRBg21.1 | 0 - 5 | NA | 0.58 | NA |
| | 2FRBg25.1 | 0 - 5 | NA | 1.61 | NA |
| | 2FRBg25.2 | 5 - 15 | NA | 1.32 | NA |
| Reach Average | | | 1.18 | 0.59 | 2.36 |
| Entire River Average | | | 1.17 | 0.55 | 2.46 |

APPENDIX H

**PCB CONGENERS AND HOMOLOG GROUP LIST
(ATSDR, 1997a)**

3. CHEMICAL AND PHYSICAL INFORMATIONS

Table 3-2. Chemical Identity of Polychlorinated Biphenyl Congeners

| IUPAC No. ^a | Compound/substituents | CAS No. ^b |
|------------------------|-----------------------|----------------------|
| Monochlorobiphenyl | | 27323-18-8 |
| 1 | 2 | 2051-60-7 |
| 2 | 3 | 2051-60-8 |
| 3 | 4 | 2051-60-9 |
| Dichlorobiphenyl | | 25512-42-9 |
| 4 | 2,2' | 13029-08-8 |
| 5 | 2,3 | 16605-91-7 |
| 6 | 2,3' | 25569-80-6 |
| 7 | 2,4 | 33284-50-3 |
| 8 | 2,4' | 34883-43-7 |
| 9 | 2,5 | 34883-39-1 |
| 10 | 2,6 | 33146-45-1 |
| 11 | 3,3' | 2050-67-1 |
| 12 | 3,4 | 2974-92-7 |
| 13 | 3,4' | 2974-90-5 |
| 14 | 3,5 | 34883-41-5 |
| 15 | 4,4' | 2050-68-2 |
| Trichlorobiphenyl | | 25323-68-6 |
| 16 | 2,2',3 | 38444-78-9 |
| 17 | 2,2',4 | 37680-66-3 |
| 18 | 2,2',5 | 37680-65-2 |
| 19 | 2,2',6 | 38444-73-4 |
| 20 | 2,3,3' | 38444-84-7 |
| 21 | 2,3,4 | 55702-46-0 |
| 22 | 2,3,4' | 38444-85-8 |
| 23 | 2,3,5 | 55720-44-0 |
| 24 | 2,3,6 | 58702-45-9 |
| 25 | 2,3',4 | 55712-37-3 |
| 26 | 2,3',5 | 38444-81-4 |
| 27 | 2,3,6 | 38444-76-7 |
| 28 | 2,4,4' | 7012-37-5 |
| 29 | 2,4,5 | 15862-07-4 |
| 30 | 2,4,6 | 35693-92-6 |
| 31 | 2,4',5 | 16606-02-3 |
| 32 | 2,4',6 | 38444-77-8 |
| 33 | 2',3,4 | 38444-86-9 |
| 34 | 2',3,5 | 37680-68-5 |
| 35 | 3,3',4 | 37680-69-6 |

PCBs

3. CHEMICAL AND PHYSICAL INFORMATION

Table 3-2. Chemical Identity of Polychlorinated Biphenyl Congeners (continued)

| IUPAC No. ^a | Compound/substituents | CAS No. ^b |
|------------------------|-----------------------|----------------------|
| 36 | 3,3',5 | 38444-87-0 |
| 37 | 3,4,4' | 38444-90-5 |
| 38 | 3,4,5 | 53555-66-1 |
| 39 | 3,4',5 | 38444-88-1 |
| Tetrachlorobiphenyl | | 26914-33-0 |
| 40 | 2,2',3,3' | 38444-93-8 |
| 41 | 2,2',3,4 | 52663-59-9 |
| 42 | 2,2',3,4' | 36559-22-5 |
| 43 | 2,2',3,5 | 70362-46-8 |
| 44 | 2,2',3,5' | 41464-39-5 |
| 45 | 2,2',3,6 | 70362-45-7 |
| 46 | 2,2',3,6' | 41464-47-5 |
| 47 | 2,2',4,4' | 2437-79-8 |
| 48 | 2,2',4,5 | 70362-47-9 |
| 49 | 2,2',4,5' | 41464-40-8 |
| 50 | 2,2',4,6 | 62796-65-8 |
| 51 | 2,2',4,6' | 64194-04-7 |
| 52 | 2,2',5,5' | 35693-99-3 |
| 53 | 2,2',5,6' | 41464-41-9 |
| 54 | 2,2',6,6' | 15968-05-5 |
| 55 | 2,3,3',4 | 74338-24-2 |
| 56 | 2,3,3',4' | 41464-43-1 |
| 57 | 2,3,3',5 | 70424-67-8 |
| 58 | 2,3,3',5' | 41464-49-7 |
| 59 | 2,3,3',6 | 74472-33-6 |
| 60 | 2,3,4,4' | 33025-41-1 |
| 61 | 2,3,4,5 | 33284-53-6 |
| 62 | 2,3,4,6 | 54230-23-7 |
| 63 | 2,3,4',5 | 74472-35-8 |
| 64 | 2,3,4',6 | 52663-58-8 |
| 65 | 2,3,5,6 | 33284-54-7 |
| 66 | 2,3',4,4' | 32698-10-0 |
| 67 | 2,3',4,5 | 73575-53-8 |
| 68 | 2,3',4,5' | 73575-52-7 |
| 69 | 2,3',4,6 | 60233-24-1 |
| 70 | 2,3',4',5 | 32598-11-1 |
| 71 | 2,3',4',6 | 41464-46-4 |
| 72 | 2,3',5,5' | 41464-42-0 |

3. CHEMICAL AND PHYSICAL INFORMATIONS

Table 3-2. Chemical Identity of Polychlorinated Biphenyl Congeners (continued)

| IUPAC No. ^a | Compound/substituents | CAS No. ^b |
|------------------------|-----------------------|----------------------|
| 73 | 2,3',5',6 | 743338-23-1 |
| 74 | 2,4,4',5 | 32690-93-0 |
| 75 | 2,4,4',6 | 32598-12-2 |
| 76 | 2',3,4,5 | 70362-48-0 |
| 77 | 3,3',4,4' | 32598-13-3 |
| 78 | 3,3',4,5 | 70362-49-1 |
| 79 | 3,3',4,5' | 41464-48-6 |
| 80 | 3,3',5,5' | 33284-52-5 |
| 81 | 3,4,4',5 | 70362-50-4 |
| Pentachlorobiphenyl | | 25429-29-2 |
| 82 | 2,2',3,3',4 | 52663-62-4 |
| 83 | 2,2',3,3',5 | 60145-20-2 |
| 84 | 2,2',3,3',6 | 52663-60-2 |
| 85 | 2,2',3,4,4' | 65510-45-4 |
| 86 | 2,2',3,4,5 | 55312-69-1 |
| 87 | 2,2',3,4,5' | 38380-02-8 |
| 88 | 2,2',3,4,6 | 55215-17-3 |
| 89 | 2,2',3,4,6' | 73575-57-2 |
| 90 | 2,2',3,4',5 | 68194-07-0 |
| 91 | 2,2',3,4',6 | 58194-05-8 |
| 92 | 2,2',3,5,5' | 52663-61-3 |
| 93 | 2,2',3,5,6 | 73575-56-1 |
| 94 | 2,2',3,5,6' | 73575-55-0 |
| 95 | 2,2',3,5',6 | 38379-99-6 |
| 96 | 2,2',3,6,6' | 73575-54-9 |
| 97 | 2,2',3',4,5 | 41464-51-1 |
| 98 | 2,2',3',4,6 | 60233-25-2 |
| 99 | 2,2',3',4',5 | 38380-01-7 |
| 100 | 2,2',4',4',6 | 39485-83-1 |
| 101 | 2,2',4,5,5' | 37680-73-2 |
| 102 | 2,2',4,5,6' | 68194-06-9 |
| 103 | 2,2',4,5',6 | 61045-21-3 |
| 104 | 2,2',4,6,6' | 56558-16-8 |
| 105 | 2,3,3',4,4' | 32598-14-4 |
| 106 | 2,3,3',4,5 | 70424-69-0 |
| 107 | 2,3,3',4',5 | 70424-68-9 |
| 108 | 2,3,3',4,5' | 70362-41-3 |
| 109 | 2,3,3',4,6 | 74472-35-8 |

3. CHEMICAL AND PHYSICAL INFORMATIONS

Table 3-2. Chemical Identity of Polychlorinated Biphenyl Congeners (continued)

| IUPAC No. ^a | Compound/substituents | CAS No. ^b |
|------------------------|-----------------------|----------------------|
| 110 | 2,3,3',4',6 | 38380-03-9 |
| 111 | 2,3,3',5,5' | 39635-32-0 |
| 112 | 2,3,3',5,6 | 74472-36-9 |
| 113 | 2,3,3',5',6 | 68194-10-5 |
| 114 | 2,3,4,4',5 | 74472-37-0 |
| 115 | 2,3,4,4',6 | 74472-38-1 |
| 116 | 2,3,4,5,6 | 18259-05-7 |
| 117 | 2,3,4',5,6 | 68194-11-6 |
| 118 | 2,3',4,4',5 | 31508-00-6 |
| 119 | 2,3',4,4',6 | 56558-17-9 |
| 120 | 2,3',4,5,5' | 68194-12-7 |
| 121 | 2,3',4,5',6 | 56558-18-0 |
| 122 | 2',3,3',4,5 | 76842-07-4 |
| 123 | 2',3,4,4',5 | 65510-44-3 |
| 124 | 2',3,4,5,5' | 70424-70-3 |
| 125 | 2',3,4,5,6' | 74472-39-2 |
| 126 | 3,3',4,4',5 | 57465-28-8 |
| 127 | 3,3',4,5,5' | 39635-33-1 |
| Hexachlorobiphenyl | | 26601-64-9 |
| 128 | 2,2',3,3',4,4' | 38380-07-3 |
| 129 | 2,2',3,3',4,5 | 55215-18-4 |
| 130 | 2,2',3,3',4,5' | 52663-3-66-8 |
| 131 | 2,2',3,3',4,6 | 61798-70-7 |
| 132 | 2,2',3,3',4,6' | 38380-05-1 |
| 133 | 2,2',3,3',5,5' | 35694-04-3 |
| 134 | 2,2',3,3',5,6 | 52704-70-8 |
| 135 | 2,2',3,3',5,6' | 52744-13-5 |
| 136 | 2,2',3,3',6,6' | 38411-22-2 |
| 137 | 2,2',3,4,4',5 | 35694-06-5 |
| 138 | 2,2',3,4,4',5' | 35065-28-2 |
| 139 | 2,2',3,4,4',6 | 56030-56-9 |
| 140 | 2,2',3,4,4',6' | 59291-64-4 |
| 141 | 2,2',3,4,5,5' | 52712-04-6 |
| 142 | 2,2',3,4,5,6 | 41411-61-4 |
| 143 | 2,2',3,4,5,6' | 68194-15-0 |
| 144 | 2,2',3,4,5',6 | 68194-14-9 |
| 145 | 2,2',3,4',6,6' | 74472-40-5 |
| 146 | 2,2',3,4',5,5' | 51908-16-8 |

3. CHEMICAL AND PHYSICAL INFORMATION

Table 3-2. Chemical Identity of Polychlorinated Biphenyl Congeners (continued)

| IUPAC No. ^a | Compound/substituents | CAS No. ^b |
|----------------------------|-----------------------|----------------------|
| 147 | 2,2',3,4',5,6 | 68194-13-8 |
| 148 | 2,2',3,4',5,6' | 74472-42-7 |
| 149 | 2,2',3,4',5',6 | 38380-04-0 |
| 150 | 2,2',3,4',5,6' | 68194-08-1 |
| 151 | 2,2',3,5,5',6 | 52663-63-5 |
| 152 | 2,2',3,5,6,6' | 68194-09-2 |
| 153 | 2,2',4,4',5,5' | 35065-27-1 |
| 154 | 2,2',4,4',5,6' | 60145-22-4 |
| 155 | 2,2',4,4',6,6' | 33979-03-2 |
| 156 | 2,3,3',4,4',5 | 38380-08-4 |
| 157 | 2,3,3',4,4',5' | 69782-90-7 |
| 158 | 2,3,3',4,4',6 | 74472-42-7 |
| 159 | 2,3,3',4,5,5' | 39635-35-3 |
| 160 | 2,3,3',4,5,6 | 41411-62-5 |
| 161 | 2,3,3',4,5',6 | 74472-43-8 |
| 162 | 2,3,3',4',5,5' | 39635-34-2 |
| 163 | 2,3,3',4',5,6 | 74472-44-9 |
| 164 | 2,3,3',4',5',6 | 74472-45-0 |
| 165 | 2,3,3',5,5',6 | 74472-46-1 |
| 166 | 2,3,4,4',5,6 | 41411-63-6 |
| 167 | 2,3',4,4',5,5' | 52663-72-6 |
| 168 | 2,3',4,4',5',6 | 59291-65-5 |
| 169 | 3,3',4,4',5,5' | 32774-16-6 |
| | | 28655-71-2 |
| Heptachlorobiphenyl | | |
| 170 | 2,2',3,3',4,4',5 | 35065-30-6 |
| 171 | 2,2',3,3',4,4',6 | 52663-71-5 |
| 172 | 2,2',3,3',4,5,5' | 52663-74-8 |
| 173 | 2,2',3,3',4,5,6 | 68194-16-1 |
| 174 | 2,2',3,3',4,5,6' | 38411-25-5 |
| 175 | 2,2',3,3',4,5',6 | 40186-70-7 |
| 176 | 2,2',3,3',4,6,6' | 52663-65-7 |
| 177 | 2,2',3,3',4',5,6 | 52663-70-4 |
| 178 | 2,2',3,3',5,5',6, | 52663-67-9 |
| 179 | 2,2',3,3',5,6,6' | 52663-64-6 |
| 180 | 2,2',3,4,4',5,5' | 35065-29-3 |
| 181 | 2,2',3,4,4',5,6 | 74472-47-2 |
| 182 | 2,2',3,4,4',5,6' | 60145-23-5 |
| 183 | 2,2',3,4,4',5',6 | 52663-69-1 |

3. CHEMICAL AND PHYSICAL INFORMATIONS

Table 3-2. Chemical Identity of Polychlorinated Biphenyl Congeners (continued)

| IUPAC No. ^a | Compound/substituents | CAS No. ^b |
|------------------------|--------------------------|----------------------|
| 184 | 2,2',3,4,4',6,6' | 74472-48-3 |
| 185 | 2,2',3,4,5,5',6 | 52712-05-7 |
| 186 | 2,2',3,4,5,6,6' | 74472-49-4 |
| 187 | 2,2',3,4',5,5',6 | 52663-68-0 |
| 188 | 2,2',3,4',5,6,6' | 74487-85-7 |
| 189 | 2,3,3',4,4',5,5' | 39635-31-9 |
| 190 | 2,3,3',4,4',5,6 | 41411-64-7 |
| 191 | 2,3,3',4,4',5',6 | 74472-50-7 |
| 192 | 2,3,3',4,5,5',6 | 74472-51-8 |
| 193 | 2,3,3',4',5,5',6 | 69782-91-8 |
| Octachlorobiphenyl | | 31472-83-0 |
| 194 | 2,2',3,3',4,4',5,5' | 35694-08-7 |
| 195 | 2,2',3,3',4,4',5,6 | 52663-78-2 |
| 196 | 2,2',3,3',4,4',5,6' | 42740-50-1 |
| 197 | 2,2',3,3',4,4',6,6' | 33091-17-7 |
| 198 | 2,2',3,3',4,5,5',6 | 68194-17-2 |
| 199 | 2,2',3,3',4,5,5',6' | 52663-75-9 |
| 200 | 2,2',3,3',4,5,6,6' | 52663-73-7 |
| 201 | 2,2',3,3',4,5',6,6' | 40186-71-8 |
| 202 | 2,2',3,3',5,5',6,6' | 2136-99-4 |
| 203 | 2,2',3,4,4',5,5',6 | 52663-76-0 |
| 204 | 2,2',3,4,4',5,6,6' | 74472-52-9 |
| 205 | 2,3,3',4,4',5,5',6 | 74472-53-0 |
| Nonachlorobiphenyl | | 53742-07-7 |
| 206 | 2,2',3,3',4,4',5,5',6 | 40186-72-9 |
| 207 | 2,2',3,3',4,4',5,6,6' | 52663-79-3 |
| 208 | 2,2',3,3',4,5,5',6,6' | 52663-77-1 |
| Decachlorobiphenyl | | 2051-24-3 |
| 209 | 2,2',3,3',4,4',5,5',6,6' | 2051-24-3 |

^a Ballschmiter and Zell 1980^b Erickson 1985

3. CHEMICAL AND PHYSICAL INFORMATIONS

Table 3-4. Approximate Weight Percent of Chlorobiphenyl in Some Aroclors

| Empirical Formula | Aroclor 1016 | Aroclor 1221 | Aroclor 1232 ^a | Aroclor 1242 | Aroclor 1248 ^a |
|--|--------------|--------------|---------------------------|--------------|---------------------------|
| C ₁₂ H ₁₀ | Not detected | 10 | Not detected | Not detected | Not detected |
| C ₁₂ H ₉ Cl | 2 | 50 | 26 | 1 | Not detected |
| C ₁₂ H ₈ Cl ₂ | 19 | 35 | 29 | 13 | 1 |
| C ₁₂ H ₇ Cl ₃ | 57 | 4 | 24 | 45 | 2 |
| C ₁₂ H ₆ Cl ₄ | 22 | 1 | 15 | 31 | 49 |
| C ₁₂ H ₅ Cl ₅ | Not detected | Not detected | Not detected | 10 | 27 |
| C ₁₂ H ₄ Cl ₆ | Not detected | Not detected | Not detected | Not detected | 2 |
| C ₁₂ H ₃ Cl ₇ | Not detected | Not detected | Not detected | Not detected | Not detected |
| C ₁₂ H ₂ Cl ₈ | Not detected | Not detected | Not detected | Not detected | Not detected |
| C ₁₂ H ₁ Cl ₉ | Not detected | Not detected | Not detected | Not detected | Not detected |
| Average molecular mass | 257.9 | 200.7 | 232.2 | 266.5 | 299.5 |

| Empirical formula | Aroclor 1254 ^a | Aroclor 1260 | Aroclor 1262 | Aroclor 1268 |
|--|---------------------------|--------------|--------------|--------------|
| C ₁₂ H ₁₀ | Not detected | Not detected | No data | No data |
| C ₁₂ H ₉ Cl | Not detected | Not detected | No data | No data |
| C ₁₂ H ₈ Cl ₂ | Not detected | Not detected | No data | No data |
| C ₁₂ H ₇ Cl ₃ | 1 | Not detected | No data | No data |
| C ₁₂ H ₆ Cl ₄ | 15 | Not detected | No data | No data |
| C ₁₂ H ₅ Cl ₅ | 53 | 12 | No data | No data |
| C ₁₂ H ₄ Cl ₆ | 26 | 42 | No data | No data |
| C ₁₂ H ₃ Cl ₇ | 4 | 38 | No data | No data |
| C ₁₂ H ₂ Cl ₈ | Not detected | 7 | No data | No data |
| C ₁₂ H ₁ Cl ₉ | Not detected | 1 | No data | No data |
| Average molecular mass | 328.4 | 375.7 | 389 | 453 |

^a Compounds that contributed the remaining composition were not given.

Source: derived from Ballschmiter et al. 1989

Table 3-5. PCB Congener Compositions (in mol%) in Aroclors^a

| IUPAC No. | Chlorine substitution pattern | Aroclor | | | | |
|--------------|-------------------------------------|---------|-------|----------------|-------|-------|
| | | 1242 | 1016 | 1248 | 1254 | 1260 |
| | BP | 0.01 | 0.50 | — ^b | — | — |
| 1 | 2 | 0.68 | 0.80 | — | — | — |
| 2 | 3 | 0.04 | 0.10 | — | — | — |
| 3 | 4 | 0.22 | 1.00 | — | — | — |
| 4 | 2,2' | 3.99 | 4.36 | 0.25 | — | — |
| 6 | 2,3' | 1.24 | 1.37 | 0.69 | 0.07 | — |
| 7 | 2,4 | 1.04 | 1.16 | — | — | — |
| 8 | 2,4' | 8.97 | 10.30 | 0.18 | — | — |
| 9 | 2,5 | 0.31 | 0.34 | trace | — | — |
| 10 | 2,6 | 0.13 | 0.20 | — | — | — |
| 12 | 3,4 | 0.09 | 0.11 | — | — | — |
| 13 | 3,4' | 0.12 | 0.12 | — | — | — |
| 14 | 3,5 | 0.35 | 0.37 | — | — | — |
| 15 | 4,4' | 0.99 | 1.07 | — | — | — |
| 16 | 2,3,2' | 3.25 | 3.50 | 0.84 | — | — |
| 17 | 2,4,2' | 2.92 | 3.14 | 0.19 | — | — |
| 18 | 2,5,2' | 9.36 | 10.87 | 9.95 | 0.07 | — |
| 19 | 2,6,2' | 0.97 | 1.08 | — | — | — |
| 20 | 2,3,3' | 3.64 | 3.99 | — | — | — |
| 22 | 2,3,4' | 2.64 | 2.80 | 1.24 | trace | trace |
| 25 | 2,4,3' | 1.68 | 1.79 | — | — | — |
| 26 | 2,5,3' | 0.55 | 0.62 | 0.75 | — | — |
| 27 | 2,6,3' | 0.54 | 0.58 | — | — | — |
| 28 | 2,4,4' | 13.30 | 14.48 | trace | — | — |
| 31 | 2,5,4' | 4.53 | 4.72 | 9.31 | 0.72 | — |
| 32 | 2,6,4' | 2.15 | 2.31 | 1.46 | — | — |
| 33 | 3,4,2' | 2.83 | 3.08 | — | — | — |
| 35 | 3,4,3' | 0.66 | 0.38 | — | — | — |
| 37 | 3,4,4' | 1.62 | 1.89 | 1.28 | 0.20 | 0.09 |
| 39 | 3,5,4' | 1.03 | 1.08 | — | — | — |
| 40 | 2,3,2',3' | 0.15 | 0.18 | 1.12 | 0.26 | 0.04 |
| 41 | 2,3,4,2' | 1.67 | 2.00 | — | — | — |
| 42 | 2,3,2',4' | — | — | 7.05 | 2.18 | 0.66 |
| 43 | 2,3,5,2' | 0.44 | 0.47 | — | — | — |
| 44 | 2,3,2',5' | 1.06 | 1.14 | — | — | — |

3. CHEMICAL AND PHYSICAL INFORMATIONS

Table 3-5. PCB Congener Compositions (in mol%) in Aroclors^a (continued)

| IUPAC No. | Chlorine substitution pattern | Aroclor | | | | |
|--------------|-------------------------------------|---------|-------|-------|-------|-------|
| | | 1242 | 1016 | 1248 | 1254 | 1260 |
| 45 | 2,3,6,2' | 0.90 | 1.00 | 5.73 | 0.15 | — |
| 46 | 2,3,2',6' | 0.31 | 0.33 | — | — | — |
| 47 | 2,4,2',4' | 1.65 | 1.8 | 3.18 | 0.52 | 0.88 |
| 48 | 2,4,5,2' | 1.33 | 1.41 | — | — | — |
| 49 | 2,4,2',5' | 3.28 | 3.48 | 3.81 | 1.63 | 0.44 |
| 52 | 2,5,2',5' | 4.08 | 4.35 | 8.36 | 4.36 | 1.91 |
| 53 | 2,5,2',6' | 0.97 | 1.07 | 6.30 | 0.13 | — |
| 54 | 2,6,2',6' | 0.17 | 0.19 | — | — | — |
| 55 | 2,3,4,3' | — | — | 0.11 | 0.43 | 0.12 |
| 56 | 2,3,3',4' | 0.60 | trace | 0.18 | 0.03 | — |
| 60 | 2,3,4,4' | 0.21 | — | — | — | — |
| 66 | 2,4,3',4' | 0.81 | 0.14 | 4.95 | 2.24 | 0.22 |
| 70 | 2,5,3',4' | 1.11 | — | 6.38 | 4.75 | 0.85 |
| 71 | 2,6,3',4' | — | — | 0.65 | — | — |
| 72 | 2,5,3',5' | 0.33 | — | 2.10 | 1.01 | 0.28 |
| 74 | 2,4,5,4' | 2.02 | 1.35 | 0.25 | 0.30 | 0.09 |
| 75 | 2,4,6,4' | 2.18 | 2.40 | — | — | — |
| 76 | 3,4,5,2' | trace | — | trace | 0.18 | 0.01 |
| 77 | 3,4,3',4' | 0.34 | — | 0.47 | 0.12 | 0.04 |
| 78 | 3,4,5,3' | 0.52 | — | — | — | — |
| 79 | 3,4,3',5' | 0.24 | — | trace | 0.23 | 0.04 |
| 80 | 3,5,3',5' | — | — | trace | trace | trace |
| 81 | 3,4,5,4' | 0.28 | — | — | — | — |
| 83 | 2,3,5,2',3' | — | — | trace | 0.32 | 0.09 |
| 84 | 2,3,6,2',3' | 0.38 | 0.01 | 0.71 | 1.72 | 0.69 |
| 85 | 2,3,4,2',4' | 0.40 | — | 0.55 | 2.15 | 0.31 |
| 87 | 2,3,4,2',5' | 0.09 | — | 1.05 | 3.81 | 1.10 |
| 91 | 2,3,6,2',4' | trace | — | 1.78 | 5.00 | 3.22 |
| 92 | 2,3,5,2',5' | 0.12 | 0.20 | 0.63 | 0.21 | — |
| 95 | 2,3,6,2',5' | 0.53 | 0.18 | — | — | — |
| 97 | 2,4,5,2',3' | — | — | 0.78 | 2.59 | 0.63 |
| 98 | 2,4,6,2',3' | 0.13 | 0.04 | — | — | — |
| 99 | 2,4,5,2',4' | 0.55 | — | 2.52 | 6.10 | 0.82 |
| 101 | 2,4,5,2',5' | 0.27 | — | 1.50 | 6.98 | 5.04 |
| 102 | 2,4,5,2',6' | — | — | trace | trace | trace |
| 105 | 2,3,4,3',4' | 0.25 | — | — | — | — |

Table 3-5. PCB Congener Compositions (in mol%) in Aroclors^a (continued)

| IUPAC No. | Chlorine substitution pattern | Aroclor | | | | |
|--------------|-------------------------------------|----------------|-------|----------------|-------|-------|
| | | 1242 | 1016 | 1248 | 1254 | 1260 |
| 106 | 2,3,4,5,3' | — | — | — | 0.40 | 0.06 |
| 108 | 2,3,4,3',5' | 0.46 | 0.16 | 0.02 | 0.55 | 0.14 |
| 110 | 2,3,6,3',4' | — ^c | — | 1.69 | 8.51 | 3.57 |
| 113 | 2,3,6,3',5' | 0.39 | 0.01 | 3.10 | trace | 0.01 |
| 114 | 2,3,4,5,4' | — | — | — | 0.25 | 0.03 |
| 118 | 2,4,5,3',4' | — ^c | — | — ^c | 8.09 | 2.00 |
| 120 | 2,4,5,3',5' | 0.31 | — | trace | 0.15 | 3.01 |
| 121 | 2,4,6,3',5' | 0.92 | — | 4.32 | 3.51 | 0.57 |
| 122 | 3,4,5,2',3' | — | — | trace | 0.76 | 1.88 |
| 123 | 3,4,5,2',4' | 0.36 | — | — | — | — |
| 126 | 3,4,5,3',4' | 0.03 | — | — | 0.16 | 1.59 |
| 127 | 3,4,5,3',5' | 0.05 | — | — | — | — |
| 128 | 2,3,4,2',3',4' | — | — | — | 1.31 | 0.47 |
| 131 | 2,3,4,6,2',3' | — | — | — | 0.14 | 0.01 |
| 132 | 2,3,4,2',3',6' | — | — | trace | 2.00 | 2.77 |
| 133 | 2,3,5,2',3',5' | — | — | 1.13 | 0.03 | 0.06 |
| 134 | 2,3,5,6,2',3' | — | — | 0.11 | 0.38 | 1.01 |
| 135 | 2,3,5,2',3',6' | — | trace | 0.20 | 0.29 | — |
| 136 | 2,3,6,2',3',6' | — | — | 0.20 | 0.34 | 1.12 |
| 138 | 2,3,4,2',4',5' | 0.08 | — | 0.19 | 4.17 | 5.01 |
| 143 | 2,3,4,5,2',6' | 0.07 | — | — | — | — |
| 148 | 2,3,5,2',4',6' | — | — | 0.12 | 0.07 | 0.06R |
| 149 | 2,4,5,2',3',6' | — | — | 0.77 | 3.59 | 9.52 |
| 151 | 2,3,5,6,2',5' | — | — | trace | 0.33 | 0.06 |
| 153 | 2,4,5,2',4',5' | 0.02 | — | 0.13 | 3.32 | 8.22 |
| 154 | 2,4,5,4',6' | — | — | — | 0.14 | — |
| 156 | 2,3,4,5,3',4' | — | — | — | — | 0.41 |
| 157 | 2,3,4,3',4',5' | — | — | — | 0.18 | 0.03 |
| 158 | 2,3,4,6,3',4' | — | — | — | 0.46 | 0.18 |
| 159 | 2,4,5,2',3',5' | — | — | — | 0.75 | 1.48 |
| 163 | 2,3,5,6,3',4' | — | — | — | — | trace |
| 167 | 2,4,5,3',4',5' | — | — | — | 0.21 | 0.17 |
| 168 | 2,4,6,3',4',5' | — | — | 0.56 | 4.23 | 0.59 |
| 170 | 2,3,4,5,2',3',4' | — | — | — | 0.43 | 0.62 |
| 171 | 2,3,4,6,2',3',4' | — | — | — | 0.30 | 4.31 |
| 174 | 2,3,4,5,2',3',6' | — | — | — | trace | 0.09 |

3. CHEMICAL AND PHYSICAL INFORMATIONS

Table 3-5. PCB Congener Compositions (in mol%) in Aroclors^a (continued)

| IUPAC No. | Chlorine substitution pattern | Aroclor | | | | |
|-----------|-------------------------------|---------|------|-------|-------|-------|
| | | 1242 | 1016 | 1248 | 1254 | 1260 |
| 176 | 2,3,4,6,2',3',6' | — | — | 0.09 | trace | 0.57 |
| 177 | 2,3,5,6,2',3',4' | — | — | — | — | trace |
| 179 | 2,3,5,6,2',3',6' | — | — | — | 0.56 | 0.83 |
| 180 | 2,3,4,5,2',4',5' | — | — | — | 0.76 | 7.20 |
| 181 | 2,3,4,5,6,2',4' | — | — | — | 0.28 | 2.72 |
| 182 | 2,3,4,5,2',4',6' | — | — | — | trace | 0.47 |
| 183 | 2,3,4,6,2',4',5' | — | — | — | 1.16 | 2.58 |
| 185 | 2,3,4,5,6,2',5' | — | — | — | 1.11 | 5.65 |
| 186 | 2,3,4,5,6,2',6' | — | — | trace | trace | 0.37 |
| 187 | 2,3,5,6,2',4',5' | — | — | — | 0.48 | 1.12 |
| 189 | 2,3,4,5,3',4',5' | — | — | — | — | 0.13 |
| 190 | 2,3,4,5,6,3',4' | — | — | — | — | 0.02 |
| 192 | 2,3,4,5,6,3',5' | — | — | — | 0.20 | 0.97 |
| 193 | 2,3,5,6,3',4',5' | — | — | — | 2.30 | — |
| 194 | 2,3,4,5,2',3',4',5' | — | — | — | — | 2.21 |
| 195 | 2,3,4,5,6,2',3',4' | — | — | — | — | trace |
| 196 | 2,3,4,5,2',3',4',6' | — | — | — | — | 0.79 |
| 197 | 2,3,4,6,2',3',4',6' | — | — | — | — | 0.30 |
| 198 | 2,3,4,5,6,2',3',5' | — | — | — | 1.00 | 0.15 |
| 199 | 2,3,4,5,6,2',3',6' | — | — | — | — | 0.38 |
| 200 | 2,3,4,6,2',3',5',6' | — | — | — | trace | 0.15 |
| 201 | 2,3,4,5,2',3',5',6' | — | — | — | — | 1.54 |
| 202 | 2,3,5,6,2',3',5',6' | — | — | — | — | 0.31 |
| 203 | 2,3,4,5,6,2',4',5' | — | — | — | — | 0.08 |
| 204 | 2,3,4,5,6,2',4',6' | — | — | — | trace | 0.13 |
| 205 | 2,3,4,5,6,3',4',5' | — | — | — | — | 0.01 |
| 206 | 2,3,4,5,6,2',3',4',5' | — | — | — | — | 0.51 |
| 207 | 2,3,4,5,6,2',3',4',6' | — | — | — | — | 1.15 |
| 208 | 2,3,4,5,6,2',3',5',6' | — | — | — | — | 0.18 |

^a WHO 1993; the congener 169 (3,4,5,3',4',5') was not found in Aroclors; Albro and Parker 1979; Albro et al. 1981

^b — Not detected

^c Presence of these congeners has been reported by other investigators (see Section 3.2)

APPENDIX I

PCB CONGENER RESULTS FOR EACH REACH AND ZONE

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|-----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| Lake Winnebago | 3 | PCB Congener 20/33 | 3 | 2 | 2.9815 | 2.9815 | 10.35 | 28.81% |
| Lake Winnebago | 4 | PCB Congener 53 | 3 | 2 | 1.4685 | 1.4685 | | 14.19% |
| Lake Winnebago | 7 | PCB Congener 170 | 3 | 1 | 2.8000 | 2.8 | | 27.05% |
| Lake Winnebago | 8 | PCB Congener 194 | 3 | 1 | 3.1000 | 3.1 | | 29.95% |
| LLBdM | 2 | PCB Congener 4/10 | 4 | 2 | 30.5000 | 1,462.1401 | 15,218.45 | 9.61% |
| LLBdM | 2 | PCB Congener 5 | 6 | 1 | 4.3000 | | | |
| LLBdM | 2 | PCB Congener 6 | 33 | 30 | 192.6767 | | | |
| LLBdM | 2 | PCB Congener 7 | 26 | 24 | 16.4775 | | | |
| LLBdM | 2 | PCB Congener 7/9 | 4 | 2 | 11.1000 | | | |
| LLBdM | 2 | PCB Congener 8 | 6 | 6 | 98.0167 | | | |
| LLBdM | 2 | PCB Congener 8/5 | 29 | 29 | 1,014.6276 | | | |
| LLBdM | 2 | PCB Congener 12 | 6 | 3 | 12.8667 | | | |
| LLBdM | 2 | PCB Congener 15 | 2 | 2 | 6.4500 | | | |
| LLBdM | 2 | PCB Congener 15 | 4 | 4 | 75.1250 | | | |
| LLBdM | 3 | PCB Congener 16 | 6 | 6 | 61.1333 | 5,119.7345 | | 33.64% |
| LLBdM | 3 | PCB Congener 16/32 | 29 | 29 | 472.6828 | | | |
| LLBdM | 3 | PCB Congener 17 | 31 | 31 | 382.4806 | | | |
| LLBdM | 3 | PCB Congener 17 | 4 | 4 | 75.1250 | | | |
| LLBdM | 3 | PCB Congener 18 | 35 | 35 | 282.4314 | | | |
| LLBdM | 3 | PCB Congener 19 | 27 | 25 | 37.8344 | | | |
| LLBdM | 3 | PCB Congener 20 | 2 | 2 | 1.5000 | | | |
| LLBdM | 3 | PCB Congener 20/33 | 4 | 4 | 60.1325 | | | |
| LLBdM | 3 | PCB Congener 22 | 35 | 34 | 414.8941 | | | |
| LLBdM | 3 | PCB Congener 24/27 | 27 | 26 | 70.7658 | | | |
| LLBdM | 3 | PCB Congener 25 | 6 | 6 | 35.0333 | | | |
| LLBdM | 3 | PCB Congener 26 | 35 | 35 | 330.3137 | | | |
| LLBdM | 3 | PCB Congener 27 | 6 | 6 | 9.5500 | | | |
| LLBdM | 3 | PCB Congener 28 | 6 | 6 | 157.1667 | | | |
| LLBdM | 3 | PCB Congener 28/31 | 29 | 29 | 2,030.8690 | | | |
| LLBdM | 3 | PCB Congener 31 | 6 | 6 | 146.3333 | | | |
| LLBdM | 3 | PCB Congener 33 | 31 | 29 | 355.4586 | | | |
| LLBdM | 3 | PCB Congener 37 | 6 | 6 | 29.9833 | | | |
| LLBdM | 3 | PCB Congener 37 | 32 | 32 | 166.0466 | | | |
| LLBdM | 4 | PCB Congener 40 | 35 | 33 | 72.4085 | 3,927.6891 | | 25.81% |
| LLBdM | 4 | PCB Congener 41 | 6 | 6 | 31.9500 | | | |
| LLBdM | 4 | PCB Congener 41/64/71 | 29 | 28 | 371.6679 | | | |
| LLBdM | 4 | PCB Congener 42 | 2 | 2 | 7.0000 | | | |
| LLBdM | 4 | PCB Congener 42 | 32 | 32 | 166.0466 | | | |
| LLBdM | 4 | PCB Congener 44 | 35 | 35 | 336.6543 | | | |
| LLBdM | 4 | PCB Congener 45 | 34 | 27 | 61.9519 | | | |
| LLBdM | 4 | PCB Congener 46 | 28 | 27 | 32.5119 | | | |
| LLBdM | 4 | PCB Congener 47 | 2 | 2 | 14.5000 | | | |
| LLBdM | 4 | PCB Congener 47/48 | 29 | 29 | 337.4307 | | | |
| LLBdM | 4 | PCB Congener 47/75 | 4 | 4 | 42.1750 | | | |
| LLBdM | 4 | PCB Congener 49 | 35 | 31 | 341.9971 | | | |
| LLBdM | 4 | PCB Congener 52 | 35 | 35 | 357.1523 | | | |
| LLBdM | 4 | PCB Congener 53 | 2 | 2 | 5.8000 | | | |
| LLBdM | 4 | PCB Congener 53 | 4 | 4 | 29.6175 | | | |
| LLBdM | 4 | PCB Congener 56 | 2 | 2 | 6.5500 | | | |
| LLBdM | 4 | PCB Congener 56/60 | 28 | 28 | 218.5429 | | | |
| LLBdM | 4 | PCB Congener 56/60 | 4 | 4 | 30.8200 | | | |
| LLBdM | 4 | PCB Congener 59 | 6 | 5 | 4.8600 | | | |
| LLBdM | 4 | PCB Congener 66 | 2 | 2 | 23.5000 | | | |
| LLBdM | 4 | PCB Congener 66 | 33 | 33 | 433.3045 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| LLBdM | 4 | PCB Congener 70 | 6 | 6 | 46.8333 | | | |
| LLBdM | 4 | PCB Congener 70/76 | 29 | 29 | 466.2069 | | | |
| LLBdM | 4 | PCB Congener 74 | 35 | 35 | 150.9143 | | | |
| LLBdM | 4 | PCB Congener 77 | 18 | 14 | 14.0071 | | | |
| LLBdM | 4 | PCB Congener 77 | 33 | 33 | 318.7452 | | | |
| LLBdM | 4 | PCB Congener 81 | 16 | 10 | 0.1359 | | | |
| LLBdM | 4 | PCB Congener 81 | 4 | 4 | 4.4055 | | | |
| LLBdM | 5 | PCB Congener 82 | 32 | 27 | 50.5307 | 2,428.7749 | | 15.96% |
| LLBdM | 5 | PCB Congener 83 | 6 | 1 | 4.9000 | | | |
| LLBdM | 5 | PCB Congener 84 | 6 | 6 | 12.1333 | | | |
| LLBdM | 5 | PCB Congener 84/92 | 28 | 28 | 331.4357 | | | |
| LLBdM | 5 | PCB Congener 85 | 33 | 27 | 89.3789 | | | |
| LLBdM | 5 | PCB Congener 87 | 29 | 29 | 200.2893 | | | |
| LLBdM | 5 | PCB Congener 87/115 | 4 | 4 | 8.9445 | | | |
| LLBdM | 5 | PCB Congener 91 | 33 | 32 | 87.6006 | | | |
| LLBdM | 5 | PCB Congener 92 | 2 | 2 | 5.0500 | | | |
| LLBdM | 5 | PCB Congener 92 | 4 | 4 | 15.1800 | | | |
| LLBdM | 5 | PCB Congener 95 | 2 | 2 | 25.5000 | | | |
| LLBdM | 5 | PCB Congener 95 | 33 | 33 | 433.3045 | | | |
| LLBdM | 5 | PCB Congener 97 | 34 | 33 | 110.7209 | | | |
| LLBdM | 5 | PCB Congener 99 | 34 | 34 | 153.8388 | | | |
| LLBdM | 5 | PCB Congener 101 | 34 | 34 | 271.5029 | | | |
| LLBdM | 5 | PCB Congener 105 | 18 | 16 | 6.7875 | | | |
| LLBdM | 5 | PCB Congener 107 | 6 | 2 | 3.4500 | | | |
| LLBdM | 5 | PCB Congener 110 | 2 | 2 | 31.0000 | | | |
| LLBdM | 5 | PCB Congener 110 | 33 | 33 | 318.7452 | | | |
| LLBdM | 5 | PCB Congener 114 | 16 | 13 | 1.5623 | | | |
| LLBdM | 5 | PCB Congener 118 | 46 | 46 | 257.0804 | | | |
| LLBdM | 5 | PCB Congener 123 | 14 | 2 | 0.7400 | | | |
| LLBdM | 5 | PCB Congener 123 | 4 | 4 | 7.5500 | | | |
| LLBdM | 5 | PCB Congener 126 | 18 | 8 | 0.0971 | | | |
| LLBdM | 5 | PCB Congener 126 | 4 | 1 | 1.4520 | | | |
| LLBdM | 6 | PCB Congener 128 | 6 | 6 | 5.3333 | 1,501.9168 | | 9.87% |
| LLBdM | 6 | PCB Congener 129 | 4 | 1 | 1.4520 | | | |
| LLBdM | 6 | PCB Congener 132/153 | 29 | 29 | 426.6759 | | | |
| LLBdM | 6 | PCB Congener 135 | 6 | 3 | 3.0333 | | | |
| LLBdM | 6 | PCB Congener 135/144 | 28 | 28 | 52.3596 | | | |
| LLBdM | 6 | PCB Congener 136 | 32 | 27 | 33.1633 | | | |
| LLBdM | 6 | PCB Congener 137/176 | 19 | 3 | 7.3833 | | | |
| LLBdM | 6 | PCB Congener 138 | 6 | 6 | 22.3167 | | | |
| LLBdM | 6 | PCB Congener 141 | 31 | 20 | 59.2355 | | | |
| LLBdM | 6 | PCB Congener 146 | 24 | 22 | 107.8409 | | | |
| LLBdM | 6 | PCB Congener 149 | 30 | 30 | 198.9993 | | | |
| LLBdM | 6 | PCB Congener 149 | 4 | 4 | 7.5500 | | | |
| LLBdM | 6 | PCB Congener 151 | 33 | 29 | 52.8990 | | | |
| LLBdM | 6 | PCB Congener 153 | 6 | 6 | 18.9667 | | | |
| LLBdM | 6 | PCB Congener 156 | 18 | 15 | 2.0360 | | | |
| LLBdM | 6 | PCB Congener 157 | 14 | 10 | 0.4483 | | | |
| LLBdM | 6 | PCB Congener 163/138 | 28 | 28 | 490.5786 | | | |
| LLBdM | 6 | PCB Congener 167 | 18 | 12 | 2.0450 | | | |
| LLBdM | 6 | PCB Congener 168 | 6 | 3 | 9.6000 | | | |
| LLBdM | 7 | PCB Congener 170 | 18 | 14 | 4.6121 | 448.7165 | | 2.95% |
| LLBdM | 7 | PCB Congener 170/190 | 26 | 26 | 116.9692 | | | |
| LLBdM | 7 | PCB Congener 202/171 | 23 | 22 | 13.6455 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| LLBdM | 7 | PCB Congener 172/197 | 13 | 13 | 11.4177 | | | |
| LLBdM | 7 | PCB Congener 174 | 32 | 26 | 44.9985 | | | |
| LLBdM | 7 | PCB Congener 137/176 | 19 | 3 | 7.3833 | | | |
| LLBdM | 7 | PCB Congener 177 | 31 | 26 | 34.6904 | | | |
| LLBdM | 7 | PCB Congener 178 | 21 | 19 | 22.3979 | | | |
| LLBdM | 7 | PCB Congener 178 | 4 | 1 | 1.4520 | | | |
| LLBdM | 7 | PCB Congener 180 | 46 | 45 | 67.6460 | | | |
| LLBdM | 7 | PCB Congener 182/187 | 27 | 26 | 58.0008 | | | |
| LLBdM | 7 | PCB Congener 183 | 31 | 24 | 38.6992 | | | |
| LLBdM | 7 | PCB Congener 185 | 16 | 9 | 13.0444 | | | |
| LLBdM | 7 | PCB Congener 187 | 6 | 3 | 13.6000 | | | |
| LLBdM | 7 | PCB Congener 189 | 18 | 8 | 0.1595 | | | |
| LLBdM | 8 | PCB Congener 194 | 31 | 27 | 25.6481 | 226.5845 | | 1.49% |
| LLBdM | 8 | PCB Congener 195 | 6 | 1 | 4.9000 | | | |
| LLBdM | 8 | PCB Congener 195 | 24 | 24 | 18.8104 | | | |
| LLBdM | 8 | PCB Congener 196 | 6 | 2 | 32.0000 | | | |
| LLBdM | 8 | PCB Congener 196/203 | 23 | 23 | 61.7609 | | | |
| LLBdM | 8 | PCB Congener 172/197 | 13 | 13 | 11.4177 | | | |
| LLBdM | 8 | PCB Congener 199 | 16 | 8 | 2.6913 | | | |
| LLBdM | 8 | PCB Congener 201 | 32 | 29 | 45.7107 | | | |
| LLBdM | 8 | PCB Congener 202 | 6 | 1 | 10.0000 | | | |
| LLBdM | 8 | PCB Congener 202/171 | 23 | 22 | 13.6455 | | | |
| LLBdM | 9 | PCB Congener 206 | 31 | 27 | 37.5815 | 89.8919 | | 0.59% |
| LLBdM | 9 | PCB Congener 208 | 6 | 2 | 33.5000 | | | |
| LLBdM | 9 | PCB Congener 208 | 24 | 24 | 18.8104 | | | |
| LLBdM | 10 | PCB Congener 209 | 6 | 1 | 13.0000 | 13.0000 | | 0.09% |
| App. - LR | 1 | PCB Congener 1 | 8 | 2 | 4.5000 | 4.5000 | 16,599.37 | 0.03% |
| App. - LR | 2 | PCB Congener 4/10 | 7 | 4 | 176.0750 | 1,975.7597 | | 11.90% |
| App. - LR | 2 | PCB Congener 6 | 15 | 15 | 195.2133 | | | |
| App. - LR | 2 | PCB Congener 7 | 8 | 8 | 2.1613 | | | |
| App. - LR | 2 | PCB Congener 7/9 | 7 | 6 | 54.4167 | | | |
| App. - LR | 2 | PCB Congener 8 | 7 | 7 | 837.4143 | | | |
| App. - LR | 2 | PCB Congener 8/5 | 8 | 8 | 137.3000 | | | |
| App. - LR | 2 | PCB Congener 12 | 8 | 6 | 23.8167 | | | |
| App. - LR | 2 | PCB Congener 15 | 8 | 8 | 549.3625 | | | |
| App. - LR | 3 | PCB Congener 16 | 8 | 8 | 938.6000 | 8,591.7803 | | 51.76% |
| App. - LR | 3 | PCB Congener 16/32 | 8 | 8 | 55.9125 | | | |
| App. - LR | 3 | PCB Congener 17 | 8 | 8 | 40.1750 | | | |
| App. - LR | 3 | PCB Congener 17 | 8 | 8 | 549.3625 | | | |
| App. - LR | 3 | PCB Congener 18 | 16 | 16 | 843.1875 | | | |
| App. - LR | 3 | PCB Congener 19 | 14 | 12 | 79.9925 | | | |
| App. - LR | 3 | PCB Congener 20/33 | 7 | 7 | 810.3841 | | | |
| App. - LR | 3 | PCB Congener 22 | 15 | 15 | 355.5600 | | | |
| App. - LR | 3 | PCB Congener 24/27 | 8 | 8 | 6.7125 | | | |
| App. - LR | 3 | PCB Congener 25 | 7 | 7 | 287.1857 | | | |
| App. - LR | 3 | PCB Congener 26 | 15 | 15 | 236.9333 | | | |
| App. - LR | 3 | PCB Congener 27 | 7 | 6 | 150.5333 | | | |
| App. - LR | 3 | PCB Congener 28 | 7 | 7 | 1,975.5714 | | | |
| App. - LR | 3 | PCB Congener 28/31 | 8 | 8 | 328.7500 | | | |
| App. - LR | 3 | PCB Congener 31 | 7 | 7 | 1,642.2857 | | | |
| App. - LR | 3 | PCB Congener 33 | 8 | 8 | 103.0875 | | | |
| App. - LR | 3 | PCB Congener 37 | 4 | 4 | 7.5000 | | | |
| App. - LR | 3 | PCB Congener 37 | 15 | 15 | 180.0467 | | | |
| App. - LR | 4 | PCB Congener 40 | 15 | 15 | 124.9600 | 4,755.9229 | | 28.65% |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|-----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| App. - LR | 4 | PCB Congener 41 | 7 | 7 | 447.6571 | | | |
| App. - LR | 4 | PCB Congener 41/64/71 | 8 | 8 | 63.4375 | | | |
| App. - LR | 4 | PCB Congener 42 | 15 | 15 | 180.0467 | | | |
| App. - LR | 4 | PCB Congener 44 | 15 | 15 | 380.5333 | | | |
| App. - LR | 4 | PCB Congener 45 | 15 | 11 | 130.9382 | | | |
| App. - LR | 4 | PCB Congener 46 | 7 | 7 | 5.6000 | | | |
| App. - LR | 4 | PCB Congener 47/48 | 8 | 8 | 41.7250 | | | |
| App. - LR | 4 | PCB Congener 47/75 | 7 | 7 | 252.3714 | | | |
| App. - LR | 4 | PCB Congener 49 | 15 | 11 | 468.1818 | | | |
| App. - LR | 4 | PCB Congener 52 | 15 | 15 | 447.1200 | | | |
| App. - LR | 4 | PCB Congener 53 | 7 | 7 | 399.1444 | | | |
| App. - LR | 4 | PCB Congener 56/60 | 8 | 8 | 85.1375 | | | |
| App. - LR | 4 | PCB Congener 56/60 | 7 | 7 | 282.4624 | | | |
| App. - LR | 4 | PCB Congener 59 | 7 | 5 | 118.5200 | | | |
| App. - LR | 4 | PCB Congener 66 | 15 | 15 | 254.1600 | | | |
| App. - LR | 4 | PCB Congener 70 | 7 | 7 | 658.0000 | | | |
| App. - LR | 4 | PCB Congener 70/76 | 8 | 8 | 103.3125 | | | |
| App. - LR | 4 | PCB Congener 74 | 15 | 15 | 209.3333 | | | |
| App. - LR | 4 | PCB Congener 77 | 9 | 5 | 11.1740 | | | |
| App. - LR | 4 | PCB Congener 77 | 15 | 15 | 79.2200 | | | |
| App. - LR | 4 | PCB Congener 81 | 9 | 5 | 0.1694 | | | |
| App. - LR | 4 | PCB Congener 81 | 7 | 5 | 12.7182 | | | |
| App. - LR | 5 | PCB Congener 82 | 15 | 13 | 39.0723 | 976.3604 | | 5.88% |
| App. - LR | 5 | PCB Congener 84 | 7 | 6 | 85.0500 | | | |
| App. - LR | 5 | PCB Congener 84/92 | 8 | 8 | 20.4875 | | | |
| App. - LR | 5 | PCB Congener 85 | 14 | 10 | 18.1600 | | | |
| App. - LR | 5 | PCB Congener 87 | 8 | 8 | 12.5250 | | | |
| App. - LR | 5 | PCB Congener 87/115 | 7 | 5 | 25.8218 | | | |
| App. - LR | 5 | PCB Congener 91 | 15 | 9 | 22.8333 | | | |
| App. - LR | 5 | PCB Congener 92 | 7 | 7 | 139.1233 | | | |
| App. - LR | 5 | PCB Congener 95 | 15 | 15 | 254.1600 | | | |
| App. - LR | 5 | PCB Congener 97 | 15 | 14 | 35.3214 | | | |
| App. - LR | 5 | PCB Congener 99 | 15 | 11 | 60.4364 | | | |
| App. - LR | 5 | PCB Congener 101 | 16 | 16 | 92.0000 | | | |
| App. - LR | 5 | PCB Congener 105 | 13 | 9 | 18.4489 | | | |
| App. - LR | 5 | PCB Congener 110 | 15 | 15 | 79.2200 | | | |
| App. - LR | 5 | PCB Congener 114 | 13 | 5 | 0.2012 | | | |
| App. - LR | 5 | PCB Congener 118 | 21 | 21 | 54.2305 | | | |
| App. - LR | 5 | PCB Congener 123 | 8 | 7 | 19.2179 | | | |
| App. - LR | 5 | PCB Congener 126 | 9 | 2 | 0.0510 | | | |
| App. - LR | 6 | PCB Congener 128 | 8 | 3 | 1.4733 | 221.7616 | | 1.34% |
| App. - LR | 6 | PCB Congener 132/153 | 8 | 8 | 23.1375 | | | |
| App. - LR | 6 | PCB Congener 135 | 8 | 3 | 1.2433 | | | |
| App. - LR | 6 | PCB Congener 135/144 | 8 | 8 | 2.3788 | | | |
| App. - LR | 6 | PCB Congener 136 | 14 | 5 | 1.3240 | | | |
| App. - LR | 6 | PCB Congener 137/176 | 1 | 1 | 0.3450 | | | |
| App. - LR | 6 | PCB Congener 138 | 8 | 8 | 69.3125 | | | |
| App. - LR | 6 | PCB Congener 141 | 14 | 5 | 2.2600 | | | |
| App. - LR | 6 | PCB Congener 146 | 6 | 3 | 7.1333 | | | |
| App. - LR | 6 | PCB Congener 149 | 8 | 8 | 10.3000 | | | |
| App. - LR | 6 | PCB Congener 149 | 8 | 7 | 19.2179 | | | |
| App. - LR | 6 | PCB Congener 151 | 16 | 8 | 3.1338 | | | |
| App. - LR | 6 | PCB Congener 153 | 8 | 8 | 58.1900 | | | |
| App. - LR | 6 | PCB Congener 156 | 13 | 7 | 0.6941 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| App. - LR | 6 | PCB Congener 157 | 5 | 5 | 0.1190 | | | |
| App. - LR | 6 | PCB Congener 163/138 | 8 | 8 | 18.9625 | | | |
| App. - LR | 6 | PCB Congener 167 | 13 | 5 | 0.7366 | | | |
| App. - LR | 6 | PCB Congener 168 | 8 | 1 | 1.8000 | | | |
| App. - LR | 7 | PCB Congener 170 | 13 | 6 | 2.0150 | 50.0656 | | 0.30% |
| App. - LR | 7 | PCB Congener 170/190 | 8 | 6 | 5.7333 | | | |
| App. - LR | 7 | PCB Congener 202/171 | 6 | 6 | 0.9792 | | | |
| App. - LR | 7 | PCB Congener 174 | 16 | 8 | 3.6463 | | | |
| App. - LR | 7 | PCB Congener 137/176 | 1 | 1 | 0.3450 | | | |
| App. - LR | 7 | PCB Congener 177 | 16 | 9 | 2.5989 | | | |
| App. - LR | 7 | PCB Congener 178 | 3 | 3 | 1.8667 | | | |
| App. - LR | 7 | PCB Congener 180 | 21 | 18 | 18.0544 | | | |
| App. - LR | 7 | PCB Congener 182/187 | 8 | 8 | 5.2125 | | | |
| App. - LR | 7 | PCB Congener 183 | 14 | 6 | 2.9467 | | | |
| App. - LR | 7 | PCB Congener 185 | 9 | 1 | 2.2000 | | | |
| App. - LR | 7 | PCB Congener 187 | 8 | 1 | 1.1000 | | | |
| App. - LR | 7 | PCB Congener 189 | 13 | 3 | 0.0677 | | | |
| App. - LR | 7 | PCB Congener 192 | 4 | 1 | 3.3000 | | | |
| App. - LR | 8 | PCB Congener 194 | 15 | 11 | 2.1282 | 18.1534 | | 0.11% |
| App. - LR | 8 | PCB Congener 195 | 4 | 4 | 2.0000 | | | |
| App. - LR | 8 | PCB Congener 196 | 7 | 1 | 2.7000 | | | |
| App. - LR | 8 | PCB Congener 196/203 | 7 | 7 | 5.9571 | | | |
| App. - LR | 8 | PCB Congener 201 | 15 | 9 | 4.3889 | | | |
| App. - LR | 8 | PCB Congener 202/171 | 6 | 6 | 0.9792 | | | |
| App. - LR | 9 | PCB Congener 206 | 15 | 9 | 3.0667 | 5.0667 | | 0.03% |
| App. - LR | 9 | PCB Congener 208 | 4 | 4 | 2.0000 | | | |
| LR - DP | 1 | PCB Congener 3 | 14 | 5 | 96.2000 | 96.2000 | 9,156.20 | 1.05% |
| LR - DP | 2 | PCB Congener 4 | 2 | 2 | 109.0000 | 1,051.9877 | | 11.49% |
| LR - DP | 2 | PCB Congener 4/10 | 12 | 3 | 36.0667 | | | |
| LR - DP | 2 | PCB Congener 6 | 31 | 28 | 41.0571 | | | |
| LR - DP | 2 | PCB Congener 7 | 19 | 17 | 4.3076 | | | |
| LR - DP | 2 | PCB Congener 7/9 | 12 | 5 | 9.2600 | | | |
| LR - DP | 2 | PCB Congener 8 | 14 | 14 | 169.3786 | | | |
| LR - DP | 2 | PCB Congener 8/5 | 17 | 17 | 351.1176 | | | |
| LR - DP | 2 | PCB Congener 9 | 2 | 2 | 14.7000 | | | |
| LR - DP | 2 | PCB Congener 12 | 14 | 6 | 6.8500 | | | |
| LR - DP | 2 | PCB Congener 15 | 2 | 2 | 239.5000 | | | |
| LR - DP | 2 | PCB Congener 15 | 12 | 11 | 70.7500 | | | |
| LR - DP | 3 | PCB Congener 16 | 14 | 13 | 165.1538 | 3,539.6776 | | 38.66% |
| LR - DP | 3 | PCB Congener 16/32 | 17 | 17 | 178.6941 | | | |
| LR - DP | 3 | PCB Congener 17 | 19 | 19 | 156.5579 | | | |
| LR - DP | 3 | PCB Congener 17 | 12 | 11 | 70.7500 | | | |
| LR - DP | 3 | PCB Congener 18 | 31 | 31 | 228.3323 | | | |
| LR - DP | 3 | PCB Congener 19 | 27 | 16 | 22.9813 | | | |
| LR - DP | 3 | PCB Congener 20 | 2 | 2 | 204.5000 | | | |
| LR - DP | 3 | PCB Congener 20/33 | 12 | 11 | 98.0636 | | | |
| LR - DP | 3 | PCB Congener 22 | 31 | 31 | 182.5968 | | | |
| LR - DP | 3 | PCB Congener 24/27 | 17 | 17 | 18.1376 | | | |
| LR - DP | 3 | PCB Congener 25 | 14 | 12 | 46.6667 | | | |
| LR - DP | 3 | PCB Congener 26 | 31 | 29 | 84.3138 | | | |
| LR - DP | 3 | PCB Congener 27 | 14 | 12 | 25.8167 | | | |
| LR - DP | 3 | PCB Congener 28 | 14 | 14 | 380.1714 | | | |
| LR - DP | 3 | PCB Congener 28/31 | 17 | 17 | 881.2941 | | | |
| LR - DP | 3 | PCB Congener 29 | 14 | 1 | 8.3000 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|-----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| LR - DP | 3 | PCB Congener 31 | 14 | 14 | 314.2429 | | | |
| LR - DP | 3 | PCB Congener 33 | 19 | 19 | 321.2105 | | | |
| LR - DP | 3 | PCB Congener 37 | 14 | 13 | 87.3923 | | | |
| LR - DP | 3 | PCB Congener 37 | 29 | 28 | 64.5018 | | | |
| LR - DP | 4 | PCB Congener 40 | 31 | 27 | 42.8852 | 3,289.5029 | | 35.93% |
| LR - DP | 4 | PCB Congener 41 | 14 | 13 | 87.1846 | | | |
| LR - DP | 4 | PCB Congener 41/64/71 | 16 | 16 | 170.9375 | | | |
| LR - DP | 4 | PCB Congener 42 | 2 | 2 | 188.5000 | | | |
| LR - DP | 4 | PCB Congener 42 | 29 | 28 | 64.5018 | | | |
| LR - DP | 4 | PCB Congener 44 | 31 | 30 | 159.7767 | | | |
| LR - DP | 4 | PCB Congener 45 | 31 | 30 | 30.7780 | | | |
| LR - DP | 4 | PCB Congener 46 | 17 | 17 | 15.4618 | | | |
| LR - DP | 4 | PCB Congener 47 | 2 | 2 | 228.5000 | | | |
| LR - DP | 4 | PCB Congener 47/48 | 17 | 17 | 109.0529 | | | |
| LR - DP | 4 | PCB Congener 47/75 | 12 | 11 | 45.7545 | | | |
| LR - DP | 4 | PCB Congener 49 | 31 | 27 | 136.5370 | | | |
| LR - DP | 4 | PCB Congener 52 | 31 | 30 | 148.2433 | | | |
| LR - DP | 4 | PCB Congener 53 | 2 | 2 | 99.0000 | | | |
| LR - DP | 4 | PCB Congener 53 | 12 | 11 | 48.3000 | | | |
| LR - DP | 4 | PCB Congener 56 | 2 | 2 | 257.0000 | | | |
| LR - DP | 4 | PCB Congener 56/60 | 17 | 17 | 184.1176 | | | |
| LR - DP | 4 | PCB Congener 56/60 | 12 | 11 | 38.2935 | | | |
| LR - DP | 4 | PCB Congener 59 | 14 | 4 | 18.3000 | | | |
| LR - DP | 4 | PCB Congener 66 | 2 | 2 | 525.0000 | | | |
| LR - DP | 4 | PCB Congener 66 | 29 | 28 | 128.6250 | | | |
| LR - DP | 4 | PCB Congener 70 | 14 | 14 | 150.0286 | | | |
| LR - DP | 4 | PCB Congener 70/76 | 17 | 17 | 259.0000 | | | |
| LR - DP | 4 | PCB Congener 74 | 31 | 30 | 88.2467 | | | |
| LR - DP | 4 | PCB Congener 75 | 2 | 2 | 8.6500 | | | |
| LR - DP | 4 | PCB Congener 77 | 23 | 15 | 22.1000 | | | |
| LR - DP | 4 | PCB Congener 77 | 29 | 28 | 30.1857 | | | |
| LR - DP | 4 | PCB Congener 81 | 22 | 10 | 0.5194 | | | |
| LR - DP | 4 | PCB Congener 81 | 12 | 11 | 4.0230 | | | |
| LR - DP | 5 | PCB Congener 82 | 31 | 25 | 10.7272 | 761.3397 | | 8.32% |
| LR - DP | 5 | PCB Congener 83 | 14 | 2 | 13.5500 | | | |
| LR - DP | 5 | PCB Congener 84 | 14 | 9 | 19.5889 | | | |
| LR - DP | 5 | PCB Congener 84/92 | 17 | 17 | 41.3118 | | | |
| LR - DP | 5 | PCB Congener 85 | 31 | 23 | 14.2078 | | | |
| LR - DP | 5 | PCB Congener 87 | 19 | 19 | 22.8632 | | | |
| LR - DP | 5 | PCB Congener 87/115 | 12 | 11 | 8.1679 | | | |
| LR - DP | 5 | PCB Congener 91 | 31 | 22 | 13.9455 | | | |
| LR - DP | 5 | PCB Congener 92 | 2 | 2 | 14.7000 | | | |
| LR - DP | 5 | PCB Congener 92 | 12 | 11 | 18.8610 | | | |
| LR - DP | 5 | PCB Congener 95 | 2 | 2 | 136.5000 | | | |
| LR - DP | 5 | PCB Congener 95 | 29 | 28 | 128.6250 | | | |
| LR - DP | 5 | PCB Congener 97 | 31 | 25 | 19.7960 | | | |
| LR - DP | 5 | PCB Congener 99 | 31 | 28 | 22.8321 | | | |
| LR - DP | 5 | PCB Congener 101 | 31 | 29 | 39.8069 | | | |
| LR - DP | 5 | PCB Congener 105 | 23 | 20 | 12.1820 | | | |
| LR - DP | 5 | PCB Congener 107 | 14 | 3 | 6.7667 | | | |
| LR - DP | 5 | PCB Congener 110 | 2 | 2 | 127.5000 | | | |
| LR - DP | 5 | PCB Congener 110 | 29 | 28 | 30.1857 | | | |
| LR - DP | 5 | PCB Congener 114 | 21 | 9 | 4.5382 | | | |
| LR - DP | 5 | PCB Congener 118 | 40 | 39 | 34.2692 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| LR - DP | 5 | PCB Congener 119 | 14 | 1 | 6.8000 | | | |
| LR - DP | 5 | PCB Congener 123 | 11 | 1 | 5.9000 | | | |
| LR - DP | 5 | PCB Congener 123 | 12 | 8 | 6.4000 | | | |
| LR - DP | 5 | PCB Congener 126 | 23 | 5 | 0.2256 | | | |
| LR - DP | 5 | PCB Congener 126 | 12 | 2 | 1.0890 | | | |
| LR - DP | 6 | PCB Congener 128 | 14 | 5 | 9.0600 | 221.8007 | | 2.42% |
| LR - DP | 6 | PCB Congener 129 | 12 | 2 | 1.0890 | | | |
| LR - DP | 6 | PCB Congener 132/153 | 17 | 17 | 37.8706 | | | |
| LR - DP | 6 | PCB Congener 135 | 14 | 5 | 10.7200 | | | |
| LR - DP | 6 | PCB Congener 135/144 | 15 | 15 | 3.6440 | | | |
| LR - DP | 6 | PCB Congener 136 | 27 | 7 | 6.7571 | | | |
| LR - DP | 6 | PCB Congener 137/176 | 4 | 3 | 0.7000 | | | |
| LR - DP | 6 | PCB Congener 138 | 14 | 13 | 23.9154 | | | |
| LR - DP | 6 | PCB Congener 141 | 29 | 16 | 5.5131 | | | |
| LR - DP | 6 | PCB Congener 146 | 13 | 7 | 26.0429 | | | |
| LR - DP | 6 | PCB Congener 149 | 19 | 19 | 21.5158 | | | |
| LR - DP | 6 | PCB Congener 149 | 12 | 8 | 6.4000 | | | |
| LR - DP | 6 | PCB Congener 151 | 31 | 21 | 6.5586 | | | |
| LR - DP | 6 | PCB Congener 153 | 14 | 13 | 21.0308 | | | |
| LR - DP | 6 | PCB Congener 156 | 23 | 14 | 2.5757 | | | |
| LR - DP | 6 | PCB Congener 157 | 11 | 9 | 0.2768 | | | |
| LR - DP | 6 | PCB Congener 157 | 12 | 1 | 1.0000 | | | |
| LR - DP | 6 | PCB Congener 163/138 | 17 | 17 | 26.8471 | | | |
| LR - DP | 6 | PCB Congener 167 | 23 | 9 | 1.1089 | | | |
| LR - DP | 6 | PCB Congener 168 | 14 | 4 | 9.1750 | | | |
| LR - DP | 7 | PCB Congener 170 | 23 | 14 | 8.2129 | 99.0852 | | 1.08% |
| LR - DP | 7 | PCB Congener 170/190 | 16 | 13 | 9.6000 | | | |
| LR - DP | 7 | PCB Congener 171 | 14 | 2 | 8.8500 | | | |
| LR - DP | 7 | PCB Congener 202/171 | 14 | 14 | 1.8036 | | | |
| LR - DP | 7 | PCB Congener 172/197 | 1 | 1 | 0.9000 | | | |
| LR - DP | 7 | PCB Congener 174 | 31 | 21 | 8.0348 | | | |
| LR - DP | 7 | PCB Congener 176 | 14 | 3 | 4.6667 | | | |
| LR - DP | 7 | PCB Congener 137/176 | 4 | 3 | 0.7000 | | | |
| LR - DP | 7 | PCB Congener 177 | 29 | 19 | 5.9474 | | | |
| LR - DP | 7 | PCB Congener 178 | 6 | 4 | 1.9250 | | | |
| LR - DP | 7 | PCB Congener 178 | 12 | 2 | 1.0890 | | | |
| LR - DP | 7 | PCB Congener 179 | 14 | 1 | 6.6000 | | | |
| LR - DP | 7 | PCB Congener 180 | 40 | 36 | 14.1722 | | | |
| LR - DP | 7 | PCB Congener 182/187 | 17 | 17 | 9.2088 | | | |
| LR - DP | 7 | PCB Congener 183 | 29 | 17 | 5.9482 | | | |
| LR - DP | 7 | PCB Congener 187 | 14 | 9 | 11.2778 | | | |
| LR - DP | 7 | PCB Congener 189 | 23 | 8 | 0.1489 | | | |
| LR - DP | 8 | PCB Congener 194 | 30 | 27 | 5.0407 | 50.9273 | | 0.56% |
| LR - DP | 8 | PCB Congener 195 | 14 | 2 | 4.8500 | | | |
| LR - DP | 8 | PCB Congener 195 | 16 | 16 | 3.4728 | | | |
| LR - DP | 8 | PCB Congener 196 | 14 | 3 | 11.7000 | | | |
| LR - DP | 8 | PCB Congener 196/203 | 16 | 16 | 9.7638 | | | |
| LR - DP | 8 | PCB Congener 172/197 | 1 | 1 | 0.9000 | | | |
| LR - DP | 8 | PCB Congener 200 | 12 | 1 | 1.0000 | | | |
| LR - DP | 8 | PCB Congener 201 | 31 | 26 | 8.2631 | | | |
| LR - DP | 8 | PCB Congener 202 | 14 | 3 | 4.1333 | | | |
| LR - DP | 8 | PCB Congener 202/171 | 14 | 14 | 1.8036 | | | |
| LR - DP | 9 | PCB Congener 206 | 30 | 21 | 7.5300 | 31.2600 | | 0.34% |
| LR - DP | 9 | PCB Congener 207 | 14 | 1 | 2.1000 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|-----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| LR - DP | 9 | PCB Congener 208 | 14 | 7 | 18.1571 | | | |
| LR - DP | 9 | PCB Congener 208 | 16 | 16 | 3.4728 | | | |
| LR - DP | 10 | PCB Congener 209 | 14 | 5 | 14.4200 | 14.4200 | | 0.16% |
| DP - GB | 1 | PCB Congener 3 | 5 | 1 | 10.0000 | 10.0000 | 4,066.09 | 0.25% |
| DP - GB | 2 | PCB Congener 6 | 8 | 8 | 45.9250 | 546.9683 | | 13.45% |
| DP - GB | 2 | PCB Congener 7 | 3 | 3 | 3.8333 | | | |
| DP - GB | 2 | PCB Congener 7/9 | 5 | 3 | 9.1000 | | | |
| DP - GB | 2 | PCB Congener 8 | 5 | 5 | 128.5000 | | | |
| DP - GB | 2 | PCB Congener 8/5 | 3 | 3 | 296.6667 | | | |
| DP - GB | 2 | PCB Congener 12 | 5 | 3 | 8.2333 | | | |
| DP - GB | 2 | PCB Congener 15 | 5 | 5 | 54.7100 | | | |
| DP - GB | 3 | PCB Congener 16 | 5 | 5 | 84.7000 | 1,993.0505 | | 49.02% |
| DP - GB | 3 | PCB Congener 16/32 | 3 | 3 | 110.0000 | | | |
| DP - GB | 3 | PCB Congener 17 | 3 | 3 | 77.0000 | | | |
| DP - GB | 3 | PCB Congener 17 | 5 | 5 | 54.7100 | | | |
| DP - GB | 3 | PCB Congener 18 | 8 | 8 | 129.2125 | | | |
| DP - GB | 3 | PCB Congener 19 | 8 | 7 | 10.0000 | | | |
| DP - GB | 3 | PCB Congener 20/33 | 5 | 5 | 70.5376 | | | |
| DP - GB | 3 | PCB Congener 22 | 8 | 8 | 86.7875 | | | |
| DP - GB | 3 | PCB Congener 24/27 | 3 | 3 | 9.0333 | | | |
| DP - GB | 3 | PCB Congener 25 | 5 | 5 | 39.0400 | | | |
| DP - GB | 3 | PCB Congener 26 | 8 | 5 | 54.4000 | | | |
| DP - GB | 3 | PCB Congener 27 | 5 | 4 | 15.8000 | | | |
| DP - GB | 3 | PCB Congener 28 | 5 | 5 | 242.4000 | | | |
| DP - GB | 3 | PCB Congener 28/31 | 3 | 3 | 586.6667 | | | |
| DP - GB | 3 | PCB Congener 31 | 5 | 5 | 183.9400 | | | |
| DP - GB | 3 | PCB Congener 33 | 3 | 3 | 136.6667 | | | |
| DP - GB | 3 | PCB Congener 37 | 5 | 5 | 71.6000 | | | |
| DP - GB | 3 | PCB Congener 37 | 8 | 8 | 30.5563 | | | |
| DP - GB | 4 | PCB Congener 40 | 8 | 8 | 22.2225 | 1,091.3948 | | 26.84% |
| DP - GB | 4 | PCB Congener 41 | 5 | 5 | 43.6800 | | | |
| DP - GB | 4 | PCB Congener 41/64/71 | 3 | 3 | 96.0000 | | | |
| DP - GB | 4 | PCB Congener 42 | 8 | 8 | 30.5563 | | | |
| DP - GB | 4 | PCB Congener 44 | 8 | 8 | 79.2875 | | | |
| DP - GB | 4 | PCB Congener 45 | 8 | 6 | 17.8333 | | | |
| DP - GB | 4 | PCB Congener 46 | 3 | 3 | 9.2000 | | | |
| DP - GB | 4 | PCB Congener 47/48 | 3 | 3 | 65.0000 | | | |
| DP - GB | 4 | PCB Congener 47/75 | 5 | 5 | 40.0200 | | | |
| DP - GB | 4 | PCB Congener 49 | 8 | 7 | 72.7714 | | | |
| DP - GB | 4 | PCB Congener 52 | 8 | 8 | 90.3875 | | | |
| DP - GB | 4 | PCB Congener 53 | 5 | 5 | 34.7424 | | | |
| DP - GB | 4 | PCB Congener 56/60 | 3 | 3 | 98.6667 | | | |
| DP - GB | 4 | PCB Congener 56/60 | 5 | 5 | 32.9238 | | | |
| DP - GB | 4 | PCB Congener 59 | 5 | 4 | 7.7750 | | | |
| DP - GB | 4 | PCB Congener 66 | 8 | 8 | 53.8188 | | | |
| DP - GB | 4 | PCB Congener 70 | 5 | 5 | 69.1800 | | | |
| DP - GB | 4 | PCB Congener 70/76 | 3 | 3 | 143.3333 | | | |
| DP - GB | 4 | PCB Congener 74 | 8 | 8 | 44.9125 | | | |
| DP - GB | 4 | PCB Congener 77 | 26 | 24 | 13.9667 | | | |
| DP - GB | 4 | PCB Congener 77 | 8 | 8 | 20.4875 | | | |
| DP - GB | 4 | PCB Congener 81 | 21 | 16 | 0.0823 | | | |
| DP - GB | 4 | PCB Congener 81 | 5 | 5 | 4.5474 | | | |
| DP - GB | 5 | PCB Congener 82 | 8 | 8 | 6.8150 | 242.9705 | | 5.98% |
| DP - GB | 5 | PCB Congener 84 | 5 | 4 | 11.6250 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| DP - GB | 5 | PCB Congener 84/92 | 3 | 3 | 24.6667 | | | |
| DP - GB | 5 | PCB Congener 85 | 8 | 3 | 8.8333 | | | |
| DP - GB | 5 | PCB Congener 87 | 3 | 3 | 13.3333 | | | |
| DP - GB | 5 | PCB Congener 87/115 | 5 | 5 | 9.2326 | | | |
| DP - GB | 5 | PCB Congener 91 | 8 | 3 | 5.4333 | | | |
| DP - GB | 5 | PCB Congener 92 | 5 | 5 | 16.2162 | | | |
| DP - GB | 5 | PCB Congener 95 | 8 | 8 | 53.8188 | | | |
| DP - GB | 5 | PCB Congener 97 | 8 | 7 | 9.2429 | | | |
| DP - GB | 5 | PCB Congener 99 | 8 | 7 | 13.2143 | | | |
| DP - GB | 5 | PCB Congener 101 | 8 | 8 | 24.3250 | | | |
| DP - GB | 5 | PCB Congener 105 | 26 | 25 | 5.8464 | | | |
| DP - GB | 5 | PCB Congener 110 | 8 | 8 | 20.4875 | | | |
| DP - GB | 5 | PCB Congener 114 | 23 | 17 | 1.0948 | | | |
| DP - GB | 5 | PCB Congener 118 | 26 | 26 | 12.7077 | | | |
| DP - GB | 5 | PCB Congener 123 | 21 | 3 | 1.1867 | | | |
| DP - GB | 5 | PCB Congener 123 | 5 | 4 | 4.8125 | | | |
| DP - GB | 5 | PCB Congener 126 | 26 | 5 | 0.0786 | | | |
| DP - GB | 6 | PCB Congener 128 | 7 | 4 | 5.8250 | 123.2046 | | 3.03% |
| DP - GB | 6 | PCB Congener 132/153 | 3 | 3 | 28.3333 | | | |
| DP - GB | 6 | PCB Congener 135/144 | 3 | 3 | 2.9333 | | | |
| DP - GB | 6 | PCB Congener 136 | 8 | 2 | 2.7500 | | | |
| DP - GB | 6 | PCB Congener 138 | 5 | 5 | 14.4600 | | | |
| DP - GB | 6 | PCB Congener 141 | 8 | 3 | 2.5000 | | | |
| DP - GB | 6 | PCB Congener 146 | 3 | 3 | 15.0000 | | | |
| DP - GB | 6 | PCB Congener 149 | 3 | 3 | 11.8667 | | | |
| DP - GB | 6 | PCB Congener 149 | 5 | 4 | 4.8125 | | | |
| DP - GB | 6 | PCB Congener 151 | 8 | 3 | 3.6000 | | | |
| DP - GB | 6 | PCB Congener 153 | 5 | 5 | 12.4000 | | | |
| DP - GB | 6 | PCB Congener 156 | 26 | 21 | 0.6490 | | | |
| DP - GB | 6 | PCB Congener 157 | 21 | 16 | 0.0491 | | | |
| DP - GB | 6 | PCB Congener 163/138 | 3 | 3 | 17.6667 | | | |
| DP - GB | 6 | PCB Congener 167 | 23 | 18 | 0.3589 | | | |
| DP - GB | 7 | PCB Congener 170 | 23 | 18 | 1.3228 | 31.1289 | | 0.77% |
| DP - GB | 7 | PCB Congener 174 | 8 | 1 | 4.5000 | | | |
| DP - GB | 7 | PCB Congener 177 | 8 | 1 | 12.0000 | | | |
| DP - GB | 7 | PCB Congener 180 | 26 | 24 | 4.1154 | | | |
| DP - GB | 7 | PCB Congener 182/187 | 3 | 3 | 6.4667 | | | |
| DP - GB | 7 | PCB Congener 183 | 7 | 2 | 2.6500 | | | |
| DP - GB | 7 | PCB Congener 189 | 23 | 6 | 0.0740 | | | |
| DP - GB | 8 | PCB Congener 194 | 8 | 5 | 4.5200 | 21.9200 | | 0.54% |
| DP - GB | 8 | PCB Congener 195 | 2 | 1 | 1.7000 | | | |
| DP - GB | 8 | PCB Congener 196 | 5 | 1 | 4.8000 | | | |
| DP - GB | 8 | PCB Congener 196/203 | 3 | 3 | 5.8667 | | | |
| DP - GB | 8 | PCB Congener 201 | 8 | 3 | 5.0333 | | | |
| DP - GB | 9 | PCB Congener 206 | 8 | 2 | 3.7500 | 5.4500 | | 0.13% |
| DP - GB | 9 | PCB Congener 208 | 2 | 1 | 1.7000 | | | |
| GB Zone 2 | 1 | PCB Congener 1 | 4 | 1 | 18.7590 | 23.9435 | 550.68 | 4.35% |
| GB Zone 2 | 1 | PCB Congener 3 | 4 | 2 | 5.1845 | | | |
| GB Zone 2 | 2 | PCB Congener 4/10 | 4 | 3 | 0.4050 | 38.3098 | | 6.96% |
| GB Zone 2 | 2 | PCB Congener 7 | 4 | 3 | 1.0197 | | | |
| GB Zone 2 | 2 | PCB Congener 8/5 | 4 | 3 | 35.3137 | | | |
| GB Zone 2 | 2 | PCB Congener 12/13 | 4 | 2 | 1.5715 | | | |
| GB Zone 2 | 3 | PCB Congener 16/32 | 4 | 3 | 11.1297 | 153.4626 | | 27.87% |
| GB Zone 2 | 3 | PCB Congener 17 | 4 | 3 | 7.5823 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|-----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| GB Zone 2 | 3 | PCB Congener 18 | 4 | 4 | 10.2173 | | | |
| GB Zone 2 | 3 | PCB Congener 19 | 4 | 1 | 1.0520 | | | |
| GB Zone 2 | 3 | PCB Congener 21 | 4 | 2 | 0.1480 | | | |
| GB Zone 2 | 3 | PCB Congener 22 | 4 | 3 | 31.5537 | | | |
| GB Zone 2 | 3 | PCB Congener 24/27 | 4 | 1 | 0.8410 | | | |
| GB Zone 2 | 3 | PCB Congener 25 | 4 | 3 | 4.4237 | | | |
| GB Zone 2 | 3 | PCB Congener 26 | 4 | 4 | 6.8410 | | | |
| GB Zone 2 | 3 | PCB Congener 28/31 | 4 | 4 | 50.9130 | | | |
| GB Zone 2 | 3 | PCB Congener 29 | 4 | 4 | 0.1865 | | | |
| GB Zone 2 | 3 | PCB Congener 33 | 4 | 4 | 21.6815 | | | |
| GB Zone 2 | 3 | PCB Congener 37 | 4 | 1 | 6.8930 | | | |
| GB Zone 2 | 4 | PCB Congener 40 | 4 | 3 | 5.8867 | 218.3708 | | 39.65% |
| GB Zone 2 | 4 | PCB Congener 41/64/71 | 4 | 4 | 16.7390 | | | |
| GB Zone 2 | 4 | PCB Congener 42 | 4 | 1 | 6.8930 | | | |
| GB Zone 2 | 4 | PCB Congener 43 | 4 | 1 | 3.5020 | | | |
| GB Zone 2 | 4 | PCB Congener 45 | 4 | 4 | 2.3463 | | | |
| GB Zone 2 | 4 | PCB Congener 46 | 4 | 1 | 4.5510 | | | |
| GB Zone 2 | 4 | PCB Congener 47/48 | 4 | 2 | 12.8135 | | | |
| GB Zone 2 | 4 | PCB Congener 49 | 4 | 3 | 18.4027 | | | |
| GB Zone 2 | 4 | PCB Congener 52 | 4 | 3 | 18.9647 | | | |
| GB Zone 2 | 4 | PCB Congener 53 | 4 | 3 | 1.5893 | | | |
| GB Zone 2 | 4 | PCB Congener 56/60 | 4 | 4 | 37.3703 | | | |
| GB Zone 2 | 4 | PCB Congener 63 | 4 | 3 | 2.2890 | | | |
| GB Zone 2 | 4 | PCB Congener 66 | 4 | 4 | 18.3600 | | | |
| GB Zone 2 | 4 | PCB Congener 70/76 | 4 | 4 | 45.1458 | | | |
| GB Zone 2 | 4 | PCB Congener 74 | 4 | 4 | 13.7695 | | | |
| GB Zone 2 | 4 | PCB Congener 77 | 11 | 11 | 3.2344 | | | |
| GB Zone 2 | 4 | PCB Congener 77 | 4 | 4 | 6.3291 | | | |
| GB Zone 2 | 4 | PCB Congener 81 | 15 | 12 | 0.1848 | | | |
| GB Zone 2 | 5 | PCB Congener 82 | 4 | 3 | 2.4943 | 72.0384 | | 13.08% |
| GB Zone 2 | 5 | PCB Congener 83 | 4 | 3 | 0.7897 | | | |
| GB Zone 2 | 5 | PCB Congener 84/92 | 4 | 3 | 7.9770 | | | |
| GB Zone 2 | 5 | PCB Congener 85 | 4 | 4 | 4.0218 | | | |
| GB Zone 2 | 5 | PCB Congener 87 | 4 | 4 | 4.2118 | | | |
| GB Zone 2 | 5 | PCB Congener 89 | 4 | 3 | 0.1213 | | | |
| GB Zone 2 | 5 | PCB Congener 91 | 4 | 3 | 1.5403 | | | |
| GB Zone 2 | 5 | PCB Congener 95 | 4 | 4 | 18.3600 | | | |
| GB Zone 2 | 5 | PCB Congener 97 | 4 | 4 | 1.9723 | | | |
| GB Zone 2 | 5 | PCB Congener 99 | 4 | 4 | 3.5993 | | | |
| GB Zone 2 | 5 | PCB Congener 100 | 4 | 3 | 1.3547 | | | |
| GB Zone 2 | 5 | PCB Congener 101 | 4 | 4 | 5.3833 | | | |
| GB Zone 2 | 5 | PCB Congener 105 | 11 | 10 | 2.0212 | | | |
| GB Zone 2 | 5 | PCB Congener 105 | 4 | 4 | 5.6258 | | | |
| GB Zone 2 | 5 | PCB Congener 107 | 4 | 2 | 1.6135 | | | |
| GB Zone 2 | 5 | PCB Congener 110 | 4 | 4 | 6.3291 | | | |
| GB Zone 2 | 5 | PCB Congener 114 | 11 | 8 | 0.1876 | | | |
| GB Zone 2 | 5 | PCB Congener 118 | 15 | 14 | 4.0893 | | | |
| GB Zone 2 | 5 | PCB Congener 119 | 4 | 2 | 0.1755 | | | |
| GB Zone 2 | 5 | PCB Congener 124 | 4 | 4 | 0.1222 | | | |
| GB Zone 2 | 5 | PCB Congener 126 | 11 | 5 | 0.0486 | | | |
| GB Zone 2 | 6 | PCB Congener 128 | 4 | 3 | 0.7257 | 19.2484 | | 3.50% |
| GB Zone 2 | 6 | PCB Congener 129 | 4 | 4 | 0.2425 | | | |
| GB Zone 2 | 6 | PCB Congener 130 | 4 | 4 | 0.3435 | | | |
| GB Zone 2 | 6 | PCB Congener 131 | 4 | 3 | 0.1850 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|------------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| GB Zone 2 | 6 | PCB Congener 132/153 | 4 | 4 | 5.6258 | | | |
| GB Zone 2 | 6 | PCB Congener 135/144/1 | 4 | 4 | 0.3666 | | | |
| GB Zone 2 | 6 | PCB Congener 137/176 | 4 | 2 | 0.0663 | | | |
| GB Zone 2 | 6 | PCB Congener 138/158/1 | 4 | 4 | 4.7700 | | | |
| GB Zone 2 | 6 | PCB Congener 141 | 4 | 4 | 0.6868 | | | |
| GB Zone 2 | 6 | PCB Congener 146 | 4 | 3 | 1.6010 | | | |
| GB Zone 2 | 6 | PCB Congener 149 | 4 | 3 | 3.1307 | | | |
| GB Zone 2 | 6 | PCB Congener 151 | 4 | 4 | 0.7355 | | | |
| GB Zone 2 | 6 | PCB Congener 156 | 11 | 8 | 0.1841 | | | |
| GB Zone 2 | 6 | PCB Congener 156 | 4 | 3 | 0.2705 | | | |
| GB Zone 2 | 6 | PCB Congener 157 | 11 | 7 | 0.0560 | | | |
| GB Zone 2 | 6 | PCB Congener 167 | 15 | 11 | 0.2586 | | | |
| GB Zone 2 | 7 | PCB Congener 170 | 11 | 11 | 0.4102 | 11.8947 | | 2.16% |
| GB Zone 2 | 7 | PCB Congener 170/190 | 4 | 4 | 2.2940 | | | |
| GB Zone 2 | 7 | PCB Congener 171 | 4 | 3 | 0.2705 | | | |
| GB Zone 2 | 7 | PCB Congener 172/197 | 4 | 1 | 0.0990 | | | |
| GB Zone 2 | 7 | PCB Congener 173 | 4 | 1 | 0.1060 | | | |
| GB Zone 2 | 7 | PCB Congener 174 | 4 | 3 | 1.6693 | | | |
| GB Zone 2 | 7 | PCB Congener 175 | 4 | 1 | 0.4480 | | | |
| GB Zone 2 | 7 | PCB Congener 137/176 | 4 | 2 | 0.0663 | | | |
| GB Zone 2 | 7 | PCB Congener 177 | 4 | 3 | 1.3483 | | | |
| GB Zone 2 | 7 | PCB Congener 178 | 4 | 4 | 0.2425 | | | |
| GB Zone 2 | 7 | PCB Congener 180 | 15 | 13 | 1.2165 | | | |
| GB Zone 2 | 7 | PCB Congener 182/187 | 4 | 4 | 1.5838 | | | |
| GB Zone 2 | 7 | PCB Congener 183 | 4 | 3 | 1.0497 | | | |
| GB Zone 2 | 7 | PCB Congener 185 | 4 | 3 | 0.3050 | | | |
| GB Zone 2 | 7 | PCB Congener 189 | 15 | 3 | 0.1027 | | | |
| GB Zone 2 | 7 | PCB Congener 191 | 4 | 1 | 0.3950 | | | |
| GB Zone 2 | 7 | PCB Congener 193 | 4 | 3 | 0.2880 | | | |
| GB Zone 2 | 8 | PCB Congener 194 | 4 | 2 | 0.9255 | 10.2542 | | 1.86% |
| GB Zone 2 | 8 | PCB Congener 195 | 4 | 1 | 0.8890 | | | |
| GB Zone 2 | 8 | PCB Congener 196/203 | 4 | 2 | 2.7965 | | | |
| GB Zone 2 | 8 | PCB Congener 172/197 | 4 | 1 | 0.0990 | | | |
| GB Zone 2 | 8 | PCB Congener 198 | 4 | 1 | 0.2140 | | | |
| GB Zone 2 | 8 | PCB Congener 199 | 4 | 2 | 0.0970 | | | |
| GB Zone 2 | 8 | PCB Congener 201 | 4 | 3 | 3.3247 | | | |
| GB Zone 2 | 8 | PCB Congener 202 | 4 | 3 | 0.2705 | | | |
| GB Zone 2 | 8 | PCB Congener 205 | 4 | 1 | 1.6380 | | | |
| GB Zone 2 | 9 | PCB Congener 206 | 4 | 1 | 1.9400 | 2.8290 | | 0.51% |
| GB Zone 2 | 9 | PCB Congener 208 | 4 | 1 | 0.8890 | | | |
| GB Zone 2 | 10 | PCB Congener 209 | 4 | 1 | 0.3270 | 0.3270 | | 0.06% |
| GB Zone 3A | 1 | PCB Congener 1 | 13 | 7 | 4.3679 | 11.1525 | 472.59 | 2.36% |
| GB Zone 3A | 1 | PCB Congener 3 | 13 | 3 | 6.7847 | | | |
| GB Zone 3A | 2 | PCB Congener 4/10 | 13 | 10 | 0.5980 | 25.5100 | | 5.40% |
| GB Zone 3A | 2 | PCB Congener 6 | 13 | 3 | 1.1973 | | | |
| GB Zone 3A | 2 | PCB Congener 7 | 13 | 13 | 0.8700 | | | |
| GB Zone 3A | 2 | PCB Congener 8/5 | 13 | 12 | 22.0933 | | | |
| GB Zone 3A | 2 | PCB Congener 12/13 | 13 | 11 | 0.7515 | | | |
| GB Zone 3A | 3 | PCB Congener 16/32 | 13 | 13 | 5.4380 | 119.4549 | | 25.28% |
| GB Zone 3A | 3 | PCB Congener 17 | 13 | 12 | 5.5622 | | | |
| GB Zone 3A | 3 | PCB Congener 18 | 13 | 13 | 8.3340 | | | |
| GB Zone 3A | 3 | PCB Congener 19 | 13 | 8 | 0.3958 | | | |
| GB Zone 3A | 3 | PCB Congener 21 | 13 | 8 | 0.4376 | | | |
| GB Zone 3A | 3 | PCB Congener 22 | 13 | 12 | 19.4934 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|-----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| GB Zone 3A | 3 | PCB Congener 24/27 | 13 | 8 | 0.6333 | | | |
| GB Zone 3A | 3 | PCB Congener 25 | 13 | 12 | 3.3117 | | | |
| GB Zone 3A | 3 | PCB Congener 26 | 13 | 12 | 5.9483 | | | |
| GB Zone 3A | 3 | PCB Congener 28/31 | 13 | 12 | 44.8115 | | | |
| GB Zone 3A | 3 | PCB Congener 29 | 13 | 8 | 0.3323 | | | |
| GB Zone 3A | 3 | PCB Congener 33 | 13 | 13 | 17.4222 | | | |
| GB Zone 3A | 3 | PCB Congener 37 | 13 | 2 | 7.3348 | | | |
| GB Zone 3A | 4 | PCB Congener 40 | 13 | 11 | 3.6555 | 180.7664 | | 38.25% |
| GB Zone 3A | 4 | PCB Congener 41/64/71 | 13 | 12 | 16.9991 | | | |
| GB Zone 3A | 4 | PCB Congener 42 | 13 | 2 | 7.3348 | | | |
| GB Zone 3A | 4 | PCB Congener 43 | 13 | 7 | 0.6460 | | | |
| GB Zone 3A | 4 | PCB Congener 45 | 13 | 11 | 1.5173 | | | |
| GB Zone 3A | 4 | PCB Congener 46 | 13 | 8 | 1.3573 | | | |
| GB Zone 3A | 4 | PCB Congener 47/48 | 13 | 7 | 5.9351 | | | |
| GB Zone 3A | 4 | PCB Congener 49 | 13 | 13 | 11.1544 | | | |
| GB Zone 3A | 4 | PCB Congener 52 | 13 | 12 | 12.8433 | | | |
| GB Zone 3A | 4 | PCB Congener 53 | 13 | 10 | 0.9681 | | | |
| GB Zone 3A | 4 | PCB Congener 56/60 | 13 | 13 | 34.8905 | | | |
| GB Zone 3A | 4 | PCB Congener 63 | 13 | 9 | 1.9296 | | | |
| GB Zone 3A | 4 | PCB Congener 66 | 13 | 13 | 20.5125 | | | |
| GB Zone 3A | 4 | PCB Congener 70/76 | 13 | 13 | 39.3267 | | | |
| GB Zone 3A | 4 | PCB Congener 74 | 13 | 13 | 14.2927 | | | |
| GB Zone 3A | 4 | PCB Congener 77 | 2 | 2 | 0.0420 | | | |
| GB Zone 3A | 4 | PCB Congener 77 | 13 | 13 | 6.9159 | | | |
| GB Zone 3A | 4 | PCB Congener 81 | 15 | 14 | 0.4457 | | | |
| GB Zone 3A | 5 | PCB Congener 82 | 13 | 13 | 1.8583 | 80.5422 | | 17.04% |
| GB Zone 3A | 5 | PCB Congener 83 | 13 | 12 | 0.6175 | | | |
| GB Zone 3A | 5 | PCB Congener 84/92 | 13 | 13 | 6.6253 | | | |
| GB Zone 3A | 5 | PCB Congener 85 | 13 | 13 | 3.4478 | | | |
| GB Zone 3A | 5 | PCB Congener 87 | 13 | 12 | 4.7658 | | | |
| GB Zone 3A | 5 | PCB Congener 89 | 13 | 8 | 0.3065 | | | |
| GB Zone 3A | 5 | PCB Congener 91 | 13 | 11 | 1.4870 | | | |
| GB Zone 3A | 5 | PCB Congener 95 | 13 | 13 | 20.5125 | | | |
| GB Zone 3A | 5 | PCB Congener 97 | 13 | 13 | 2.6139 | | | |
| GB Zone 3A | 5 | PCB Congener 99 | 13 | 13 | 4.5394 | | | |
| GB Zone 3A | 5 | PCB Congener 100 | 13 | 10 | 1.2399 | | | |
| GB Zone 3A | 5 | PCB Congener 101 | 13 | 13 | 6.6102 | | | |
| GB Zone 3A | 5 | PCB Congener 105 | 2 | 1 | 1.6000 | | | |
| GB Zone 3A | 5 | PCB Congener 105 | 13 | 13 | 6.9265 | | | |
| GB Zone 3A | 5 | PCB Congener 107 | 13 | 9 | 1.4812 | | | |
| GB Zone 3A | 5 | PCB Congener 110 | 13 | 13 | 6.9159 | | | |
| GB Zone 3A | 5 | PCB Congener 114 | 2 | 1 | 0.0930 | | | |
| GB Zone 3A | 5 | PCB Congener 114 | 13 | 4 | 0.4580 | | | |
| GB Zone 3A | 5 | PCB Congener 118 | 15 | 11 | 7.6610 | | | |
| GB Zone 3A | 5 | PCB Congener 119 | 13 | 11 | 0.5803 | | | |
| GB Zone 3A | 5 | PCB Congener 124 | 13 | 11 | 0.2020 | | | |
| GB Zone 3A | 6 | PCB Congener 128 | 13 | 12 | 1.0562 | 26.4454 | | 5.60% |
| GB Zone 3A | 6 | PCB Congener 129 | 13 | 12 | 0.5795 | | | |
| GB Zone 3A | 6 | PCB Congener 130 | 13 | 11 | 0.7565 | | | |
| GB Zone 3A | 6 | PCB Congener 131 | 13 | 9 | 0.3253 | | | |
| GB Zone 3A | 6 | PCB Congener 132/153 | 13 | 13 | 6.9265 | | | |
| GB Zone 3A | 6 | PCB Congener 134 | 13 | 4 | 0.4580 | | | |
| GB Zone 3A | 6 | PCB Congener 135/144/ | 13 | 11 | 0.6061 | | | |
| GB Zone 3A | 6 | PCB Congener 137/176 | 13 | 7 | 0.0931 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|------------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| GB Zone 3A | 6 | PCB Congener 138/158/1 | 13 | 13 | 7.4459 | | | |
| GB Zone 3A | 6 | PCB Congener 141 | 13 | 11 | 1.0925 | | | |
| GB Zone 3A | 6 | PCB Congener 146 | 13 | 12 | 1.6548 | | | |
| GB Zone 3A | 6 | PCB Congener 149 | 13 | 13 | 3.0812 | | | |
| GB Zone 3A | 6 | PCB Congener 151 | 13 | 13 | 0.9704 | | | |
| GB Zone 3A | 6 | PCB Congener 156 | 2 | 1 | 0.4800 | | | |
| GB Zone 3A | 6 | PCB Congener 156 | 13 | 10 | 0.2503 | | | |
| GB Zone 3A | 6 | PCB Congener 157 | 2 | 1 | 0.0830 | | | |
| GB Zone 3A | 6 | PCB Congener 167 | 15 | 8 | 0.5859 | | | |
| GB Zone 3A | 7 | PCB Congener 170 | 2 | 1 | 0.7800 | 14.1858 | | 3.00% |
| GB Zone 3A | 7 | PCB Congener 170/190 | 13 | 10 | 2.1566 | | | |
| GB Zone 3A | 7 | PCB Congener 171 | 13 | 10 | 0.2503 | | | |
| GB Zone 3A | 7 | PCB Congener 172/197 | 13 | 7 | 0.2560 | | | |
| GB Zone 3A | 7 | PCB Congener 173 | 13 | 1 | 0.0160 | | | |
| GB Zone 3A | 7 | PCB Congener 174 | 13 | 12 | 1.6533 | | | |
| GB Zone 3A | 7 | PCB Congener 175 | 13 | 8 | 0.4044 | | | |
| GB Zone 3A | 7 | PCB Congener 137/176 | 13 | 7 | 0.0931 | | | |
| GB Zone 3A | 7 | PCB Congener 177 | 13 | 12 | 1.1793 | | | |
| GB Zone 3A | 7 | PCB Congener 178 | 13 | 12 | 0.5795 | | | |
| GB Zone 3A | 7 | PCB Congener 180 | 15 | 13 | 2.0082 | | | |
| GB Zone 3A | 7 | PCB Congener 182/187 | 13 | 12 | 2.4848 | | | |
| GB Zone 3A | 7 | PCB Congener 183 | 13 | 12 | 1.2763 | | | |
| GB Zone 3A | 7 | PCB Congener 185 | 13 | 6 | 0.2582 | | | |
| GB Zone 3A | 7 | PCB Congener 189 | 15 | 6 | 0.1915 | | | |
| GB Zone 3A | 7 | PCB Congener 191 | 13 | 1 | 0.2640 | | | |
| GB Zone 3A | 7 | PCB Congener 193 | 13 | 9 | 0.3344 | | | |
| GB Zone 3A | 8 | PCB Congener 194 | 13 | 10 | 0.8751 | 10.9323 | | 2.31% |
| GB Zone 3A | 8 | PCB Congener 195 | 13 | 11 | 1.0118 | | | |
| GB Zone 3A | 8 | PCB Congener 196/203 | 13 | 7 | 2.2071 | | | |
| GB Zone 3A | 8 | PCB Congener 172/197 | 13 | 7 | 0.2560 | | | |
| GB Zone 3A | 8 | PCB Congener 198 | 13 | 5 | 4.0386 | | | |
| GB Zone 3A | 8 | PCB Congener 199 | 13 | 5 | 0.1062 | | | |
| GB Zone 3A | 8 | PCB Congener 201 | 13 | 9 | 2.1871 | | | |
| GB Zone 3A | 8 | PCB Congener 202 | 13 | 10 | 0.2503 | | | |
| GB Zone 3A | 9 | PCB Congener 206 | 13 | 11 | 1.5355 | 2.7247 | | 0.58% |
| GB Zone 3A | 9 | PCB Congener 207 | 13 | 9 | 0.1773 | | | |
| GB Zone 3A | 9 | PCB Congener 208 | 13 | 11 | 1.0118 | | | |
| GB Zone 3A | 10 | PCB Congener 209 | 13 | 11 | 0.8754 | 0.8754 | | 0.19% |
| GB Zone 3B | 1 | PCB Congener 1 | 33 | 20 | 8.4662 | 19.3947 | 663.37 | 2.92% |
| GB Zone 3B | 1 | PCB Congener 3 | 33 | 6 | 10.9285 | | | |
| GB Zone 3B | 2 | PCB Congener 4/10 | 33 | 30 | 0.7071 | 40.3738 | | 6.09% |
| GB Zone 3B | 2 | PCB Congener 6 | 33 | 4 | 1.0568 | | | |
| GB Zone 3B | 2 | PCB Congener 7 | 33 | 26 | 1.3457 | | | |
| GB Zone 3B | 2 | PCB Congener 8/5 | 33 | 33 | 35.4009 | | | |
| GB Zone 3B | 2 | PCB Congener 12/13 | 33 | 29 | 1.8634 | | | |
| GB Zone 3B | 3 | PCB Congener 16/32 | 33 | 32 | 9.3606 | 161.1755 | | 24.30% |
| GB Zone 3B | 3 | PCB Congener 17 | 33 | 31 | 7.5766 | | | |
| GB Zone 3B | 3 | PCB Congener 18 | 33 | 33 | 12.9205 | | | |
| GB Zone 3B | 3 | PCB Congener 19 | 33 | 24 | 0.4962 | | | |
| GB Zone 3B | 3 | PCB Congener 21 | 33 | 15 | 0.4486 | | | |
| GB Zone 3B | 3 | PCB Congener 22 | 33 | 33 | 32.7182 | | | |
| GB Zone 3B | 3 | PCB Congener 24/27 | 33 | 17 | 1.2219 | | | |
| GB Zone 3B | 3 | PCB Congener 25 | 33 | 33 | 4.6565 | | | |
| GB Zone 3B | 3 | PCB Congener 26 | 33 | 33 | 8.8535 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|------------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| GB Zone 3B | 3 | PCB Congener 28/31 | 33 | 33 | 48.1308 | | | |
| GB Zone 3B | 3 | PCB Congener 29 | 33 | 23 | 0.3081 | | | |
| GB Zone 3B | 3 | PCB Congener 33 | 33 | 33 | 29.0042 | | | |
| GB Zone 3B | 3 | PCB Congener 37 | 33 | 3 | 5.4797 | | | |
| GB Zone 3B | 4 | PCB Congener 40 | 33 | 29 | 6.1076 | 249.9880 | | 37.68% |
| GB Zone 3B | 4 | PCB Congener 41/64/71 | 33 | 33 | 22.3618 | | | |
| GB Zone 3B | 4 | PCB Congener 42 | 33 | 3 | 5.4797 | | | |
| GB Zone 3B | 4 | PCB Congener 43 | 33 | 15 | 0.8621 | | | |
| GB Zone 3B | 4 | PCB Congener 45 | 33 | 24 | 2.9187 | | | |
| GB Zone 3B | 4 | PCB Congener 46 | 33 | 17 | 1.6671 | | | |
| GB Zone 3B | 4 | PCB Congener 47/48 | 33 | 12 | 23.4554 | | | |
| GB Zone 3B | 4 | PCB Congener 49 | 33 | 28 | 15.8384 | | | |
| GB Zone 3B | 4 | PCB Congener 52 | 33 | 28 | 17.0753 | | | |
| GB Zone 3B | 4 | PCB Congener 53 | 33 | 22 | 1.6578 | | | |
| GB Zone 3B | 4 | PCB Congener 56/60 | 33 | 33 | 43.9080 | | | |
| GB Zone 3B | 4 | PCB Congener 63 | 33 | 21 | 2.5280 | | | |
| GB Zone 3B | 4 | PCB Congener 66 | 33 | 33 | 24.4177 | | | |
| GB Zone 3B | 4 | PCB Congener 70/76 | 33 | 33 | 48.3495 | | | |
| GB Zone 3B | 4 | PCB Congener 74 | 33 | 33 | 22.2603 | | | |
| GB Zone 3B | 4 | PCB Congener 77 | 4 | 4 | 0.6100 | | | |
| GB Zone 3B | 4 | PCB Congener 77 | 33 | 33 | 9.9160 | | | |
| GB Zone 3B | 4 | PCB Congener 81 | 37 | 32 | 0.5747 | | | |
| GB Zone 3B | 5 | PCB Congener 82 | 33 | 32 | 2.5683 | 112.7651 | | 17.00% |
| GB Zone 3B | 5 | PCB Congener 83 | 33 | 33 | 0.9134 | | | |
| GB Zone 3B | 5 | PCB Congener 84/92 | 33 | 29 | 11.0144 | | | |
| GB Zone 3B | 5 | PCB Congener 85 | 33 | 33 | 5.1421 | | | |
| GB Zone 3B | 5 | PCB Congener 87 | 33 | 33 | 6.1528 | | | |
| GB Zone 3B | 5 | PCB Congener 89 | 33 | 26 | 0.2863 | | | |
| GB Zone 3B | 5 | PCB Congener 91 | 33 | 31 | 1.8598 | | | |
| GB Zone 3B | 5 | PCB Congener 95 | 33 | 33 | 24.4177 | | | |
| GB Zone 3B | 5 | PCB Congener 97 | 33 | 33 | 3.2681 | | | |
| GB Zone 3B | 5 | PCB Congener 99 | 33 | 33 | 6.5144 | | | |
| GB Zone 3B | 5 | PCB Congener 100 | 33 | 30 | 2.0197 | | | |
| GB Zone 3B | 5 | PCB Congener 101 | 33 | 33 | 9.6962 | | | |
| GB Zone 3B | 5 | PCB Congener 105 | 4 | 4 | 0.5675 | | | |
| GB Zone 3B | 5 | PCB Congener 105 | 33 | 33 | 10.3893 | | | |
| GB Zone 3B | 5 | PCB Congener 107 | 33 | 26 | 2.7028 | | | |
| GB Zone 3B | 5 | PCB Congener 110 | 33 | 33 | 9.9160 | | | |
| GB Zone 3B | 5 | PCB Congener 114 | 4 | 2 | 0.0655 | | | |
| GB Zone 3B | 5 | PCB Congener 114 | 33 | 7 | 0.2910 | | | |
| GB Zone 3B | 5 | PCB Congener 118 | 37 | 33 | 13.7993 | | | |
| GB Zone 3B | 5 | PCB Congener 119 | 33 | 27 | 0.8160 | | | |
| GB Zone 3B | 5 | PCB Congener 124 | 33 | 29 | 0.3644 | | | |
| GB Zone 3B | 6 | PCB Congener 128 | 33 | 33 | 1.1920 | 36.6579 | | 5.53% |
| GB Zone 3B | 6 | PCB Congener 129 | 33 | 32 | 0.5620 | | | |
| GB Zone 3B | 6 | PCB Congener 130 | 33 | 32 | 1.1079 | | | |
| GB Zone 3B | 6 | PCB Congener 131 | 33 | 23 | 0.4337 | | | |
| GB Zone 3B | 6 | PCB Congener 132/153 | 33 | 33 | 10.3893 | | | |
| GB Zone 3B | 6 | PCB Congener 134 | 33 | 7 | 0.2910 | | | |
| GB Zone 3B | 6 | PCB Congener 135/144/1 | 33 | 29 | 1.0931 | | | |
| GB Zone 3B | 6 | PCB Congener 137/176 | 33 | 11 | 0.1496 | | | |
| GB Zone 3B | 6 | PCB Congener 138/158/1 | 33 | 32 | 11.2302 | | | |
| GB Zone 3B | 6 | PCB Congener 141 | 33 | 33 | 1.4413 | | | |
| GB Zone 3B | 6 | PCB Congener 146 | 33 | 30 | 2.2344 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| GB Zone 3B | 6 | PCB Congener 149 | 33 | 33 | 4.3656 | | | |
| GB Zone 3B | 6 | PCB Congener 151 | 33 | 33 | 1.3791 | | | |
| GB Zone 3B | 6 | PCB Congener 156 | 4 | 1 | 0.0680 | | | |
| GB Zone 3B | 6 | PCB Congener 156 | 33 | 22 | 0.3380 | | | |
| GB Zone 3B | 6 | PCB Congener 157 | 4 | 1 | 0.0420 | | | |
| GB Zone 3B | 6 | PCB Congener 167 | 37 | 22 | 0.3405 | | | |
| GB Zone 3B | 7 | PCB Congener 170 | 4 | 3 | 0.1500 | 19.4548 | | 2.93% |
| GB Zone 3B | 7 | PCB Congener 170/190 | 33 | 30 | 3.8601 | | | |
| GB Zone 3B | 7 | PCB Congener 171 | 33 | 22 | 0.3380 | | | |
| GB Zone 3B | 7 | PCB Congener 172/197 | 33 | 17 | 0.3779 | | | |
| GB Zone 3B | 7 | PCB Congener 173 | 33 | 2 | 0.1060 | | | |
| GB Zone 3B | 7 | PCB Congener 174 | 33 | 31 | 2.1793 | | | |
| GB Zone 3B | 7 | PCB Congener 175 | 33 | 18 | 0.3732 | | | |
| GB Zone 3B | 7 | PCB Congener 137/176 | 33 | 11 | 0.1496 | | | |
| GB Zone 3B | 7 | PCB Congener 177 | 33 | 31 | 2.2413 | | | |
| GB Zone 3B | 7 | PCB Congener 178 | 33 | 32 | 0.5620 | | | |
| GB Zone 3B | 7 | PCB Congener 180 | 37 | 34 | 3.1355 | | | |
| GB Zone 3B | 7 | PCB Congener 182/187 | 33 | 33 | 2.9870 | | | |
| GB Zone 3B | 7 | PCB Congener 183 | 33 | 33 | 1.8023 | | | |
| GB Zone 3B | 7 | PCB Congener 185 | 33 | 13 | 0.2805 | | | |
| GB Zone 3B | 7 | PCB Congener 189 | 37 | 13 | 0.2145 | | | |
| GB Zone 3B | 7 | PCB Congener 191 | 33 | 16 | 0.3550 | | | |
| GB Zone 3B | 7 | PCB Congener 193 | 33 | 14 | 0.3427 | | | |
| GB Zone 3B | 8 | PCB Congener 194 | 33 | 26 | 1.4137 | 17.0732 | | 2.57% |
| GB Zone 3B | 8 | PCB Congener 195 | 33 | 27 | 1.6040 | | | |
| GB Zone 3B | 8 | PCB Congener 196/203 | 33 | 21 | 6.0761 | | | |
| GB Zone 3B | 8 | PCB Congener 172/197 | 33 | 17 | 0.3779 | | | |
| GB Zone 3B | 8 | PCB Congener 198 | 33 | 4 | 0.5108 | | | |
| GB Zone 3B | 8 | PCB Congener 199 | 33 | 21 | 0.2794 | | | |
| GB Zone 3B | 8 | PCB Congener 201 | 33 | 25 | 6.2843 | | | |
| GB Zone 3B | 8 | PCB Congener 202 | 33 | 22 | 0.3380 | | | |
| GB Zone 3B | 8 | PCB Congener 205 | 33 | 1 | 0.1890 | | | |
| GB Zone 3B | 9 | PCB Congener 206 | 33 | 26 | 2.9855 | 4.8507 | | 0.73% |
| GB Zone 3B | 9 | PCB Congener 207 | 33 | 17 | 0.2612 | | | |
| GB Zone 3B | 9 | PCB Congener 208 | 33 | 27 | 1.6040 | | | |
| GB Zone 3B | 10 | PCB Congener 209 | 33 | 26 | 1.6382 | 1.6382 | | 0.25% |
| GB Zone 4 | 1 | PCB Congener 1 | 27 | 13 | 2.0243 | 4.9621 | 102.58 | 4.84% |
| GB Zone 4 | 1 | PCB Congener 3 | 27 | 6 | 2.9378 | | | |
| GB Zone 4 | 2 | PCB Congener 4/10 | 27 | 26 | 0.3325 | 3.7919 | | 3.70% |
| GB Zone 4 | 2 | PCB Congener 6 | 27 | 4 | 0.4548 | | | |
| GB Zone 4 | 2 | PCB Congener 7 | 27 | 17 | 0.1330 | | | |
| GB Zone 4 | 2 | PCB Congener 8/5 | 27 | 20 | 2.6141 | | | |
| GB Zone 4 | 2 | PCB Congener 12/13 | 27 | 15 | 0.2575 | | | |
| GB Zone 4 | 3 | PCB Congener 16/32 | 27 | 26 | 0.7281 | 18.6330 | | 18.16% |
| GB Zone 4 | 3 | PCB Congener 17 | 27 | 8 | 0.6011 | | | |
| GB Zone 4 | 3 | PCB Congener 18 | 27 | 26 | 1.2855 | | | |
| GB Zone 4 | 3 | PCB Congener 19 | 27 | 14 | 0.0851 | | | |
| GB Zone 4 | 3 | PCB Congener 21 | 27 | 16 | 0.1250 | | | |
| GB Zone 4 | 3 | PCB Congener 22 | 27 | 25 | 1.9688 | | | |
| GB Zone 4 | 3 | PCB Congener 24/27 | 27 | 13 | 0.2820 | | | |
| GB Zone 4 | 3 | PCB Congener 25 | 27 | 26 | 0.4237 | | | |
| GB Zone 4 | 3 | PCB Congener 26 | 27 | 26 | 0.5198 | | | |
| GB Zone 4 | 3 | PCB Congener 28/31 | 27 | 27 | 10.1617 | | | |
| GB Zone 4 | 3 | PCB Congener 29 | 27 | 8 | 0.2581 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|-----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| GB Zone 4 | 3 | PCB Congener 33 | 27 | 27 | 1.8574 | | | |
| GB Zone 4 | 3 | PCB Congener 37 | 27 | 3 | 0.3367 | | | |
| GB Zone 4 | 4 | PCB Congener 40 | 27 | 15 | 0.3694 | 30.7348 | | 29.96% |
| GB Zone 4 | 4 | PCB Congener 41/64/71 | 27 | 27 | 1.8457 | | | |
| GB Zone 4 | 4 | PCB Congener 42 | 27 | 3 | 0.3367 | | | |
| GB Zone 4 | 4 | PCB Congener 43 | 27 | 5 | 0.0940 | | | |
| GB Zone 4 | 4 | PCB Congener 45 | 27 | 19 | 0.4082 | | | |
| GB Zone 4 | 4 | PCB Congener 46 | 27 | 16 | 0.2266 | | | |
| GB Zone 4 | 4 | PCB Congener 47/48 | 27 | 13 | 1.1982 | | | |
| GB Zone 4 | 4 | PCB Congener 49 | 27 | 23 | 1.4382 | | | |
| GB Zone 4 | 4 | PCB Congener 52 | 27 | 27 | 1.7965 | | | |
| GB Zone 4 | 4 | PCB Congener 53 | 27 | 7 | 0.2114 | | | |
| GB Zone 4 | 4 | PCB Congener 56/60 | 27 | 25 | 5.9397 | | | |
| GB Zone 4 | 4 | PCB Congener 63 | 27 | 22 | 0.4315 | | | |
| GB Zone 4 | 4 | PCB Congener 66 | 27 | 27 | 5.6385 | | | |
| GB Zone 4 | 4 | PCB Congener 70/76 | 27 | 27 | 5.8307 | | | |
| GB Zone 4 | 4 | PCB Congener 74 | 27 | 27 | 3.0719 | | | |
| GB Zone 4 | 4 | PCB Congener 77 | 4 | 2 | 0.0250 | | | |
| GB Zone 4 | 4 | PCB Congener 77 | 27 | 27 | 1.6262 | | | |
| GB Zone 4 | 4 | PCB Congener 81 | 31 | 27 | 0.2466 | | | |
| GB Zone 4 | 5 | PCB Congener 82 | 27 | 26 | 0.3182 | 24.2265 | | 23.62% |
| GB Zone 4 | 5 | PCB Congener 83 | 27 | 22 | 0.2520 | | | |
| GB Zone 4 | 5 | PCB Congener 84/92 | 27 | 18 | 2.1434 | | | |
| GB Zone 4 | 5 | PCB Congener 85 | 27 | 27 | 1.1971 | | | |
| GB Zone 4 | 5 | PCB Congener 87 | 27 | 26 | 0.8422 | | | |
| GB Zone 4 | 5 | PCB Congener 89 | 27 | 7 | 0.1423 | | | |
| GB Zone 4 | 5 | PCB Congener 91 | 27 | 11 | 0.2488 | | | |
| GB Zone 4 | 5 | PCB Congener 95 | 27 | 27 | 5.6385 | | | |
| GB Zone 4 | 5 | PCB Congener 97 | 27 | 27 | 1.0555 | | | |
| GB Zone 4 | 5 | PCB Congener 99 | 27 | 24 | 1.6614 | | | |
| GB Zone 4 | 5 | PCB Congener 100 | 27 | 4 | 0.6223 | | | |
| GB Zone 4 | 5 | PCB Congener 101 | 27 | 25 | 2.0490 | | | |
| GB Zone 4 | 5 | PCB Congener 105 | 4 | 2 | 0.0480 | | | |
| GB Zone 4 | 5 | PCB Congener 105 | 27 | 27 | 2.2273 | | | |
| GB Zone 4 | 5 | PCB Congener 107 | 27 | 24 | 0.4522 | | | |
| GB Zone 4 | 5 | PCB Congener 110 | 27 | 27 | 1.6262 | | | |
| GB Zone 4 | 5 | PCB Congener 114 | 27 | 6 | 0.1770 | | | |
| GB Zone 4 | 5 | PCB Congener 118 | 31 | 28 | 3.0913 | | | |
| GB Zone 4 | 5 | PCB Congener 119 | 27 | 27 | 0.3317 | | | |
| GB Zone 4 | 5 | PCB Congener 124 | 27 | 24 | 0.1022 | | | |
| GB Zone 4 | 6 | PCB Congener 128 | 27 | 27 | 0.5943 | 9.9753 | | 9.72% |
| GB Zone 4 | 6 | PCB Congener 129 | 27 | 23 | 0.2113 | | | |
| GB Zone 4 | 6 | PCB Congener 130 | 27 | 24 | 0.3299 | | | |
| GB Zone 4 | 6 | PCB Congener 131 | 27 | 12 | 0.1402 | | | |
| GB Zone 4 | 6 | PCB Congener 132/153 | 27 | 27 | 2.2273 | | | |
| GB Zone 4 | 6 | PCB Congener 134 | 27 | 6 | 0.1770 | | | |
| GB Zone 4 | 6 | PCB Congener 135/144/ | 27 | 24 | 0.3067 | | | |
| GB Zone 4 | 6 | PCB Congener 137/176 | 27 | 2 | 0.0458 | | | |
| GB Zone 4 | 6 | PCB Congener 138/158/ | 27 | 24 | 3.3320 | | | |
| GB Zone 4 | 6 | PCB Congener 141 | 27 | 26 | 0.3175 | | | |
| GB Zone 4 | 6 | PCB Congener 146 | 27 | 24 | 0.6942 | | | |
| GB Zone 4 | 6 | PCB Congener 149 | 27 | 27 | 0.9519 | | | |
| GB Zone 4 | 6 | PCB Congener 151 | 27 | 27 | 0.3213 | | | |
| GB Zone 4 | 6 | PCB Congener 156 | 4 | 1 | 0.0089 | | | |

Appendix I. PCB Congener Data and Homolog Results in Sediment

| Location (Reach/Zone) | PCB Homolog Group # | PCB Congener | Total Number of Samples | Total Number Detected | RI Mean Result (ug/kg) | Cumulative Homolog Result | Reach Zone Total PCB | Percent of Reach/Zone Total PCB |
|-----------------------|---------------------|----------------------|-------------------------|-----------------------|------------------------|---------------------------|----------------------|---------------------------------|
| GB Zone 4 | 6 | PCB Congener 156 | 27 | 14 | 0.0860 | | | |
| GB Zone 4 | 6 | PCB Congener 167 | 31 | 20 | 0.2310 | | | |
| GB Zone 4 | 7 | PCB Congener 170/190 | 27 | 25 | 0.6776 | 4.5844 | | 4.47% |
| GB Zone 4 | 7 | PCB Congener 171 | 27 | 14 | 0.0860 | | | |
| GB Zone 4 | 7 | PCB Congener 172/197 | 27 | 15 | 0.0682 | | | |
| GB Zone 4 | 7 | PCB Congener 173 | 27 | 3 | 0.0700 | | | |
| GB Zone 4 | 7 | PCB Congener 174 | 27 | 24 | 0.4304 | | | |
| GB Zone 4 | 7 | PCB Congener 175 | 27 | 20 | 0.1761 | | | |
| GB Zone 4 | 7 | PCB Congener 137/176 | 27 | 2 | 0.0458 | | | |
| GB Zone 4 | 7 | PCB Congener 177 | 27 | 20 | 0.4112 | | | |
| GB Zone 4 | 7 | PCB Congener 178 | 27 | 23 | 0.2113 | | | |
| GB Zone 4 | 7 | PCB Congener 180 | 31 | 23 | 0.8312 | | | |
| GB Zone 4 | 7 | PCB Congener 182/187 | 27 | 27 | 0.7720 | | | |
| GB Zone 4 | 7 | PCB Congener 183 | 27 | 22 | 0.5081 | | | |
| GB Zone 4 | 7 | PCB Congener 189 | 31 | 3 | 0.1043 | | | |
| GB Zone 4 | 7 | PCB Congener 191 | 27 | 7 | 0.0991 | | | |
| GB Zone 4 | 7 | PCB Congener 193 | 27 | 16 | 0.0932 | | | |
| GB Zone 4 | 8 | PCB Congener 194 | 27 | 25 | 0.5615 | 4.1716 | | 4.07% |
| GB Zone 4 | 8 | PCB Congener 195 | 27 | 22 | 0.3888 | | | |
| GB Zone 4 | 8 | PCB Congener 196/203 | 27 | 18 | 0.5237 | | | |
| GB Zone 4 | 8 | PCB Congener 172/197 | 27 | 15 | 0.0682 | | | |
| GB Zone 4 | 8 | PCB Congener 198 | 27 | 12 | 1.7107 | | | |
| GB Zone 4 | 8 | PCB Congener 199 | 27 | 2 | 0.0415 | | | |
| GB Zone 4 | 8 | PCB Congener 201 | 27 | 22 | 0.7913 | | | |
| GB Zone 4 | 8 | PCB Congener 202 | 27 | 14 | 0.0860 | | | |
| GB Zone 4 | 9 | PCB Congener 206 | 27 | 22 | 0.6337 | 1.1377 | | 1.11% |
| GB Zone 4 | 9 | PCB Congener 207 | 27 | 16 | 0.1152 | | | |
| GB Zone 4 | 9 | PCB Congener 208 | 27 | 22 | 0.3888 | | | |
| GB Zone 4 | 10 | PCB Congener 209 | 27 | 24 | 0.3598 | 0.3598 | | 0.35% |