

Ecosystem Responses to Mercury Contamination

Indicators of Change

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4 Monitoring and Evaluating Trends in Methylmercury Accumulation in Aquatic Biota

James G. Wiener, R.A. Bodaly, Steven S. Brown, Marc Lucotte, Michael C. Newman, Donald B. Porcella, Robin J. Reash, and Edward B. Swain

ABSTRACT

The monitoring of mercury in aquatic food webs supporting the production of fish and wildlife is directly relevant to concerns about health and ecological risks of methylmercury (MeHg) exposure. We present a framework for monitoring concentrations of mercury in aquatic biota, with emphasis on assessing responses to changes in loadings of mercury from atmospheric deposition and other sources. In this chapter, we (1) identify specific attributes (criteria) of indicators that would be useful for discerning temporal trends and spatial patterns in the concentration of mercury in aquatic biota, (2) critically evaluate and rank candidate biological indicators useful for monitoring trends in mercury, (3) outline approaches for sampling and analysis of recommended biological indicators, (4) identify ancillary data needs and potential confounding factors that should be considered or documented to ensure the defensible interpretation of data on monitored biological indicators, and (5) consider the environmental settings (waterbody type and geographic location) that would be most sensitive for detecting changes in atmospheric deposition of mercury. Criteria were applied to ensure that the biological indicators selected are useful, relevant, and sufficiently diagnostic to detect a change in mercury bioaccumulation in response to altered mercury loadings. Toxicological problems with mercury in aquatic ecosystems result from biotic exposure to MeHg, a highly toxic compound that readily accumulates in exposed organisms and can biomagnify to high concentrations in organisms atop aquatic food webs. Biotic monitoring should, therefore, focus on assessing trends in bioaccumulation of MeHg; in samples from trophic levels below fish, this requires the determination of MeHg. We considered six general groups of

aquatic biological indicators: piscivorous fish, prey fish, benthic invertebrates, zooplankton, phytoplankton, and periphyton. Piscivorous fish and 1-year-old prey fish, all analyzed individually, are considered the preferred aquatic biological indicators for trend monitoring. For piscivorous fish, total-mercury determinations on axial muscle (preferably without skin), sampled annually, would indicate gradual (multi-year) trends in MeHg that are directly relevant to humans who eat sport fish. For prey fish, annual sampling and analysis of either whole fish or axial muscle for total mercury should indicate annual changes in exposure to MeHg. In North America, the historical record for MeHg in piscivorous fish (from data on total mercury in filets or axial muscle) extends about 35 years, much longer than the comparatively sparse historical record for MeHg in water and aquatic biota of lower trophic levels. The analytical method (cold vapor atomic absorption spectrophotometry) that produced most of the historic data on total mercury in piscivorous fish is valid, and the potential utility of these existing data for trend analysis merits careful consideration. Benthic invertebrates have been monitored and analyzed for mercury more extensively in estuarine systems than in fresh waters. The consumption of estuarine macroinvertebrates, such as oysters, clams, shrimp, and crabs, is also a direct pathway for human exposure to MeHg. Determination of MeHg in shellfish and macrocrustaceans could, therefore, be useful for trend monitoring in estuaries. The importance of MeHg uptake and transfer at the base of the food web is recognized. However, the utility of periphyton, phytoplankton, zooplankton, and freshwater benthic invertebrates for trend monitoring is diminished by interpretational complexities associated with large temporal variation in the biotic composition and MeHg content among samples, and by our limited understanding of the processes and variables that affect concentrations of MeHg in these groups. Many anthropogenic and natural factors, independent of the bulk loading of mercury from atmospheric deposition, can strongly influence the concentrations of MeHg in aquatic biota. Our ability to discern linkages between MeHg concentrations in aquatic biota and changing external loadings of mercury to aquatic systems will depend on knowledge of such factors and on the minimization of their confounding effects in biotic monitoring programs.

4.1 INTRODUCTION

Elevated mercury concentrations in fish tissue adversely affect the quality of fishery resources in many inland, coastal, and marine waters. In the United States, mercury was responsible for 76% of all fish-consumption advisories in 2004, and 44 states and 1 territory had advisories attributed to mercury (USEPA 2005). Growing awareness of the mercury problem has prompted increasing efforts to survey mercury in fish, and the number of statewide fish-consumption advisories issued for coastal waters, lakes, and rivers in the United States has increased substantially during the past decade (Wiener et al. 2003; USEPA 2005). In Canada, mercury accounted for more than 97% (2572) of all fish-consumption advisories listed in 1997 (USEPA 2001). Nearly all of the mercury in fish is methylmercury (MeHg), a highly toxic compound that readily crosses biological membranes, accumulates in exposed organisms, and can biomagnify to high concentrations in aquatic food webs (Grieb et al. 1990; Bloom 1992; Francesconi and Lenanton 1992; Wiener et al. 2003).

Atmospheric deposition is an important source of total mercury in many surface waters (Fitzgerald et al. 1998; Bindler et al. 2001; Lin et al. 2001; Lamborg et al. 2002), and it has been widely inferred that a significant portion of the MeHg bioaccumulated in many aquatic ecosystems is derived from mercury entering the surface water or its watershed in atmospheric deposition (Johansson et al. 1991; Watras et al. 1994; Rolfhus and Fitzgerald 1995; Jackson 1997; Monteiro and Furness 1997; Downs et al. 1998). In many remote and semi-remote areas of the Northern Hemisphere that lack in-watershed sources of anthropogenic mercury, the rate of mercury accumulation in lacustrine sediments has increased by a factor of 2 to 5 or more since the mid-1800s or early 1900s, based on analyses of dated cores of sediment and peat (Swain et al. 1992; Lucotte et al. 1995; Lockhart et al. 1998; Lorey and Driscoll 1999; Bindler et al. 2001; Lamborg et al. 2002; Shotyk et al. 2005). Some cores from semi-remote sites show evidence of recent declines in mercury deposition, possibly associated with decreasing emissions of anthropogenic mercury (Engstrom and Swain 1997; Benoit et al. 1998; Bindler et al. 2001; Shotyk et al. 2003, 2005).

We present a framework for monitoring concentrations of mercury in aquatic biota, with emphasis on assessing responses to changes in loadings of mercury from atmospheric deposition and other sources. The monitoring of mercury in aquatic food webs supporting the production of fish and wildlife is directly relevant to societal concerns about this toxic metal. Much of the scientific effort on mercury contamination of aquatic food webs has been prompted by the human health risks of MeHg exposure (Myers and Davidson 1998; Mahaffey 2000; Clarkson 2002), given that the consumption of finfish and shellfish is the primary exposure pathway in humans (NRC 2000; Mahaffey et al. 2004). Consumption of fish is also an important pathway of MeHg exposure for wildlife atop aquatic food webs (Heinz 1996; Wolfe et al. 1998; Wiener et al. 2003).

4.2 OBJECTIVES

This chapter focuses on monitoring trends in bioaccumulation in relation to anticipated changes in emissions of mercury from anthropogenic sources. Aquatic biota, however, are exposed to mercury from multiple sources, including historic anthropogenic, current anthropogenic, and natural sources. We identify aquatic biological indicators that can provide evidence of a temporal change in bioaccumulation of mercury (estimated from concentrations in tissue or whole organisms) from all sources. The objectives of this chapter are fivefold:

- 1) To identify specific attributes (criteria) of indicators that would be useful for discerning temporal trends and spatial patterns in the concentration of mercury in aquatic biota
- 2) To critically evaluate and rank candidate biological indicators useful for monitoring trends in mercury
- 3) To outline approaches for sampling and analysis of recommended biological indicators

- 4) To identify ancillary data needs and potential confounding factors that should be considered or documented to ensure the defensible interpretation of data on monitored biological indicators
- 5) To identify the environmental settings (water body type and geographic location) that would be most sensitive for detecting changes in atmospheric deposition of mercury in a trend-monitoring program

Many factors other than the bulk loading of mercury from atmospheric deposition can strongly influence the concentrations of MeHg in aquatic biota. Our ability to discern trends in MeHg concentrations in aquatic biota that are linked to changing loadings of mercury will depend on knowledge of such factors and on the minimization of their confounding effects in biotic monitoring programs for mercury.

4.3 AQUATIC BIOLOGICAL INDICATORS

4.3.1 CRITERIA TO SELECT INDICATORS

The selection of biological indicators should be guided by criteria to ensure that the indicators are relevant, useful, and sufficiently diagnostic to detect a change in the concentration of mercury in whole organisms or specific tissue(s) over multi-year or decadal time scales. We identified 9 criteria for the selection of biological indicators:

- 1) *Relevance.* A key criterion in the selection of biological indicators is relevance to human and ecological health and to the development of policy. Fish are directly relevant, for example, given that consumption of fish is the primary pathway for exposure to MeHg. The concentration of MeHg in fish is also a key variable in the issuance of fish-consumption advisories.
- 2) *Historical data on the indicator.* Existing information on the statistical variation, bias, and other interpretational attributes of potential biological indicators should be examined and considered in the design of a sampling program for assessing trends in mercury bioaccumulation.
- 3) *Clear recognition of confounding factors.* Many human and natural factors that are unrelated to bulk loadings of mercury from the atmosphere or other sources can strongly influence concentrations of mercury in aquatic biota. A *confounding factor* is here defined as a variable that interferes with the isolation of the effects of mercury loading on temporal trends in mercury concentrations in monitored biota within a water body or group of waters. Our ability to discern biotic trends in mercury concentrations linked to changing loadings of mercury to aquatic systems will require both knowledge of potential confounding factors and the minimization of their confounding effects in monitoring programs (see Section 4.6.2).
- 4) *Knowledge of intrinsic co-variables.* Concentrations of mercury in fish are typically correlated with age or body size. An understanding of, and ability to account for, the effects of such intrinsic variables is essential for evaluating contaminant trends.

- 5) *Broad geographic distribution.* When feasible, monitoring should focus on bioindicator species or taxa having a broad geographic range, to assess trends and patterns across large geographic areas.
- 6) *Importance in trophic transfer of MeHg.* Preference should be given to bioindicator organisms that have an important role in the trophic transfer of MeHg within food webs.
- 7) *Constrained feeding ecology and trophic position.* Temporal shifts in feeding habits or trophic position can alter dietary MeHg uptake, complicating the interpretation of trends in mercury concentration. An *a priori* knowledge of feeding ecology should facilitate the selection of suitable candidate bioindicator species during the design of a monitoring program. If a bioindicator is known to undergo ontogenetic shifts in diet, it would be advisable to limit sampling and analysis to a given life stage.
- 8) *Temporal response to changes in mercury loadings.* Long-lived bioindicators with long response times (several years) to changing mercury loadings could be more susceptible to the influence of confounding factors, possibly reducing their utility for detecting effects of altered loadings. Thus, short-lived (≤ 1 -year old), as well as long-lived, organisms should be considered when selecting potential bioindicators.
- 9) *Impacts of sampling on the target population.* Continued removal of individuals could eventually affect the concentration of mercury in members of the sampled population, creating artifacts in trend data. The potential effects of sampling on the target population, and potential approaches for reducing such effects, should be considered. Nonlethal techniques for sampling of fish tissue (Baker et al. 2004; Peterson et al. 2005) may be required when sampling protected populations or when sampling in protected environments, such as national parks.

4.3.2 CANDIDATE AQUATIC BIOLOGICAL INDICATORS

Six general groups of aquatic biological indicators were considered: 1) piscivorous fish, 2) prey fish, 3) benthic invertebrates, 4) zooplankton, 5) phytoplankton, and 6) periphyton. We begin with a brief summary of our understanding of mercury bioaccumulation and trophic transfer in aquatic ecosystems, to provide essential background information for the subsequent sections on aquatic biological indicators. This summary is based on selected recent reviews (Jackson 1998; Morel et al. 1998; Benoit et al. 2003; Wiener et al. 2003) and the original reports cited.

In a toxicological sense, the primary problem with mercury in aquatic ecosystems can be defined as biotic exposure to, or bioaccumulation of, MeHg. Trend monitoring of mercury in aquatic biota should accordingly focus on MeHg, which readily crosses biological membranes and accumulates to concentrations in aquatic organisms that vastly exceed those in surface water. In fish, concentrations of MeHg commonly exceed those in water by a factor of 10^6 to 10^7 or more. Most of the mercury in the aquatic environment is inorganic, yet nearly all (>95%) of the mercury accumulated in fish is MeHg (Grieb et al. 1990; Bloom 1992; Francesconi and Lenanton 1992), obtained almost entirely via dietary uptake (Rodgers 1994; Hall

et al. 1997; Harris and Bodaly 1998). Relative to MeHg, inorganic mercury is less readily transferred through successive trophic levels and does not biomagnify (Watras et al. 1998).

Methylmercury biomagnifies in aquatic food webs, and patterns in biomagnification are similar even among aquatic systems that differ in type of water body, mercury source, and pollution intensity. The transfer of MeHg in the upper trophic levels of aquatic food webs is almost entirely via dietary uptake, whereas direct uptake from water can be important for some lower food-chain organisms, such as phytoplankton and zooplankton. The concentration of MeHg increases up the food web from water and lower trophic levels to fish and piscivores, and the fraction of total mercury present as MeHg also increases with increasing trophic level through fish. The fraction of total mercury present as MeHg can vary greatly within trophic levels below fish. In aquatic invertebrates, for example, the MeHg fraction can range from about 10% to more than 90% of total mercury. It is, therefore, essential to determine MeHg (rather than total mercury) in biological samples from trophic levels below fish, including phytoplankton, periphyton, benthic invertebrates, and zooplankton. In fish, determination of total mercury, which requires less analytical effort and expense than MeHg, provides reliable estimates of MeHg concentration.

The greatest increase in MeHg concentration occurs in the trophic step between water and algae. Bioaccumulation factors between water and seston, for example, often range from about 10^5 to about 10^6 , whereas ratios of MeHg concentrations between successive trophic levels above algae are generally less than 10^1 . Within an assemblage of fish, concentrations of MeHg increase with ascending trophic level, and variation in trophic position accounts for much of the variation in mercury concentration among species within a given water body. Concentrations of MeHg in fish also increase with increasing age or size because of the very slow rate of elimination relative to the rapid rate of uptake and because larger fish can consume larger prey with higher concentrations of MeHg. Much of the MeHg accumulated in fish is stored in skeletal muscle, tightly bound to sulfhydryl groups in protein.

Although the entry of MeHg into the base of the food web and its subsequent transfer in the lowest trophic levels are poorly understood, it is evident that the concentration of MeHg in all trophic levels is strongly correlated with its supply from methylating environments. In fish, for example, much of the modern spatial variation in mercury concentrations (within a given trophic level) can be attributed to variation in factors and processes that affect the microbial production of MeHg and its entry intooxic waters.

4.3.2.1 Fish

We consider 2 groups of fish as candidate biological indicators for monitoring trends in MeHg: 1) piscivorous (fish-eating) fish and 2) small prey fish (often termed “forage fish”). Analyses of piscivorous fish reflect the concentrations of MeHg near the top of aquatic food webs, providing a useful measure of potential dietary exposure in humans who eat fish. Small prey fish are eaten by a variety of piscivorous fish, birds, and mammals, but are generally not important in the diet of humans. Humans eat large omnivorous fish; however, the substantial confounding effects of the large

spatial and temporal variation in the diet and trophic position of omnivorous fish diminish their potential utility for trend analysis.

4.3.2.1.1 *Piscivorous Fish*

Piscivorous fish can accumulate high concentrations of MeHg, and information on MeHg in piscivorous fish is directly relevant to the public and the policy community. Piscivorous fish are present in most surface waters and can be obtained with moderate sampling effort with a variety of active and passive gear. Sampling would generally not affect target populations except in very small lakes and streams, where nonlethal sampling would lessen impacts on target populations.

In the United States, the threshold mercury concentration for commercial sale of fish is determined by the Food and Drug Administration, whereas consumption advice for recreational (noncommercial) fish is developed by individual states and tribes. Mercury data collected for development of fish-consumption advisories are typically from analyses of filets (axial muscle tissue, with or without skin) for total mercury, with concentrations expressed on a wet-weight basis. Analysis of filets for total mercury yields a valid estimate of MeHg concentration (Grieb et al. 1990; Bloom 1992), whether the analyzed sample consists of a large filet or a small mass of tissue obtained with a biopsy needle (Cizdziel et al. 2002; Baker et al. 2004).

Many piscivorous fish are important recreationally or commercially. The sampling and analysis of heavily exploited fish stocks are not recommended for trend monitoring, because intensive fishing pressure — over a period of years to decades — can substantially reduce mercury concentrations in members of heavily exploited populations (Section 4.6.1). Interpretation of temporal trends in mercury concentrations in fish, relative to changes in mercury loadings, will be more defensible if applied to fish populations and water bodies that are not subjected to intensive fishing pressure. Many commercially and recreationally important marine fish have been significantly depleted by over-fishing (Pauly et al. 2003; Coleman et al. 2004; Hutchings and Reynolds 2004). The application of trend data on mercury concentrations in heavily exploited marine fishes, such as yellowfin tuna *Thunnus albacares* (Kraepiel et al. 2003), has questionable validity as an approach for assessing temporal changes in the abundance of MeHg in the ecosystem.

In summary, piscivorous fish are present in most surface waters, require moderate sampling effort, and are the primary pathway for dietary exposure of humans to MeHg. Analyses of filets or axial muscle for total mercury should indicate gradual (multi-year) trends in the supply of MeHg. Given these attributes, nonmigratory piscivorous fish are priority candidates for monitoring MeHg. In very small lakes or streams, nonintrusive sampling methods should be used to reduce impacts on sampled populations.

4.3.2.1.2 *Prey Fish*

Prey fish are here defined as small, usually short-lived, finfish. In North American fresh waters, many are members of the cyprinid (minnow), percid (perch), and centrarchid (sunfish) families. Prey fish are widely distributed, common, and important in the transfer of MeHg to higher trophic levels, such as piscivorous fish and many fish-eating birds. MeHg concentrations in prey fish of uniform age are less

susceptible to certain, potential confounding factors, such as variation in trophic position, than are concentrations in long-lived, piscivorous fish.

Mercury concentrations in prey fish are useful indicators of relative MeHg levels in food webs supporting the production of sport fish and wildlife, information relevant to the public and the policy community. There is a sizable scientific literature on MeHg in prey fish, but they have been monitored less extensively than sport fish. Effects of removal sampling on target populations would be insignificant in all but the very smallest lakes.

This group of fish includes species that feed on zooplankton, benthic invertebrates, and occasionally, periphyton (Roseman et al. 1996; Bodaly and Fudge 1999; Gorski et al. 1999). The size of young prey fish can vary seasonally because measurable growth occurs throughout much of the year. Small prey fish, such as yearling yellow perch (*Perca flavescens*), finescale dace (*Phoxinus neogaeus*), or mimic shiner (*Notropis volucellus*), typically grow rapidly during the summer in temperate lakes, increasing their biomass by 2- to 5-fold. Mercury concentrations are usually increasing or stable during this period of growth; consequently, the total body burden of mercury in individual fish increases substantially during summer (Bodaly and Fudge 1999; Gorski et al. 1999). Concentrations of MeHg in prey fish can be expected to vary seasonally (Figure 4.1), and such variation should be considered when crafting sampling protocols for trend monitoring. Prey fish require little to moderate sampling effort, and samples taken in the early spring or fall, when growth is slow and temporal variation in mercury concentration is less, may provide the best comparisons among years and surface waters. Analysis of prey fish for total mercury or MeHg would reveal inter-annual variation in MeHg exposure (Frost et al. 1999; Gorski et al. 1999).

In summary, prey fish are present in most surface waters, require moderate sampling effort, are important in the trophic transfer of MeHg in aquatic food webs, and probably indicate annual changes in exposure to MeHg. Given these attributes,

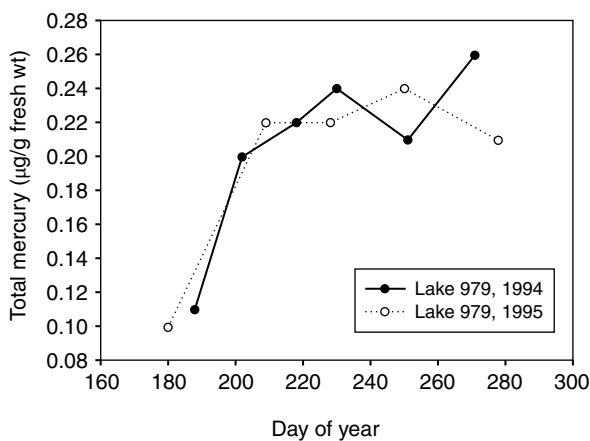


FIGURE 4.1 Whole-body concentrations of total mercury (present largely as MeHg) in caged finescale dace, showing seasonal increases in mercury concentrations during summer in Lake 979, an experimental reservoir in northwestern Ontario that was flooded during the Experimental Lakes Area Reservoir Project. (Source: Modified from Bodaly and Fudge 1999.)

prey fish are appropriate candidates for monitoring of MeHg. Seasonal variation should be considered carefully in the design of trend-monitoring protocols for prey fish.

4.3.2.1.3 Example of a Prey-Fish Indicator: Yellow Perch

Analyses of total mercury in whole bodies or axial muscle tissue of age-1 yellow perch have provided a useful measure of MeHg concentrations in food webs of many North American lakes. This widely distributed species inhabits lakes and reservoirs across much of the north-central, northeastern, and eastern United States and across the central and eastern provinces of Canada (Scott and Crossman 1973; Becker 1983). An ecologically similar congeneric species, the Eurasian perch (*Perca fluviatilis*), is distributed across much of Europe and northern Asia (Thorpe 1977).

During their first year, yellow perch have a small gape (jaw opening), which limits their diet largely to small zooplankton and small zoobenthos (Roseman et al. 1996; Lyons et al. 2000). Thus, the trophic position of age-1 yellow perch is not expected to vary substantially among sites. Generally abundant in lakes within much of its geographic range, the yellow perch is a preferred prey of certain piscivores, such as walleye (*Sander vitreus*) and common loons (*Gavia immer*), and is an important link in the food-web transfer of MeHg (Colby et al. 1979; Barr 1996). Concentrations of total mercury in age-1 or age-2 yellow perch are strongly and positively correlated with concentrations in coexisting piscivorous fish, including walleye, black bass (*Micropterus* spp.), and northern pike (*Esox lucius*) (Suns et al. 1987; Cope et al. 1990; JG Wiener, unpublished data for northern pike). Statistical analyses have shown strong relations between the total mercury concentration and burden of age-1 yellow perch and ecosystem characteristics (e.g., lake chemistry, wetland influence) or whole-lake manipulations (e.g., experimental acidification) that are known to influence the production of MeHg and its abundance in aquatic food webs (Grieb et al. 1990; Suns and Hitchin 1990; Wiener et al. 1990; Simonin et al. 1994; Frost et al. 1999; Wiener et al. 2003). Substantial mercury data are also available for the Eurasian perch (Metsaelae and Rask 1989; Andersson et al. 1995; Haines et al. 1995; Porvari 1998; Svobodova et al. 1999; Lindstroem 2001).

One-year-old yellow perch can be readily sampled in spring with small-mesh trap nets, seines, or small electroshockers fished in littoral habitat without significantly affecting their abundance or year-class strength. Age-1 fish obtained in spring have resided in a sampled lake for about 1 year. A target sample size of 15 to 30 whole, age-1 yellow perch (analyzed individually) from a given lake typically yields a standard error of the mean in the range of 1 to 6 ng/g wet weight, providing a precise estimate of mean whole-body concentration (JG Wiener, University of Wisconsin–La Crosse, data from Wisconsin and Minnesota lakes; corresponding mean concentrations range from about 20 to 200 ng/g wet weight). At age-1, the age of yellow perch can be accurately determined by examining scales taken near the area of insertion of the left pectoral fin.

4.3.2.2 Benthic Invertebrates

Benthic invertebrates are macroscopic animals that live at or near the sediment/water interface. Some benthic invertebrates, particularly mussels, readily accumulate metals, prompting their use as biological indicators of mercury contamination (Smith

and Green 1975). Benthic invertebrates have been widely used in biological monitoring of freshwater and marine habitats (Resh and McElvay 1993; Southerland and Stribling 1995; Weigel et al. 2003), providing a foundation for their use in trend-monitoring programs.

The dietary importance of benthic invertebrates to many species of fish, birds, and mammals (Vander Zanden and Vadeboncoeur 2002) signifies their importance in the trophic transfer of MeHg and their potential relevance as biological indicators. Some benthic invertebrates (e.g., oysters, clams, shrimp, crabs, and crayfish) are consumed by humans, providing a direct pathway for exposure to MeHg. In the United States, shellfish rank below fish as a source of dietary MeHg in the human population (NRC 2000; Schober et al. 2003).

Many benthic invertebrates have short life spans (≤ 1 year) but little is known about how quickly mercury concentrations in such organisms respond to changes in *external* loadings of mercury to the aquatic ecosystem. Bed sediment can be an important sink for mercury in aquatic systems if sediment-associated mercury is isolated from active biogeochemical cycling (Henry et al. 1995; Wiener and Shields 2000). Sediment can also serve as a source of MeHg in freshwater and estuarine ecosystems, given that the oxic/anoxic interface in the sediment is an important zone of mercury methylation (Gilmour et al. 1998; Benoit et al. 2003). Benthic organisms have physical contact with bed sediment, and the relative contributions of mercury from in-place sedimentary sources and of mercury from current external sources to their MeHg burdens will influence their sensitivity as indicators to altered mercury loadings from external sources. In this regard, sediment-dwelling invertebrates that feed on particles from the overlying water column — a group including many clams and aquatic insects — may be more useful than deposit feeders as indicators of external mercury loadings. The kinetics of MeHg bioaccumulation (ingestion, assimilation, and elimination) in benthic invertebrates have received little study but the limited available information should nonetheless be applied to the selection of candidate bioindicator species and to the interpretation of trend data.

The trophic position of benthic invertebrates varies widely among species. The diet is their primary pathway of contaminant exposure, and the feeding ecology of a benthic species largely determines its exposure to dissolved and particulate sedimentary contaminants (Brown et al. 2000). The dietary assimilation efficiency for MeHg (55–70%) in marine invertebrates is much higher than that for inorganic mercury (2–22%; Wang and Fisher 1999). In marine bivalves, exposure can occur through multiple pathways, including direct uptake from the overlying water or pore water and dietary uptake via ingestion of plankton and detritus from water or sediment (Thomann et al. 1995).

Benthic invertebrate communities are taxonomically and trophically complex, and their abundance and species composition in a water body often vary seasonally and among years. Sediment-dwelling invertebrates can be readily sampled but considerable effort is often required to remove benthic organisms from grab samples of sediment, to determine their taxonomic composition, and to obtain sufficient sample mass of a target taxon for analysis. Sampling would not substantially affect target populations.

In summary, benthic invertebrates are important in the trophic transfer of MeHg. The sensitivity of benthic organisms to altered mercury loadings from external sources will presumably depend on the relative contributions of mercury from in-place sediment and external sources to their total MeHg uptake. Moreover, the large effort required to produce samples that are well-defined taxonomically and trophically reduces their desirability as biotic indicators for trend monitoring in freshwater systems. Conversely, in estuarine systems, there has been more extensive monitoring and analysis of mercury in shellfish and macrocrustaceans, which could be useful in monitoring of estuaries.

4.3.2.3 Zooplankton

Zooplankton are small, often microscopic crustaceans that live in the water column. They are widely distributed, common, and important in pelagic food webs. Zooplankton are eaten by many fish and by early life stages of some fish that become piscivorous as juveniles or adults. Zooplankton are not eaten by humans but are an appropriate and relevant candidate indicator because of their importance in the trophic transfer of MeHg to fish. Sampling would not significantly affect populations or assemblages of zooplankton, even in small lakes.

Zooplankton vary seasonally and annually in abundance (Rusak et al. 2002), and respond within hours or days to changes in the supply of MeHg to the water column (Herrin et al. 1998; Paterson et al. 1998; Tsui and Wang 2004). Zooplankton can accumulate significant quantities of MeHg from food or water (Monson and Brezonik 1999; Peech Cherewyk 2002; Tsui and Wang 2004). Spatio-temporal variation in the taxonomic composition and abundance of zooplankton is large in temperate lakes (Rusak et al. 2002). Consequently, the trophic position of bulk zooplankton samples can vary greatly because these assemblages are composed of trophically diverse taxa that can eat phytoplankton, bacteria, detritus, and other zooplankton. Species assemblages can change rapidly, and a target species or taxon may not be available at certain times of the year.

The MeHg content of zooplankton varies among taxa (Back and Watras 1995), a complicating factor that can be eliminated by the determination of MeHg in individual taxa (Back et al. 1995). Zooplankton are readily sampled but samples should be checked and processed to remove phytoplankton and detritus. A single bulk sample from a plankton net can be used to characterize the open-water community of zooplankton in a lake at a particular time. The processing of samples can require substantial effort, and bulk samples of zooplankton from some surface waters contain particles that are extremely difficult to remove, diminishing the integrity of the sample.

Existing data on MeHg in zooplankton are few, and there has been little long-term monitoring of MeHg in zooplankton. In northern lakes, the concentrations of MeHg in zooplankton vary seasonally and annually (Figure 4.2), typically increasing in summer and declining in autumn (Herrin et al. 1998; Paterson et al. 1998). In thermally stratified lakes, the concentration of MeHg in zooplankton can increase markedly after the fall overturn (Figure 4.2), when MeHg in anoxic hypolimnetic

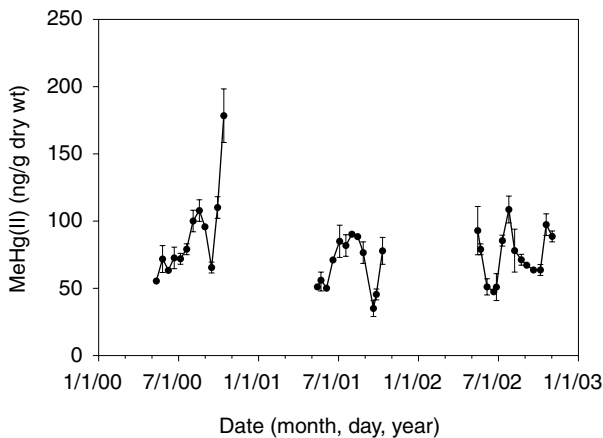


FIGURE 4.2 Concentrations of MeHg (mean \pm 1 standard error) in zooplankton from Lake 240 of the Experimental Lakes Area (northwestern Ontario, Canada), showing seasonal variation during summer and pronounced rapid increases in mean concentration after the fall overturn. (Source: Michael J. Paterson, Fisheries and Oceans Canada, Winnipeg, Manitoba, unpublished data.)

waters becomes mixed throughout the water column (Herrin et al. 1998). Such seasonal variation merits careful attention in sampling protocols for zooplankton in a trend program for mercury, regardless of whether bulk samples or specific taxa are used as a bioindicator, and may require multiple sampling events during the year. Samples taken in mid-summer would reflect conditions for bioaccumulation of MeHg during the period of maximal growth of fish but would miss the effect of the fall overturn in stratified lakes. Given such large intra- and inter-annual variation, we expect that many years of sampling and analysis would be needed to discern long-term trends in MeHg concentrations in zooplankton.

In summary, zooplankton are present in all lakes, are readily sampled, are important in the trophic transfer of MeHg, and would respond rapidly to changes in the supply of MeHg to the water column. Seasonal and interannual changes in MeHg concentration would complicate the timing of sampling and the interpretation of trend data. Pronounced short-term variability may reduce their suitability for assessing trends on the multi-year time scales expected for reductions in mercury emissions.

4.3.2.4 Phytoplankton

Phytoplankton are microscopic plants (algae) that live in the water column and are trophically situated at or near the base of pelagic food webs. Phytoplankton are taxonomically diverse, and their abundance and species composition vary seasonally, annually, and spatially. Phytoplankton are present in all aquatic systems and are considered important in the trophic transfer of MeHg. It has been generally assumed that zooplankton receive most of their MeHg via consumption of phytoplankton; however, recent research suggests that direct uptake of MeHg by zooplankton from water may be more important than previously thought (Monson and Brezonik 1999;

Peech Cherewyk 2002; Tsui and Wang 2004). Phytoplankton have a constrained trophic position; are not consumed by fish, wildlife, or humans, and are indirectly relevant to the public or the policy community. They would respond very rapidly — within minutes or hours — to changes in MeHg concentrations in water. Phytoplankton obtain MeHg directly from water (Mason et al. 1996), and algal density can influence concentrations of MeHg in phytoplankton via biomass dilution (Pickhardt et al. 2002).

Phytoplankton are easily sampled with fine-mesh nets. However, samples require considerable processing before analysis to remove zooplankton, detritus, or other particulates. Sampling would not measurably affect target populations, even in the smallest lakes.

Little is known about MeHg in phytoplankton, particularly freshwater phytoplankton (Becker and Bigham 1995). Concentrations of MeHg in freshwater phytoplankton are related to those in water but the partitioning of MeHg between water and phytoplankton is strongly affected by concentrations of dissolved organic matter (Watras et al. 1998).

In summary, phytoplankton are present in all lakes, are trophically well defined, and are a candidate indicator for trend monitoring. They would rapidly respond to changes in the supply of MeHg. However, discrete samples of phytoplankton are not easily obtained and phytoplankton are not a recommended indicator for trend monitoring, largely because of the complexities associated with interpretation of trend data for phytoplankton. Large intra- and inter-annual variations in the MeHg content of phytoplankton would complicate the interpretation of trend data and the identification of long-term trends.

4.3.2.5 Periphyton

Periphyton are microscopic and macroscopic algae that attach to and grow on solid surfaces, such as lake bottoms, rooted aquatic vegetation, and submerged woody debris. Periphyton form part of the base of littoral food webs in lakes. Periphyton communities are taxonomically diverse and the attached communities contain other organisms, such as bacteria and zooplankton, as well as detrital material. Periphyton vary seasonally and annually in both abundance and species composition.

Periphyton are widely distributed and common; however, the trophic position of the overall community is complex and variable. Periphyton have been studied little with regard to mercury cycling but would be expected to respond rapidly to changes in the supply of MeHg. Periphyton can be directly eaten by fish, and periphyton mats may be an important site for mercury methylation in some ecosystems (Cleckner et al. 1999). Methylmercury typically comprises a very small, but quite variable, fraction of the total mercury in periphyton (Cleckner et al. 1998; Bowles et al. 2001). Periphyton are relevant neither to the public nor the policy community, because they are not consumed by people or wildlife, and their importance in the cycling of mercury is unclear. Periphyton would be relatively easy to sample but samples would contain a complex mixture of plants, small invertebrates, and detritus, complicating interpretation of the MeHg concentrations therein. Sampling would not significantly affect target populations.

In summary, periphyton are present in all lakes, easy to sample, and would respond rapidly to changes in the abundance of MeHg. Periphyton are sometimes eaten directly by fish. The diverse, complex, and variable nature of the periphyton community, however, would complicate interpretation of mercury concentrations in periphyton in a monitoring program.

4.3.3 RECOMMENDED AQUATIC BIOLOGICAL INDICATORS

The criteria listed in Section 4.3.1 were applied to the selection of biological indicators to ensure their relevance and utility for assessing trends in the bioaccumulation of MeHg associated with altered loadings of mercury to aquatic systems. This evaluation, based largely on the discussion in Section 4.3.2, is summarized in Table 4.1.

We consider prey fish, piscivorous fish, and estuarine benthic invertebrates (listed in order of preference) suitable aquatic biological indicators for trend monitoring. One-year-old prey fish and piscivorous fish (analyzed individually) are the preferred biological indicators for freshwater systems. Determination of total mercury in axial muscle (preferably without skin) from piscivorous fish would indicate gradual (multi-year) trends in MeHg that are directly relevant to humans who eat sport fish. For 1-year-old prey fish, the annual sampling and analysis of either whole fish or axial muscle for total mercury would indicate annual changes in MeHg exposure in freshwater and marine ecosystems. In coastal estuaries, 1-year-old prey fish (analyzed for total mercury), as well as shellfish and macrocrustaceans (analyzed for MeHg), are recommended as biological indicators for trend monitoring. Adult piscivorous fish from heavily exploited populations or heavily fished water bodies are not recommended as biological indicators because of the potentially large confounding effects of intensive fish harvest on MeHg concentrations in such populations.

The sampling of aquatic biological indicators of different size, age, and trophic position can enhance the interpretation and understanding of temporal patterns in MeHg concentration. Concentrations of MeHg in 1-year-old prey fish will be sensitive to interannual variations in controlling factors and processes (such as mercury loadings, temperature, and hydrology), whereas older piscivorous fish will integrate and reflect temporal changes across multi-year time scales. Based on observed chronologies of mercury concentrations in aquatic biota in reservoirs during the first 30 years after flooding (Bodaly et al. 1997), we anticipate that decreases in MeHg concentrations in response to decreased atmospheric loadings would first be evident in 1-year-old prey fish, whereas decreases in concentrations in long-lived piscivores would lag a few years later.

The importance of MeHg uptake and transfer at the base of the food web is recognized and should be the subject of focused, intensive research. The utility of periphyton, phytoplankton, zooplankton, and freshwater benthic invertebrates for trend monitoring, however, is hampered by procedural and interpretational complexities associated with the large spatio-temporal variation in species assemblages, sample heterogeneity, and MeHg concentration. The utility of these groups for trend monitoring is also hampered by our very limited understanding of the processes, variables, and confounding factors that affect their bioaccumulation of MeHg.

TABLE 4.1
Recommended criteria for selection of aquatic biological indicators for monitoring and assessment of methylmercury (MeHg), and their application to candidate biological indicators

Criterion	Importance of criterion	Extent to which the aquatic biological indicator satisfies the criterion:						
		Periphyton	Phytoplankton	Zooplankton	Benthic invertebrates	Prey fish	Piscivorous fish	
Relevance to the MeHg problem	To ensure relevance to human health, ecological risk, and development of policy	Low	Low	Low	Low in fresh waters; High in coastal estuaries	Medium to High	High and direct	
Availability of historic MeHg data	To document the utility of the indicator and its probable reliability for detecting change in mercury loading	Low	Low	Low	Low to Medium	Low to Medium	High	
Recognition of extrinsic confounding factors	To facilitate the defensible interpretation of monitoring results on MeHg	Low	Low	Medium	Medium to High	Medium to High	High	
Knowledge of intrinsic co-variables	To ensure knowledge of organismal attributes that can affect MeHg concentration and complicate interpretation of results	Low	Low	Medium	Low to Medium	Medium	High	

TABLE 4.1 (continued)
Recommended criteria for selection of aquatic biological indicators for monitoring and assessment of methylmercury (MeHg), and their application to candidate biological indicators

Criterion	Importance of criterion	Extent to which the aquatic biological indicator satisfies the criterion:				
		Periphyton	Phytoplankton	Zooplankton	Benthic invertebrates	Piscivorous fish
Broad geographic distribution	To select biotic indicators that have broad spatial coverage in a regional, national, or multi-national monitoring program	Widespread, but spatially and temporally variable assemblages	Widespread, but spatially and temporally variable assemblages	Widespread, but spatially and temporally variable assemblages	Several widespread species and genera	Several widespread species and genera
Important in trophic transfer of MeHg	To select biotic indicators with a significant role in the trophic transfer of MeHg in aquatic food webs	Unclear	Important	Highly important	Highly important	Highly important
Constrained food habits or trophic position	Spatio-temporal variation in trophic position can confound and complicate interpretation of trends in MeHg concentration in the indicator	Highly constrained	Variable and complex	Variable and complex; some filter feeders	Variable and complex	Ontogenetic shifts with increasing size; less variable in adults

Temporal response to changes in abundance of MeHg	A slow rate of response to altered MeHg loading could increase the potential for interference by confounding factors, whereas a rapid response, coupled with high intraannual variability would hinder identification of multi-year trends	Hours to days	Hours to days	Hours to days	Days to seasonal in small benthos; annual to multi-annual in large benthos	Annual	Multi-annual
Effects of sampling on target population or community	To reduce effects of sampling on monitored populations and biotic communities, and to limit potential confounding biotic responses to sampling	Nonintrusive	Nonintrusive	Nonintrusive	Nonintrusive	Nonintrusive	Generally nonintrusive, except in some small lakes and streams

4.4 MONITORING AND TREND ANALYSIS

We recommend that biological indicators be sampled annually to assess responses of MeHg concentrations to changes in mercury loadings and that such sampling be limited to nonmigratory species. Although sampling at less frequent intervals (e.g., every 2 or 3 years) would decrease effort and cost per monitored water body, it would substantially reduce statistical power to detect temporal trends and would delay the detection of changes in mercury concentration (Hebert and Weseloh 2003; Bignert et al. 2004).

Freshwater piscivorous fish and prey fish should be sampled in several water bodies because data from multiple sites (i.e., clusters of sites) are needed to describe trends within a given geographic area. The primary response to decreased mercury deposition within a given area may be decreased mercury concentrations in fish, yet concentrations may remain stable or increase in some water bodies because of other factors that influence mercury cycling, MeHg production, and bioaccumulation. In Minnesota, for example, analysis of state-wide trend data on total mercury in filets of standard-sized game fish from 176 lakes showed that 49% of the lakes had statistically significant ($p < 0.05$) decreases in fish-mercury concentrations, 25% had significant increases, and 26% did not change (Bruce A. Monson, Minnesota Pollution Control Agency, St. Paul, Minnesota, personal communication). Significantly more of the Minnesota lakes exhibited decreasing mercury concentrations in fish than could be attributed to chance alone (χ^2 test, $p < 0.05$). Concentrations have also declined significantly in walleyes inhabiting boreal lakes of central Canada in the last 20 to 30 years (Johnston et al. 2003), and variation among lakes in fish-mercury trends is also evident in the Canadian data (Figure 4.3).

Concentrations of MeHg in individual fish typically increase with increasing size and age, and data on the length, weight, and age of individual fish are needed to interpret trends in mercury concentration. Piscivorous fish should, therefore, be analyzed individually. Temporal trends in mercury concentrations in piscivorous fish can be biased by variation in the size or age composition of samples taken at different times. It is, therefore, useful to estimate mercury concentrations in fish of a given age or size. In Florida, mercury concentrations in axial muscle of 3-year-old large-mouth bass (*Micropterus salmoides*) have been successfully used to examine variation in fish contamination among lakes (Lange et al. 1993) and to monitor temporal trends in the Everglades (Atkeson et al. 2003). For piscivorous fish, stratified sampling by predefined length interval could be a useful framework for sampling a target population.

One widely used approach has been to sample individual fish from a target population across a range of lengths and to apply linear regression between mercury concentration and length to estimate the concentration in a fish of some standardized length; examples include 50-cm walleye and 60-cm northern pike (Parks and Hamilton 1987; Johnston et al. 2003). This procedure yields a concentration adjusted to a specific length, along with an associated confidence interval around the estimated value. Statistical power analysis of initial data can be used to estimate the sample size (number of fish) needed to detect a change of given magnitude (Exponent 2003).

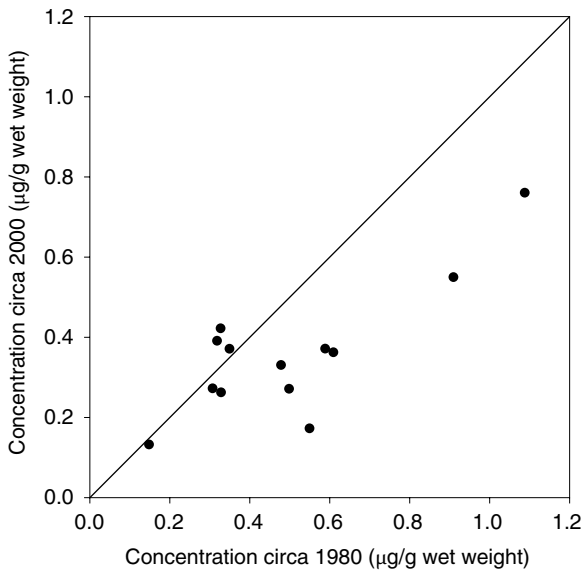


FIGURE 4.3 Recent changes in concentrations of total mercury in axial muscle of walleyes from 13 boreal lakes in northwestern Ontario and Manitoba, Canada. Standardized concentrations for 50-cm walleye sampled during 1996–2000 are plotted against standardized concentrations for fish sampled during 1977–1983 (data are for reference lakes reported in Johnston et al. 2003). Each point below the diagonal line, which has a slope of 1.0, represents a lake where the standardized concentration declined between the 2 sampling intervals.

Polynomial regression with indicator variables is another recommended statistical method for analysis of fish-mercury data. This procedure, described by Tremblay et al. (1998), allows rigorous statistical comparison of mercury-to-length relations among years and is considered superior to simple linear regression and analysis of covariance for analysis of data on mercury-length relations in fish.

For piscivorous fish, we recommend that axial muscle without skin be sampled and analyzed to avoid the additional variation that would result from differing proportions of skin and axial muscle tissue among samples. Concentrations of mercury are less in skin than in axial muscle, and the inclusion of skin in a filet can reduce the concentration of total mercury or MeHg in the sample by 5 to 12 percent (Glass et al. 1998; Serdar et al. 2001).

The historic data on total mercury in piscivorous fish are generally considered valid, and it could be useful to perform trend monitoring in carefully selected fresh waters for which historic data are acceptable and sufficient for statistical comparisons. In the late 1960s and early 1970s, the discovery of mercury-contaminated fish prompted widespread surveillance and monitoring of mercury in fish. The earliest surveys focused largely on surface waters that received mercury in polluted wastewater, but often included fish sampled from presumed “reference sites” that did not receive mercury from industrial point-source discharges. The surveillance of mercury

in fish was expanded spatially with growing awareness of the importance of atmospheric deposition as a potentially significant source of mercury in surface waters lacking on-site anthropogenic sources (Wiener et al. 2003). Historic mercury data for fish from surface waters that did not receive point-source discharges of mercury may be useful for assessing decadal trends related to atmospheric deposition. Historic fish-mercury data from analyses of composite samples (containing subsamples from multiple fish), however, has limited statistical utility for trend analysis (Exponent 2003).

We recommend that prey fish be sampled in early spring or fall, when growth is slow and temporal variation in mercury concentration is less pronounced, to facilitate comparisons among years and surface waters. Both axial muscle and whole fish have been successfully used in research applications, and analysis of either matrix would be suitable in a monitoring program for mercury. Determination of total mercury would be appropriate for muscle tissue, given that nearly all ($\geq 95\%$) of the mercury in fish muscle is MeHg. Most ($>70\%$) of the mercury in whole prey fish is MeHg (Francesconi and Lenanton 1992; Hill et al. 1996; Bodaly and Fudge 1999; Hammerschmidt et al. 1999), and we recommend that all whole prey fish be analyzed for total mercury and that a subset (from 5 to 20% of the fish, selected at random) be analyzed for MeHg.

If whole prey fish are analyzed, data should be reported as whole-body concentration and burden. Mercury *burden*, defined as the total mass of mercury accumulated in a whole fish, is calculated as the product of body weight and whole-body concentration. In prey fish of known age (e.g., age-1), mercury burden is an ecologically relevant measure of bioaccumulation, representing the mass of mercury accumulated by a fish during its residence in a monitored water body. The burden also represents the mass of mercury that a predator would ingest when eating the prey fish. Prey fish should be analyzed individually, and ancillary measurements include the total length (to nearest millimeter), fresh weight (to 0.01 g), and age of individual fish. Age can be estimated by examination of scales or other bony structures (DeVries and Frie 1996).

The trophic position of candidate benthic invertebrates for trend monitoring should be known or estimated so that samples with equivalent trophic status can be analyzed and compared. Stable-isotope methods are available for estimating trophic position (Fry 1991; Broman et al. 1992) and for assessing the influence of trophic position on mercury concentrations (Cabana and Rasmussen 1994; Kidd et al. 1995). The application of $\delta^{15}\text{N}$ and associated metrics in monitoring programs would enhance our ability to understand trends and strengthen inferences concerning linkages between trends in biotic concentrations and mercury loadings. The corrections and enhancements of Phillips (2001) and Phillips and Gregg (2001) should be considered when using data on stable isotopes to estimate the trophic status of benthic species.

The technique that has been widely applied for analyzing total mercury in aquatic biota since the late 1960s (cold vapor atomic absorption spectrophotometry) remains a valid analytical method. Accordingly, we infer that most of the historical data for total mercury in fish tissues are valid. Moreover, the historical data on total mercury concentrations in fish tissues provide defensible estimates of prior MeHg concentrations.

This is in contrast to most data collected before 1990 on total mercury and MeHg in water; most of these early data for water are not considered valid because sensitive analytical methods were not sufficiently developed and samples were typically contaminated during handling. There are some historical data on total mercury in aquatic invertebrates, but comparatively few estimates of MeHg in historic samples. Thus, the historical mercury record for piscivorous fish in North America is now about 35 years long, whereas the much sparser historical record for MeHg in water and aquatic biota in lower trophic levels is much shorter, perhaps 10 to 20 years in most regions. In comparison, the historic record of total-mercury deposition that can be examined by analyses of dated cores of sediment, peat, and glacial ice spans 2 to several centuries (Engstrom and Swain 1997; Martínez-Cortizas et al. 1999; Bindler et al. 2001; Schuster et al. 2002; Shotyk et al. 2003).

4.5 ANCILLARY DATA

The defensible interpretation of trend data in a biological monitoring program for mercury will be enhanced by the concurrent collection of relevant information on monitored airsheds, watersheds, and surface waters (see Chapters 2 and 3). The interpretation of fish-mercury data and the identification of factors causing temporal shifts in MeHg concentrations in aquatic biota would be strengthened by co-locating trend monitoring for airsheds, watersheds, and biological indicators at intensive study sites (Driscoll et al. this volume (Chapter 2), Krabbenhoft et al. this volume (Chapter 3)). Important metrics for monitored airsheds and watersheds include annual atmospheric deposition of total mercury, MeHg, and sulfate; interannual variations in rainfall and temperature; watershed area; land cover, abundance of hydrologically connected wetlands; and annual export of total mercury and MeHg from catchments to monitored surface waters. For lakes, key limnological data include lake morphometry (area, maximum depth, mean depth, percent littoral area); physicochemical characteristics of water (pH, dissolved organic carbon, sulfate, total suspended solids, chlorophyll, acid neutralizing capacity, color, and phosphorus); depth profiles of temperature and dissolved oxygen during summer stratification; hydrologic type (e.g., seepage or drainage lake); and concentrations of total mercury and MeHg in oxic water (preferably sampled when lakes are mixed). Information on the depositional chronology of total mercury in dated sediment cores would be useful for lakes with a multi-decadal, historic record of mercury concentrations in fish. Ancillary site-specific biological data include harvest of fish and shellfish; results of biotic and fishery surveys; and the age, growth rate, condition, sex, and trophic position (inferred from stable-isotope and dietary analyses) of aquatic biological indicators.

A number of recent experiments have shown that the reproduction and fitness of fish can be adversely affected by exposure to environmentally realistic concentrations of MeHg (Fjeld et al. 1998; Latif et al. 2001; Hammerschmidt et al. 2002; Drevnick and Sandheinrich 2003). In cases where trend monitoring reveals high MeHg concentrations in biological indicator organisms, subsequent field and laboratory studies should be considered to assess toxicological effects.

4.6 INTERPRETATION OF TREND-MONITORING DATA

Several biogeochemical processes, environmental factors, and human disturbances can influence the production of MeHg, its abundance in surface waters, and its concentrations in aquatic biota (Jackson 1998; Lucotte et al. 1999c; Benoit et al. 2003; Wiener et al. 2003). Thus, MeHg concentrations often vary spatially in aquatic organisms of the same species and size, even among nearby water bodies that receive similar atmospheric loadings of mercury (Wiener et al. 2003). The bioaccumulative responses of aquatic organisms to altered external loadings of mercury can, therefore, be expected to vary substantially among surface waters. Even within a given water body, internal ecosystem dynamics, anthropogenic activities unrelated to mercury loadings, and long-term external factors could confound or obscure responses in MeHg bioaccumulation to altered mercury loadings (e.g., Sorensen et al. 2005). Similarity of pertinent biogeochemical, environmental, and human factors affecting MeHg concentrations in aquatic organisms — among sampling intervals in a trend-monitoring program — would enhance our ability to detect responses to altered mercury loadings. Such similarity should not, however, be expected or assumed in a trend-monitoring program.

4.6.1 SOURCES OF VARIATION AND POTENTIAL CONFOUNDING FACTORS

Here we examine sources of variation and potential confounding factors that could obscure temporal changes in MeHg concentrations in aquatic biota in response to altered loadings of mercury. General sources of variation and potential confounding factors include interannual variations and dynamics of the monitored ecosystems, direct influences of anthropogenic and natural activities, and long-term external factors. Knowledge of such factors is prerequisite to discerning trends in MeHg concentrations in aquatic biota associated with altered loadings of mercury to aquatic systems. Awareness of potential confounding factors is also essential for minimizing or statistically accounting for their effects in a trend-monitoring program (see Chapter 3).

- 1) *Methylation and demethylation.* The methylation of inorganic mercury and demethylation of MeHg are key processes influencing the abundance of MeHg in food webs supporting the production of shellfish and fish (Benoit et al. 2003; Wiener et al. 2003). Rates of methylation and demethylation are affected by several variables, including temperature, sulfur cycling, light penetration, and the amount and quality of available organic matter (Miskimmin et al. 1992; Allan et al. 2001; Benoit et al. 2003; Kainz et al. 2003; Rencz et al. 2003). Such controlling variables can vary in response to natural events or conditions, such as water-level fluctuations and flooding, atypically cold or warm weather, increased turbidity caused by resuspension of bottom sediment, or varying abundances of allochthonous and autochthonous organic matter.
- 2) *Food-web and trophic dynamics.* Organisms at the base of aquatic food webs play a key role in the transfer of MeHg to upper trophic levels (Jackson and Harvey 1993; Plourde et al. 1997; Tremblay and Lucotte 1997; Bodaly and Fudge 1999). Within a given water body, interannual

variations in climatic and environmental conditions can affect the composition and abundance of pelagic and benthic invertebrates. Such variation in invertebrate assemblages can alter MeHg bioaccumulation in small and large fish at the end of the growing season (Parkman and Meili 1993; Montgomery et al. 2000; Gorski et al. 2003; Kainz et al. 2003).

Most fish are opportunistic organisms whose diets and trophic position can vary with food availability and size. Variation in environmental conditions among years (including water levels during spawning, availability of spawning sites, winter harshness, and summer oxygen depletion) can alter trophic structure of the fish community and the metabolism of fish, thereby affecting concentrations of MeHg in fish tissue (Lockhart et al. 1972; Meili 1991; Harris and Bodaly 1998; Scheuhammer and Graham 1999; Stafford and Haines 2001; Gorski et al. 2003). Interannual variation in growth rates of fish could influence the size-standardized mercury concentrations that are used to assess temporal trends in MeHg accumulation (Tremblay et al. 1998; Johnston et al. 2003; Sonesten 2003).

- 3) *Landscape disturbance*. Forest soils contain large inventories of mercury (Roulet et al. 1998; Lucotte et al. 1999a; Rencz et al. 2003) that can be mobilized into aquatic systems by disturbance of the forest landscape. Logging and fire can expose soil and increase the transport of mercury from soil to aquatic systems, a situation observed in northern boreal forest (Garcia and Carignan 1999; Porvari et al. 2003) and in Amazonian rainforest (Roulet et al. 1999, 2000). Agricultural practices, urban development, and road construction can also contribute to soil erosion and increased transport of mercury in soil from terrestrial to aquatic environments. Human disturbances of a watershed could also indirectly affect the production and bioaccumulation of MeHg in an aquatic system, by altering hydrologic pathways, water levels, nutrient inputs, primary production, light penetration, and trophic status.
- 4) *Fishing intensity*. Temporal variation in fishing intensity and harvest in a monitored water body could confound the analysis of trends in mercury concentrations in adult piscivorous fish. In Finnish and Scandinavian lakes, mercury concentrations in fish of given length declined notably a few years after intensive fishing had removed substantial numbers of the large adult fish (Göthberg 1983; Verta 1990). In experimental boreal lakes in Canada, the inventories of mercury in the water column and surface sediment were reduced little by intensive fishing (Surette et al. 2006). Intensive fishing may have affected the structure of the zooplankton communities (Masson and Tremblay 2002); however, the diets of the studied fishes did not change significantly after intensive fishing (Doire et al. 2002). Intensive fishing, however, significantly increased the growth rate of large predatory fish (Simoneau et al. 2005) and markedly decreased MeHg concentrations in fish (Surette et al. 2003).
- 5) *Exotic species*. The voluntary or accidental introduction of exotic species may influence the trophic structure of aquatic systems (Hrabik et al. 1998), thereby affecting MeHg concentrations in aquatic organisms in upper

trophic levels. Johnston et al. (2003), however, found that the introduction of rainbow smelt (*Osmerus mordax*) did not significantly affect concentrations of mercury in native predatory fish in lakes of central Canada.

- 6) *Reservoir impoundment.* The bioaccumulation of MeHg in aquatic organisms is greatly increased by the flooding of vegetated wetland or upland terrestrial habitats. This has been well documented in newly flooded reservoirs in temperate, boreal, and tropical regions (Bodaly et al. 1984; Johnston et al. 1991; Porvari 1995; Plourde et al. 1997; Seda and Kubecka 1997; Tremblay and Lucotte 1997; Lucotte et al. 1999b; Verdon and Tremblay 1999; Montgomery et al. 2000). Flooding of new reservoirs greatly increases the rate of mercury methylation, causing rapid increases (days to a few weeks) in concentrations of MeHg in water, zooplankton, and prey fish (Paterson et al. 1998; Bodaly and Fudge 1999). Concentrations of MeHg in the axial muscle of adult piscivorous fish can increase as much as 10-fold relative to pre-flood or reference values and remain elevated for decades after initial impoundment (Porvari 1998; Schetagne and Verdon 1999). New reservoirs also export MeHg, greatly increasing the concentrations of mercury in aquatic biota and fish inhabiting downstream aquatic environments (Johnston et al. 1991; Schetagne and Verdon 1999). In older reservoirs, concentrations of mercury in prey fish can fluctuate substantially in response to water-level fluctuations (Sorensen et al. 2005).
- 7) *Climate change.* Climate change can influence an array of environmental variables that directly and indirectly affect the biogeochemical cycling of mercury. The microbial methylation of mercury is temperature sensitive, with increasing production and concentrations of MeHg in temperate aquatic ecosystems at higher temperature (Bodaly et al. 1993). Climate change can modify the hydrological cycle, thereby altering mercury transport, transformations, and trophic structure in aquatic systems. Changes in the intensity of ultraviolet (UV) radiation at the water surface could influence both the production and photodemethylation (destruction) of MeHg in surface waters (Sellers et al. 1996, 2001; Lean and Siciliano 2003).
- 8) *Acidity and sulfate content of wet deposition.* Temporal variation in the acidity and sulfate content of wet deposition could modify the accumulation of MeHg in aquatic biota, given that numerous interactions between mercury transformations, MeHg uptake, sulfate, and aqueous pH have been reported (Spry and Wiener 1991; Odin et al. 1994; Frost et al. 1999; Scheuhammer and Graham 1999; Kelly et al. 2003). The biogeochemistry of mercury in waters with low acid neutralizing capacity and in methylating wetland environments could be affected by long-term chemical changes in wet deposition.

4.6.2 STEPS TO CONSTRAIN CONFOUNDING FACTORS AND ENHANCE INTERPRETATION

Monitoring programs should be designed to limit the effects of confounding factors that impair our ability to discern both trends in MeHg concentrations in aquatic

biota and their relation to altered atmospheric loadings of mercury. Adherence to the guidelines below should eliminate some of the pitfalls that can complicate interpretation of trend data on MeHg in aquatic biota. The application of these guidelines during the development of a monitoring program requires compilation of appropriate information on candidate study sites (surface waters and their watersheds) and their resident aquatic biota.

- 1) *Exclude very contaminated surface waters during site selection.* Concentrations of MeHg in biota inhabiting surface waters historically contaminated by wastes from industrial point-source discharges or historic mining activities can remain substantially elevated for decades (Latif et al. 2001; Wiener et al. 2003). In new reservoirs, the concentrations of MeHg in piscivorous fish remain elevated for several years after initial flooding, declining gradually over 2 to 3 decades (Bodaly et al. 1997). A trend-monitoring program intended to assess responses to changes in atmospheric loading should not include surface waters that are recovering from prior on-site elevation in abundance of either total mercury or MeHg.
- 2) *Exclude surface waters on disturbed watersheds.* Surface waters in watersheds that have recently undergone disturbance (e.g., fire or notable human development) or in watersheds where disturbance is anticipated during monitoring (e.g., logging or changing land use) should be avoided during site selection. This requires access to information on prior land use, as well as land-management plans. Given this, semi-remote surface waters in protected areas on state, provincial, or federal lands, such as national parks, would be appropriate sites for trend monitoring.
- 3) *Exclude surface waters subjected to intensive or highly variable fishing pressure.* Surface waters that are subjected to intensive fishing pressure or that are expected to undergo highly variable commercial or recreational harvest of fish or shellfish during monitoring should be excluded during site selection.
- 4) *Employ nonlethal methods when sampling small populations of piscivores.* In very small lakes or streams that support small populations, the application of procedures for nonlethal sampling (e.g., Baker et al. 2004; Peterson et al. 2005) may be advisable to avoid potential effects of removal sampling on MeHg concentrations in target populations of piscivorous fish. This determination requires advance knowledge of bioindicator organisms present at a candidate monitoring site, the approximate number (sample size) of an indicator organism to be taken during each sampling event, and the approximate size of the target population being sampled.
- 5) *Include multiple water bodies (clusters of sites) within each monitored geographic area.* Within a geographic area, the chosen biological indicators should be sampled in several water bodies of a given type (i.e., clusters of lakes or streams), because data from several sites will probably be needed to identify the *overall* trend within the area (Section 4.4). The direction of temporal trends can differ among individual water bodies because of spatio-temporal variation in other factors that influence mercury

cycling, net methylation, bioaccumulation, and concentrations in organisms. Statistical analyses should focus first on detecting temporal trends within individual sites and second on evaluating the overall direction of change across the cluster of sites within a monitored geographic area. Descriptive and exploratory statistical analyses should characterize monitored ecosystems that exhibit different trends (e.g., significant decrease, no change, or significant increase in MeHg concentrations) within a given area, as an initial step toward identifying potential causes of the observed differences in trends.

- 6) *Monitor bioaccumulation in some lakes with minimal watershed influence.* Delineating between watershed-derived mercury and atmospherically derived mercury is problematic in a biological trend-monitoring program. Wetlands can be particularly important sites of MeHg production and significant sources of total mercury, MeHg, and organic matter for adjoining surface waters (Hurley et al. 1995; St. Louis et al. 1996; Curtis 1998; Sellers et al. 2001). Concentrations of MeHg in aquatic biota should respond most rapidly to altered mercury deposition in atmospherically dominated systems, such as perched seepage lakes, which lack streams and receive little or no inflow of surface or ground water from the surrounding terrestrial environment. The production of MeHg and its bioaccumulation in such seepage lakes are therefore influenced little by watershed characteristics and processes (Watras et al. 1994; Hrabik and Watras 2002; Watras et al. 2002; Wiener et al. 2003).
- 7) *Ensure that contemporary and historic monitoring data are valid and comparable.* Mercury data from different biological monitoring programs often are not directly comparable because of variation in methods for sampling, sample preparation, and analyses. Contemporary monitoring efforts that intend to use historic data on biological indicators should take steps to ensure that 1) the historic data are valid and statistically useful and 2) the methods applied in contemporary and historic monitoring produce comparable data. Quality assurance procedures should be implemented to document the accuracy and ensure the reliability of analytical measurements. Increased uniformity in methods for sampling, sample preparation, chemical analyses, database management, and statistical summarization of data would facilitate comparison and synthesis of mercury data across geopolitical boundaries.
- 8) *Continue monitoring for a sufficient time frame.* Temporal records of mercury concentrations in aquatic biota should be sufficiently long to detect decade-scale trends that could be obscured by short-term fluctuations caused by interannual variability. This requires a commitment to sustained allocation of resources at the onset of a trend-monitoring program, for a recommended time frame of 20 to 30 years. Existing monitoring or research programs may include suitable candidate sites for monitoring, for which reliable historic data on mercury and pertinent ancillary variables are available.

Additional detailed guidance concerning the development of monitoring programs for bioaccumulative contaminants is available from the U.S. Environmental Protection Agency (2000) and from other chapters in this book (Chapters 2, 3, and 5). Provision of statistical guidance relevant to sampling design and data analyses is beyond the scope of this chapter, but such information is available from other sources (Gilbert 1987; Jorgensen and Pedersen 1994; Tremblay et al. 1998; Conquest 2000; U.S. Environmental Protection Agency 2000).

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