

Mercury in boreal freshwater fish – factors and processes governing increasing concentrations

DISSERTATION FOR THE DEGREE OF PHILOSOPHIÆ DOCTOR

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Abstract

Mercury (Hg) is a natural element, present all over the world at trace concentrations. Due to its volatility the element can undergo long-range transport in the atmosphere, and is historically accumulated in catchment soils of remote locations. Inorganic Hg can become methylated into toxic and bioaccumulative methylmercury (MeHg), which is biomagnified in aquatic food chains with potential harmful effects on organisms. Although awareness was raised concerning Hg as an environmental concern almost 6 decades ago, the complexity of the mechanisms controlling accumulation of MeHg in freshwater food chains are still largely unknown.

Due to the propensity of MeHg to accumulate, concentrations can often be low in various natural environmental matrices, e.g. water and biota at the bottom of the food chain. As is documented through studies of sample pre-treatment methods for water and biota in the present thesis, care must be taken when choosing sampling and analytical approaches to avoid erroneous results and conclusions. For water samples, using one bottle for both MeHg and total Hg (TotHg) determination, could lead to an underestimation of approximately 10 % of the TotHg concentrations. Similarly, choosing an alkaline digestion method instead of an acid extraction technique for biological material could lead to an underestimation of more than 30 % of the MeHg concentration.

In remote areas, where no local inputs of Hg exist, catchment loading of Hg to surface waters is shown to dominate over direct on-lake atmospheric Hg deposition. Hence, the factors and mechanisms controlling and affecting accumulation of Hg in freshwater fish, directly and indirectly, can be divided into three sub-groups (see *Illustration*): *i*) catchment Hg cycling; *ii*) aquatic and sediment in-lake processes; and *iii*) biological food chain processes. In the present thesis significant processes that are influencing Hg concentrations in fish are highlighted in all three sub-groups, with a specific focus on factors driving spatial and temporal trends of Hg concentrations in the aquatic phase and the food chain. Additionally, it is shown how the three sub-groups of processes are strongly interlinked and how, although on different concentration scales, processes are similar in boreal and subarctic regions.

Historically stored Hg is transported from catchment soils to surface waters with dissolved organic matter (DOM) as a transport vector. We show how variations of MeHg and TotHg

concentrations in water are strongly correlated to the concentration of DOM on a spatial scale. However, these strong spatial correlations between dissolved organic carbon (DOC), or total organic carbon (TOC), and Hg species are often not present on a temporal scale, thus highlighting the strong relationship between catchment and lake processes. This is also illustrated by the fact that reduced Hg emissions in Europe are not directly reflected in Hg food chain levels, due to catchment retention and soil accumulation of atmospheric Hg input.

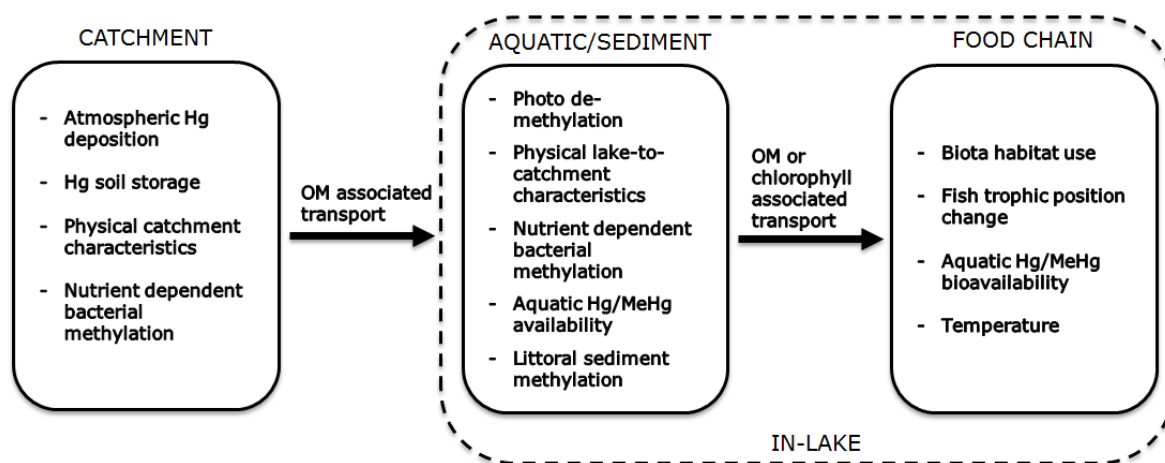


Illustration The complexity of processes involved in controlling Hg concentrations in freshwater fish species, here illustrated by the factors and mechanisms highlighted in the present thesis. Factors and mechanisms are divided into subgroups, depending on whether they occur primarily in the catchment (papers 2 and 4), in the lake (in the aquatic phase or the sediments (papers 2, 4 and 7) or in the food chain (papers 4, 5 and 6)) or whether they are responsible for transport interlinking the three subgroups (papers 2 and 6).

The catchment cycling of Hg is further complicated by the fact that in the literature wetlands have been shown to act both as sources and sinks for MeHg. We show here how intermediate nutrient status (assessed by nitrogen concentrations) in surface waters provides the highest MeHg fraction (relative to TotHg). The influence of nitrogen on methylation is likely related to bacterial methylation rather than redox processes, and is an issue that deserves more attention.

One of the most significant advancements in the understanding of in-lake Hg cycling over the last ten years is related to the de-methylation of MeHg. While methylation processes have been a focus for decades, abiotic and biotic processes of de-methylation have only recently been addressed. Surface

waters throughout Northern Europe show trends of increasing DOM levels, which leads to reduced light penetration and reduced photo de-methylation (PD). We show how DOC concentrations affect present PD of MeHg and also how it influences future MeHg budgets of pristine lake catchments in Norway. We found that, if DOC concentrations increase by 20 %, PD loss will decrease by 31 % in a humic lake.

The processes of Hg magnification through the food chain are well understood. However, the issues related to how and where MeHg enters the food chain are less known. Climate driven factors such as temperature and hydrology, as well as deposition of other elements (as nitrogen and sulphur) are thought to indirectly affect the accumulation of Hg in food chains through lake productivity, methylation rates, fish growth and changing habitat use. We show here how invertebrate habitat use and changes in fish trophic position can also significantly influence the concentrations, accumulation and magnification of MeHg in aquatic food chains. Additionally, we suggest that top predators (i.e. top-down pressure on the food chain) in these lakes could significantly change the biomagnification rates of MeHg. Together, these processes will, directly and indirectly, affect present and future concentrations of Hg in Scandinavian freshwater fish.

Although a number of mechanisms are highlighted within this thesis, we struggle to see all the possible mechanisms that are controlling the changing Hg concentrations observed in pristine freshwater fish. While we look for connections and key processes, concentrations of Hg in top predators in these pristine lakes continues to increase. In addition, concentrations vary significantly from year to year, without any clear cause, making it difficult to pinpoint the most important processes. Thousands of lakes worldwide have fish populations with Hg concentrations exceeding health advisory limits. The lack of understanding of all processes involved in controlling Hg accumulation in fish, and also how these processes interlink, limits the ability to predict future levels of Hg in fish under environmental change.

In order to further increase our understanding of what controls Hg concentrations in fish in northern ecosystems, future research needs to be focused on combined effects of climate and pollution (i.e. atmospheric deposition), as well as transport and accumulation processes of MeHg. In particular,

a better understanding of factors that drive aqueous MeHg concentrations and bioavailability is critical for improving predictions of bioaccumulation of Hg in those food chains.

List of publications

This thesis is based upon the work contained in the following papers:

Paper 1: Hans Fredrik Veiteberg Braaten, Heleen A. de Wit, Christopher Harman, Ulla Hageström and Thorjørn Larssen, 2014. *Effects of sample preparation and storage on mercury speciation in natural stream water*, International Journal of Environmental Analytical Chemistry, 94, 4, 381-384.

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Paper 4: Markus Lindholm, Heleen A. de Wit, Tor Erik Eriksen and Hans Fredrik Veiteberg Braaten, 2014. *Littoral as key habitat for mercury bioaccumulation in a humic lake*, Water, Air & Soil Pollution, 225:2141.

Paper 5: Hans Fredrik Veiteberg Braaten, Eirik Fjeld, Sigurd Rognerud, Espen Lund and Thorjørn Larssen, 2014. *Seasonal and year-to-year variation of mercury concentration in perch (*Perca fluviatilis*) in Boreal lakes*, Environmental Toxicology and Chemistry, 33, 12, 2661-2670.

Paper 6: Hans Fredrik Veiteberg Braaten, Tor Erik Eriksen, Markus Lindholm, Guttorm Christensen and Thorjørn Larssen. *Effects of water chemistry and ecology on the uptake and trophic transfer of methylmercury in boreal and subarctic Norwegian lakes*, manuscript.

Paper 7: Amanda Poste, Hans Fredrik Veiteberg Braaten, Heleen A. de Wit, Kai Sørensen and Thorjørn Larssen. *Effects of photo de-methylation on the methylmercury budget of boreal Norwegian lakes*, accepted for publication in Environmental Toxicology and Chemistry.

Abbreviations

AIC	Akaike Information Criterion
BAF	Bioaccumulation factor
BAF _Z	Zooplankton bioaccumulation factor
C/N	Carbon/nitrogen ratio
CRM	Certified reference material
CVAFS	Cold vapor atomic fluorescence spectrometry
δ ¹³ C	Ratio of heavier to lighter stable isotopes of carbon
δ ¹⁵ N	Ratio of heavier to lighter stable isotopes of nitrogen
DI	Deionized water
DOC	Dissolved organic carbon
DOM	Dissolved organic matter
DMHg	Di methylmercury
e.g.	<i>Exempli gratia</i> (for example)
EMERGE	European mountain lake ecosystems: regionalisation, diagnostic and socio-economic evaluation
EN	European Standard
FEP	Fluorinated ethylene propylene
FLPE	Fluoropolymere bottles
GC	Gas chromatography
GEM	Gaseous elemental mercury
GIS	Geographical Information System
Hg	Mercury
Hg ⁰	Elemental mercury
Hg(II)	Inorganic divalent mercury
ICD	Ice cover duration
i.e.	<i>Id est</i> (that is)
LOD	Limit of detection

m.a.s.l	Meters above sea level
MDL	Method detection limit
MeHg	Methylmercury
%MeHg	Fraction of methylmercury ($[\text{MeHg}]/[\text{TotHg}] * 100$)
MMHg	Mono methylmercury
NS	Norwegian Standard
OM	Organic matter
PAR	Photosynthetically active radiation
PD	Photo de-methylation
pH	Measure of hydronium ion concentration
PLS	Partial Least Squares
QA/QC	Quality assurance/quality control
RMSE	Root mean square error
SRB	Sulphate reducing bacteria
TMS	Trophic magnification slope
TOC	Total organic carbon
UNEP	United Nations Environmental Programme
USEPA	United States Environmental Protection Agency
UV-A/UV-B	Ultraviolet A/Ultraviolet B radiation
WHO	World Health Organisation
WMS	Web Map Services
WMO	World Meteorological Organisation

1 Introduction

Mercury (Hg) is a naturally occurring element which has a biogeochemical cycle that involves atmospheric, aquatic and terrestrial compartments throughout the world (Selin, 2009). Over the last few centuries, anthropogenic activities have altered the biogeochemical cycle of Hg (UNEP, 2002). In fact, of the more than 5700 Mg of Hg emitted into the atmosphere every year, 2320 Mg are estimated to be of direct anthropogenic origin and an additional fraction from re-emission (Pirrone et al., 2010). Environmental and health impacts of Hg are however only indirectly related to atmospheric concentrations of Hg species. It is the conversion of inorganic Hg species to the toxic and bioaccumulative organic forms, of which methylmercury (MeHg) is the most important, that is of major concern (Driscoll et al., 2013). Because of the accumulating properties of MeHg, low concentrations in the natural environment can still lead to high concentrations in the top of the food chain. So, although important in the overall budget of worldwide Hg cycling, anthropogenic activities will not be the focus of this thesis. The main goal is to identify and discuss important factors and mechanisms controlling changing Hg concentrations in freshwater environments without local Hg sources, particularly in fish.

1.1 *Hg speciation*

Identification and quantification of different species of Hg is vital to be able to ascertain toxicity, mobility and bioaccumulation within the environment. The important chemical species of Hg can be divided into elemental Hg (Hg^0), inorganic Hg and organic Hg (Leermakers et al., 2005), all of which can be exchanged in and between atmospheric, aquatic and terrestrial systems (Morel et al., 1998).

Hg^0 , or gaseous elemental Hg (GEM), is volatile, is the most stable form of Hg in the atmosphere (Schroeder et al., 1993), and can be airborne for approximately 1 year (Slemr et al., 2003). Inorganic Hg is found in oxidation state +1 and +2, where +2 (Hg(II)) is most common in the natural environment. Hg(II) is easily soluble in water and the main form of Hg in aquatic systems (Schroeder et al., 1993). Of the organic forms, mono methylmercury (MMHg, hereafter only MeHg) and di methylmercury (DMHg) are the most common forms (Tessier and Turner, 1995). MeHg is toxic and the most abundant form of Hg in most fish tissues (> 95 %, Bloom, 1992), because the specie

biomagnifies through the food chain. The biomagnifying properties of MeHg are due to the ability to accumulate in proteins faster than it is excreted (Trudel and Rasmussen, 2006).

In studies of MeHg biomagnification, the term “bioavailable forms of Hg” (i.e. bioavailability of Hg) is often used. Here, we use the term bioavailability of Hg to describe the Hg and MeHg that is available for uptake into the base of the food chain (Barkay et al., 1997). French et al. (2014), shows that the bioavailability of Hg is highly dependent on OM. In low DOC (< 8.5 mg/L) waters, Hg is mainly associated with fulvic acids and readily taken up and accumulated in the food chain. However, as DOC concentrations increase above 8.5 mg/L, Hg becomes associated with larger and less bioavailable humic acids. As we discuss later on, how Hg is bound in water (e.g. to sulphur, chloride etc.) will also affect the bioavailable and methylating properties of Hg.

1.2 Hg in freshwater ecosystems

In northern freshwater ecosystems with no direct local inputs of Hg contamination, surface water concentrations of Hg are usually low (ng/L, paper 2). In such systems, long-range transported atmospheric Hg is the main source of Hg contamination (Jackson, 1997) and has led to long-term accumulation of Hg in catchment soils (Fitzgerald et al., 1998). Because of the catchment retention, atmospheric inputs of Hg do not correlate directly to Hg in freshwaters (Larssen et al., 2008), and catchment loading of Hg dominate over direct on-lake Hg deposition (Lee et al., 1998, Lee et al., 2000). A large manipulation study in North America (The Mercury Experiment to Assess Atmospheric Loading in Canada and the United States (METAALICUS)), where Hg were added to the catchment as well as the lake, showed that an increase in Hg loading of approximately 7 times the ambient wet deposition gave increased concentrations in biota (30-40 %, including young of the year fish) over a three year period (Harris et al., 2007). Harris et al. (2007) state that “*essentially all of the increase in fish MeHg concentrations came from Hg deposited directly to the lake surface. In contrast, <1% of the Hg isotope deposited to the watershed was exported to the lake.*” Based on this, the authors suggest that lakes receiving reduced input of Hg from the atmosphere due to increased emission controls, would lower their fish Hg concentrations. The decline in the Hg content of fish would be rapid, as a result of reduced direct deposition to the lake, followed by a slow (centuries) further decline due to re-

equilibration of the catchment pools. The size of the initial response to reduced deposition will strongly depend on the lake to catchment ratio.

Since most Scandinavian lakes have a large catchment relative to the lake surface, the findings from the North American manipulation study would imply that only a small initial response to reduced atmospheric input can be expected, and the catchment pools of Hg will be of major importance compared to direct atmospheric deposition to the lake, e.g. Larssen et al. (2008), Lee et al. (2000). From Larssen et al. (2008) (and Lee et al., 2000) it is estimated that pristine catchments can contain pools of Hg 8000 (and 15500) times larger than the annual stream water output and 2000 (and 600) times larger than the input from throughfall and litterfall. The response of reduced atmospheric deposition should therefore be expected to be very slow.

In addition the slow transport of Hg through the catchment, another important reason for the often observed lack of direct relationships between atmospheric deposition of Hg and Hg concentrations in fish is the processes involved in production of MeHg in a lake-catchment system. The MeHg availability in a lake is determined by the balance between processes of methylation (production of MeHg) and de-methylation (degradation of MeHg, Benoit et al., 2003). Through methylation, inorganic Hg is transformed into toxic and bioaccumulative MeHg (Bloom, 1992). MeHg is accumulated in the aquatic food chain (Trudel and Rasmussen, 2006), and aquatic biota in northern freshwater ecosystems contain elevated concentrations of Hg, related to historical anthropogenic emissions of Hg to the atmosphere (Driscoll et al., 2013). Elevated concentrations of MeHg in aquatic food chains can potentially show harmful effects on organisms (WHO, 1991) and humans (Mergler et al., 2007) through fish consumption (UNEP, 2002).

1.2.1 Hg in freshwater fish

In thousands of North American and Scandinavian freshwater lakes, fish Hg concentrations exceed limits advised for human consumption (0.3 – 0.5 mg/kg Hg wet weight, UNEP, 2002). A compilation of multi-annual studies of Hg levels in terrestrial, freshwater and marine biota in polar and circumpolar areas in North America and Scandinavia, under coordination of the Arctic Council, suggests that neutral and rising trends of Hg are dominating (Riget et al., 2011). Riget et al. (2011)

states that data on Hg in fish covering the past one to three decades can be used to illustrate how Hg concentrations have changed in recent times and will also suggest likely near-time future trends. However, only a few time series for freshwater fish were included in the review by Riget et al. (2011).

In the present thesis the term *trend* is used to describe and illustrate how fish Hg concentrations are changing over the past three decades (1990s, 2000s and 2010s) in Norway. Increases in concentrations of Hg in freshwater fish from the 1990s onwards have been documented in Sweden (Akerblom et al., 2012), Finland (Miller et al., 2013), Norway (Fjeld and Rognerud, 2009) and Canada (Ontario, Gandhi et al., 2014), although this rising trend is not found in all regions and for all fish species. Recent studies from lakes in Sweden (Akerblom et al., 2014, Miller et al., 2013) are in fact showing declining concentrations of Hg in fish. However, despite reduced Hg emissions in several world regions (Streets et al., 2011) and reduced or unchanged atmospheric Hg deposition in Northern Europe (Wangberg et al., 2007, Harmens et al., 2008, Torseth et al., 2012) and Canada (Cole et al., 2014), there is little evidence to suggest that Hg contamination in fish is beginning to decline.

Given the mixed results on data considering changing Hg concentrations in fish, there is a clear need for more data considering year-to-year variations. In Gandhi et al. (2014), time trends were considered for different fish species (to incorporate specie-specific differences in accumulation of MeHg (Bhavsar et al., 2010)) and for different time periods (to document changing Hg trends at different decades between 1970 and 2012). It was shown that while fish Hg concentrations from 1970 to 1990 was declining, concentrations in recent decades (time periods 1985-2005 and 1995-2012) were increasing. Overall (1970-2012), patterns were shown to be neutral or declining, depending on the fish species considered (Gandhi et al., 2014a).

1.2.2 Trophic transfer of MeHg

Studies have shown that variations in MeHg exposure and uptake at the base of the food chain drive much of the variation seen in Hg concentrations at higher trophic levels (Chasar et al., 2009, de Wit et al., 2012). However, data on MeHg and dietary markers (stable carbon and nitrogen isotopes) for lower food chain compartments are lacking in the literature (Kidd et al., 2012), and little is known

regarding the environmental factors that determine the efficiency for which MeHg is taken up at the base of the food chain.

MeHg concentrations increase with trophic position (Kidd et al., 1995), calculated from the ratio of heavier to lighter stable isotopes of nitrogen ($^{15}\text{N}/^{14}\text{N} = \delta^{15}\text{N}$, Kidd et al., 1999, Peterson and Fry, 1987). The linear regression between MeHg concentrations (on a logarithmic scale) and $\delta^{15}\text{N}$ in biota describes the degree of biomagnification, i.e. the mean change in organism MeHg concentration with trophic level. The resulting Trophic Magnification Slope (TMS) is used as an indicator of the potential for biomagnification of MeHg through a food chain (Yoshinaga et al., 1992).

The ratio of stable carbon isotopes ($\delta^{13}\text{C} = ^{13}\text{C}/^{12}\text{C}$) values provide information on the major source of energy for an organism, and are used to determine which food chain the organisms belong to (Post, 2002). The three main lake habitats littoral, pelagial and profundal show contrasting quality of carbon and nutrients (Chetelat et al., 2011), leading to differences in MeHg concentrations of primary consumers depending on which zone they inhabit (Chetelat et al., 2011, paper 4). The supply of MeHg to the food chains is suggested to be affected by factors such as Hg loading (Harris et al., 2007, van der Velden et al., 2013), pH (Watras et al., 1998) and DOC (dissolved organic carbon, Rennie et al., 2005, Chasar et al., 2009).

Both physicochemical and biological factors affect MeHg bioaccumulation (and hence values of TMS). Acidity (Watras et al., 1998), concentrations of dissolved organic matter (DOM, Rolffhus et al., 2011, Chetelat et al., 2011), Hg availability (DeForest et al., 2007, de Wit et al., 2012) and lake productivity (Pickhardt et al., 2002) all affect bioaccumulation rates, as do temperature (Greenfield et al., 2001, Lavoie et al., 2013), growth rates of biota (Dittman and Driscoll, 2009), energy sources (Trudel and Rasmussen, 2006), prey contamination (Trudel and Rasmussen, 2006) and predation effects (Henderson et al., 2012, Jones et al., 2013). A global review of the environmental drivers of TMS identified latitude, DOC and productivity as important drivers, whilst a great deal of unexplained variability remained, highlighting the need for further work (Lavoie et al., 2013).

1.2.3 Hg transport, production and fate

DOM measured as DOC is the main transport vector for Hg and MeHg from catchment soils to surface waters (Grigal, 2002). Hg and other trace metals are bound to OM at the acid sites, where, for inorganic and organic Hg, the most common acidic site is thiol groups (Ravichandran, 2004, Amirbahman et al., 2002). The ionic binding between inorganic Hg (Hg^+ and Hg^{2+}) and MeHg (CH_3Hg^+) and thiol groups (reduced sulphur) in soil and aquatic OM (Ravichandran, 2004, Skjellberg et al., 2006), leads to mobilisation of Hg species from soils to streams (Mierle and Ingram, 1991) and lakes (Driscoll et al., 1995). Hence, the expression of OM as a transport vector for Hg and MeHg.

Following the arguments above, concentrations of total organic carbon (TOC) and DOC show thus strong spatial correlations with concentrations of Hg in lake surface water in Scandinavia (Meili et al., 1991, Skjellberg et al., 2003, Eklof et al., 2012) and North America (Driscoll et al., 1995, Benoit et al., 2003, Shanley et al., 2008). Fluxes of Hg in lake outlets relative to the catchment storage of Hg are usually small (Grigal, 2002, Grigal, 2003, Larssen et al., 2008), suggesting that leaching of deposited Hg from soils to surface waters is likely to continue for decades to centuries.

Processes of methylation and de-methylation in the catchment and lake determine the aqueous MeHg concentrations. Production of MeHg occurs primarily through methylation of inorganic Hg by sulphur reducing bacteria (SRB) under anoxic conditions (Morel et al., 1998), but is also shown to occur through other mechanisms (Gilmour et al., 2013). Thus the production of MeHg can take place in the catchment wetlands (St. Louis et al., 1994, Tjerngren et al., 2012b), the sediments (Benoit et al., 2003, Gilmour et al., 1998) or in the water phase itself (Xun et al., 1987).

The fraction of MeHg (as MeHg-to-TotHg ratio or %MeHg) is often used as an indicator of the environment's capability to produce MeHg (cf. methylation potential; McClain et al., 2003, Mitchell et al., 2008a). The methylation mechanism is not understood in detail, but a number of parameters have been identified as important. These parameters include the composition and activity of the microbial community, which depend on sulphur (S) chemistry, availability of inorganic Hg and OM, temperature and pH (Ullrich et al., 2001, Benoit et al., 2003). The role of OM as substrate in the methylation process is related to carbon as an electron donor when sulphate is reduced to sulphide by SRB (sometimes also Fe(III) reduced to Fe(II) by Fe reducing bacteria). The significance of both

carbon and sulphate for this process is documented through different stimulation studies, e.g. (Mitchell et al., 2008b) and (Jeremiason et al., 2006).

Factors controlling MeHg production and degradation in the aquatic environment are reviewed in (Benoit et al., 2003) and (Li and Cai, 2013). Benoit et al. (2003) states that although Hg methylation is a function of Hg concentration, the variation of methylation rates is larger than the range in Hg deposition rates, highlighting the importance of other factors as well. Of particular importance are the concentrations of sulphur and sulphide: while the SRB utilises sulphate as energy source through reduction (while oxidising carbon in OM), inorganic Hg is bound to sulphide and diffuses into the cell membrane. Hence, a pattern of increased MeHg concentrations in high methylation rate areas, are often accompanied by reduced sulphide concentrations (Benoit et al., 2003).

In addition, new studies show the importance of nutrient status on MeHg production rates in boreal wetlands (Tjerngren et al., 2012b, Tjerngren et al., 2012a). Although the idea of a nutrient influence on bacterial methylation of Hg is not new (Gilmour et al., 1998), the mechanisms behind the influence are not well understood. Tjerngren et al. (2012a) suggest that the nutrient influence is related to a higher availability of electron donors for methylating bacteria. However, Tjerngren et al. (2012a) shows that as nutrient status increases, also pH increases, and demethylation is favoured over methylation. Additional research on the influence of nutrient status on Hg cycling in general and Hg methylation in particular is clearly of great importance.

The dominant MeHg degradation process in lake systems is thought to be photo demethylation (Lehnherr and Louis, 2009).

1.2.4 Drivers of Hg in aquatic environments

In 2009 a highly significant trend towards increasing Hg concentrations in freshwater fish in boreal Norway since the 1990s, was discovered (Fjeld and Rognerud, 2009). The documented increase was surprising, as the atmospheric deposition of Hg had decreased (or showed unchanged levels) over the same period due to emission reductions in Europe (Torseth et al., 2012). Environmental features that potentially drive Hg processes in aquatic environments include catchment characteristics, lake

chemistry, climate conditions and atmospheric deposition of Hg, S and nitrogen (N), in addition to the biological features already mentioned (see *1.2.2 Trophic transfer of MeHg*).

Catchment characteristics which promote Hg leaching to freshwaters are wetlands and forests. Wetlands act as hotspots for MeHg production (Tjerngren et al., 2012b, St. Louis et al., 1996), while forests have large terrestrial Hg stores related to increased deposition from canopy scavenging of atmospheric Hg (Graydon et al., 2008). Long time-trend data of MeHg are not abundant in current literature, but records from catchments in Sweden, Finland and Canada show that temporal variations in MeHg appear to be related to hydrology and temperature driven changes in Hg methylation rates (Futter et al., 2012).

In freshwaters the elevated concentrations of Hg in fish appear to be particularly connected to humus-rich waters (Hakanson et al., 1988), which makes a connection between the recent rise in surface water DOC (Monteith et al., 2007) and increase in Hg in fish plausible, although the mechanistic explanation for this is unclear. Browning of surface waters may lead to a higher exposure of MeHg and increased energy transfer from land-derived DOC to the lower food chain, reduced MeHg in algae (Luengen et al., 2012) and reduced in-lake losses from PD (Sellers et al., 1996). Chasar et al. (2009), demonstrated that the availability of MeHg at the base of the food chain in streams is a strong determinant of MeHg in top predators. Spatial and temporal variation of MeHg in primary consumers was consistent with variations in exposure to aqueous MeHg and DOC, in addition to diet and nutrient availability in boreal streams (de Wit et al., 2012). A better understanding of factors that drive aqueous MeHg concentrations and bioavailability is therefore critical for improving predictions of bioaccumulation of Hg in the food chain.

1.3 Trends in global Hg emissions

Emissions of Hg to the atmosphere have decreased by approximately 80 % in Europe since the 1980s (Streets et al., 2011). However, due to increased emissions in Asia global emissions of Hg are currently shown to be increasing (Pirrone et al., 2010, Streets et al., 2011). Unless emission controls are widely implemented, this trend is expected to continue in the near future as a large amount of equipment phased out from industrial processes is expected to become Hg-containing waste (Pirrone

et al., 2010). In fact, a new study reveals that previously unquantified use of Hg in products and processes (so-called “commercial Hg”), has contributed a large anthropogenic source of Hg to the global environment (Horowitz et al., 2014). In November 2013, the Minamata Convention for Mercury was signed by 93 countries, aiming to protect human health and the environment from adverse effects of Hg at a global scale (UNEP, 2014).

1.4 Objectives

Following the observed increase in concentrations of Hg in freshwater fish in Norway from the early 1990s to 2008 (Fjeld and Rognerud, 2009), the main goal of this project was to confirm the trend (i.e. that 2008 was not an “outlier-year” with respect to Hg concentrations) and find the key explanatory factors and processes. In areas where no local emission of Hg exists, catchment loading of Hg is shown to dominate over direct on-lake atmospheric Hg deposition (Lee et al., 1998, Lee et al., 2000). Hence, the factors and mechanisms controlling and affecting accumulation of Hg in freshwater fish, directly and indirectly, can be divided into three sub-groups (see *Illustration*): catchment Hg cycling (1); aquatic and sediment in-lake processes (2); and biological food chain mechanisms (3). In the present project we highlighted significant processes in all three groups, with a specific focus on spatial and temporal trends of in-lake and food chain processes. Specifically, addressed are the following questions:

1. Are concentrations of Hg in freshwater fish in Norway still increasing (after 2008), and what are the potential drivers behind such a possible increase?
2. What are the key variables explaining the spatial concentration levels of Hg and MeHg, in addition to methylation potential, in Norwegian surface waters?
3. What are the main biological and physicochemical lake features, affecting the bioaccumulation and biomagnification of MeHg through boreal and subarctic lake food chains?
4. How does photochemical degradation affect concentration levels of MeHg in Norwegian surface waters today and in terms of different future DOC concentration scenarios?

Firstly, we documented the spatial distribution of TotHg, MeHg and methylation potential together with potential explanatory environmental variables in 51 Norwegian surface waters where high concentrations of Hg in fish have previously been shown to be an issue (paper 2). Secondly, a subset of the 51 lakes was used to investigate detailed mechanisms responsible for the potentially increasing Hg concentrations in fish (n = 2, paper 5), controlling factors for MeHg biomagnification (n = 4, paper 6), MeHg habitat-specific bioaccumulation (n = 1, paper 4), and the importance of present and future abiotic de-methylation of MeHg (n = 3 plus one additional lake, paper 7).

Thirdly, the importance of different sample treatment methods on analytical results were investigated for water (MeHg and TotHg, paper 1) and biota samples (MeHg, paper 3).

2 Materials and methods

2.1 Study sites

Included in the present thesis is a study of the environmental factors controlling Hg speciation and methylation potential in a total of 52 Norwegian freshwater lakes. The lakes are located in southeast and northeast Norway (Figure 1), and chosen because they represent areas where previous investigations indicate substantial concentrations of Hg in fish (Fjeld and Rognerud, 2009, Fjeld et al., 2010). In some cases fish Hg concentrations are exceeding Norwegian fish advisory limits (0.5 mg/kg, Norwegian Food Safety Authority, 2005). Of the 52 lakes we studied, 51 are included in a study of the environmental factors controlling Hg speciation in surface water (paper 2, lake ID 1-51, Figure 1). Of the 51 lakes from paper 2, we chose five lakes that were studied in more detail. Three of these are typical boreal lakes located in southeast Norway (ID 1 Breidtjern, ID 11 Tollreien and ID 32 Langtjern, paper 4, 5, 6 and 7), while the fourth lake is subarctic (ID 40 Vuorasjavri, paper 6). Additionally, one lake (ID 52 Sognsvann) was included as a clear-water lake for our PD study (paper 7).

The northern lakes ($n = 5$; ID 39 – 43) are located on a subarctic tundra plain with little topographical differences. The area is dominated by birch forest and wetlands, with average yearly air temperatures below zero (from -0.8 to -3.2 °C). The lakes in the southeast are located within generally forested catchments, dominated by coniferous tree species, with presence of wetland, and in the boreal ecotone. The mean yearly air temperature is above zero for all lakes ($n = 47$; ID 1 - 38; 44 - 52) in this area (from 1.3 to 5.8 °C).

The chosen lakes represent a wide range of physical catchment characteristics. Included potential explanatory factors for the 51 lakes included in paper 2 are elevation, lake and catchment area, lake-to-catchment ratio, wetland area and wetland-to-catchment ratio (a summary in Table 1). The surface areas of the studied lakes ranged from < 0.01 km² to 16.6 km² and the size of the catchment areas span four orders of magnitude from 0.02 km² to 268.8 km². The lakes are situated across a wide elevation range, running from 56 to 610 m.a.s.l. Seven of the southern lakes are located in close proximity, i.e. within 5 km² (Figure 1 inset; ID 32 - 38). Six of these (ID 33 - 38) are small (< 0.02 km²) and are located upstream of the seventh (ID 32). The surface area of the individual lakes,

and total wetland area, range from less than 1 % to 32 %, and from 2 % to 29 % of the total catchment area, respectively.

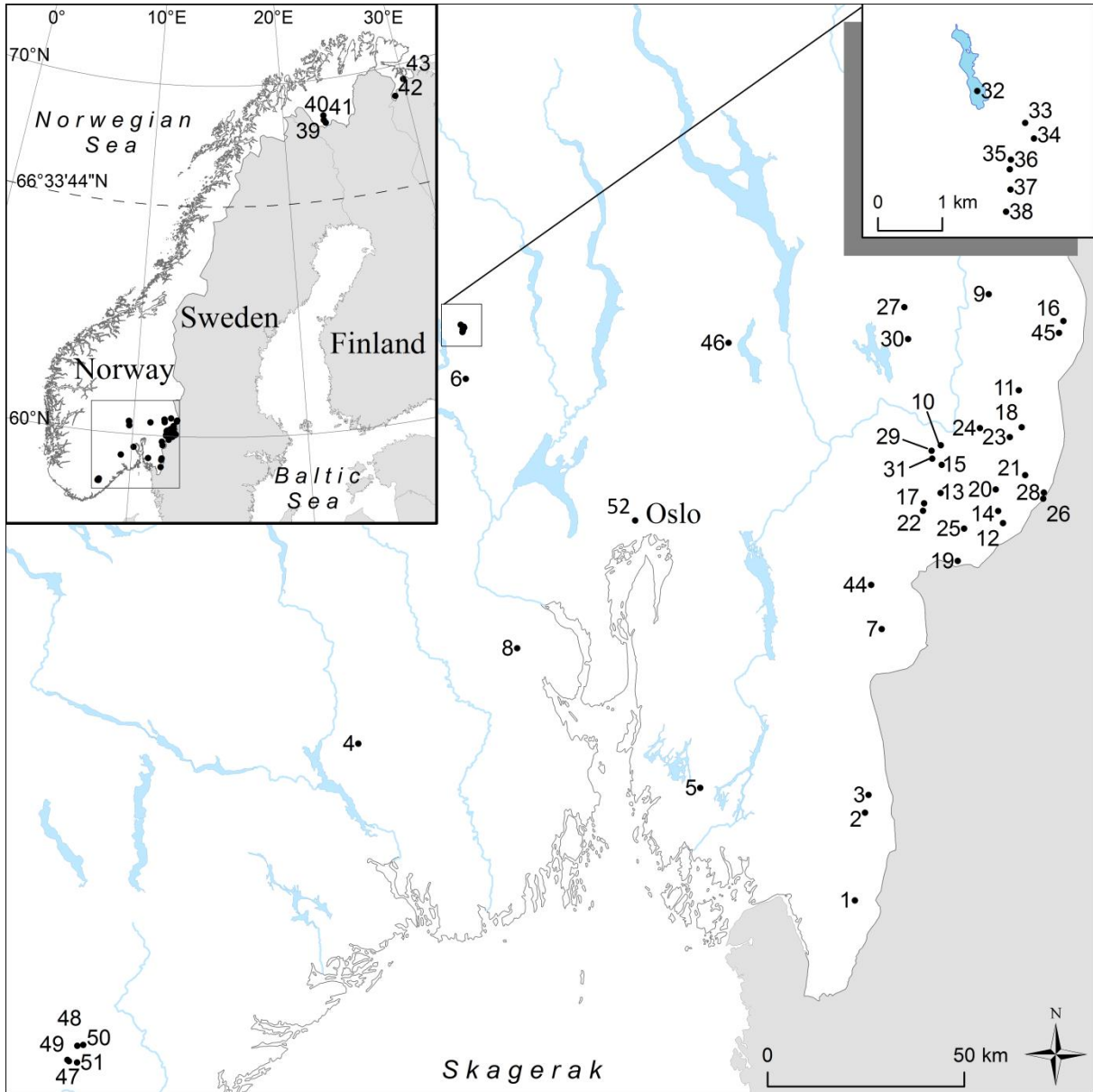


Figure 1 Geographical location of the 52 lakes included in the present study. Numbers on the map refers to lake-ID used throughout the study. The five lakes selected for more in-depth investigations are Breidtjern (ID 1), Tollreien (ID 11), Langtjern (ID 32), Vuorasjavri (ID 40) and Sognsvann (ID 52). Map modified from paper 2.

Table 1 Minimum, mean and maximum levels for all catchment characteristics, deposition patterns and climate variables included in paper 2. Data from available lakes (n = 51) are separated into lakes located in the north (n = 5, ID 39-43) and lakes located in the south (n = 46, ID 1-38, 44-51). Table copied from paper 2.

Specification	Unit	Mean value (minimum, maximum)	
		Subarctic lakes (n = 5)	Boreal lakes (n = 46)
Catchment characteristics			
Lake size	km ²	0.93 (0.20, 3.37)	0.88 (<0.01, 16.56)
Catchment size	km ²	26.67 (0.93, 60.51)	15.42 (0.02, 268.84)
Lake-to-catchment ratio	%	8.3 (0.5, 21.5)	7.4 (0.7, 31.6)
Wetland area	km ²	4.50 (0.03, 15.30)	1.14 (<0.01, 18.37)
Wetland-to-catchment ratio	%	11.4 (3.0, 25.3)	12.0 (1.7, 28.9)
Elevation	m.a.s.l	246 (56, 371)	307 (60, 610)
Deposition patterns			
Top sediment Hg	µg/g	0.16 (0.14, 0.21)	0.36 (0.30, 0.46)
N deposition	mEq/m ² /yr	10.5 (9.9, 11.9)	43.2 (33.7, 63.4)
S deposition	mEq/m ² /yr	8.0 (6.2, 10.7)	13.0 (10.3, 20.8)
Climate variables			
Run-off	mm/yr	316 (312, 324)	489 (230, 944)
Mean annual temperature	°C	-2.3 (-3.2, -0.8)	3.6 (1.3, 5.8)
Precipitation	mm/yr	372 (329, 453)	816 (653, 1182)

2.1.1 Langtjern

Langtjern (60°37' N, 9°73' E, ID 32, Figure 1), a 0.23 km² large humic lake situated at 518 m.a.s.l. in the boreal conifer forest region of southern Norway, was one of the main study site for papers 1, 4, 6 and 7. The catchment has been included in the national acid rain monitoring programme since 1972, which includes weekly monitoring of outlet chemistry for major cations and anions (Garmo et al., 2013).

The physical and chemical characteristics of Langtjern are typical for small boreal humic lakes. Maximum and mean depth in Langtjern is 12 and 2 m, respectively, and the summer thermocline is located at approximately 3 m. The catchment area amounts to 4.69 km², most of which consists of sparse pine forest (63%), mire and bogs (16%) and exposed gneiss bedrocks (16%). The lake is acidic, humic and dystrophic, with a mean annual lake outlet pH, TOC, nitrate (NO₃⁻) and total

phosphorous (Tot-P) concentration for 2009 to 2011 of respectively 5.1, 11.6 mg/L, 12 µg/L and 5 µg/L in the outlet (unpublished data). The area is acid-sensitive and acid deposition has driven the original trout population to extinction. An artificial stocked trout community is re-established, where limited numbers of farmed yearlings have been released every third year.

2.1.2 *Breidtjern and Tollreien*

Breidtjern (59°6' N 11°40' E, ID 1) and Tollreien (60°17' N 12°19' E, ID 11) are located in southeast Norway (Figure 1), representing pristine boreal areas where previous studies indicate substantial levels of Hg in freshwater fish (Fjeld and Rognerud, 2009, Fjeld et al., 2010). Both lake catchments are dominated by forest with presence of wetlands, and with no agriculture. The two lakes are different in both surface water (0.26 and 0.82 km²) and catchment area (2.1 and 34.7 km²), with Tollreien being the larger lake catchment system.

The mean yearly air temperature and precipitation is typical for southeast Norway; below 6 °C and 900 mm, respectively. Chemical deposition patterns reveal the typical south-north gradient of deposition seen in Norway, with higher deposition rates of N and S in Breidtjern (the lake located furthest south, 58.6 and 17.7 mEq/m²/yr, respectively) compared to Tollreien (35.7 and 10.8 mEq/m²/yr, respectively). On the other hand there is little difference in the top sediment Hg concentrations (0.30 and 0.33 µg/g, respectively for Breidtjern and Tollreien) and loading of Hg to the two lakes are assumed similar. This is confirmed with patterns of Hg concentrations in moss (*Hylocomium splendens*, Harmens et al., 2010).

Top consumers of the lake's food chains were perch (*Perca fluviatilis*) in Breidtjern, and perch and pike (*Esox Lucius*) in Tollreien.

2.1.3 *Vuorasjavri*

Vuorasjavri (68°58' N 23°11' E, ID 40) was selected to represent the subarctic region (Figure 1). The lake location is dominated by birch forest and wetlands. It is the largest lake (3.4 km²) and catchment (47.7 km²) of the five lakes included for in-depth analysis. The loading of Hg to the lake (top sediment Hg concentration is 0.14 µg/g), and deposition of N (10.2 mEq/m²/yr) and S (6.3 10.2 mEq/m²/yr) are

the lowest in the study, reflecting the significantly lower deposition of these compounds normally seen in northern Norway (i.e. subarctic areas). Air temperature and precipitation is also lower than in the south, representing the tundra plain described previously.

Top consumers were perch, pike, arctic charr (*Salvelinus alpinus*) and burbot (*Lota lota*) in the Vuorasjavri food chain.

2.2 *Sampling*

2.2.1 *Water sampling*

Water sampling during the ice-covered winter period (between November and April) was conducted using a water sampler (*Ruttner*, 1 L) at an approximate depth of 1 m below ice cover. The water sampler was cleaned with acid (1 % trace level grade hydrochloric acid, HCl) followed by rinses with deionized water (DI). Concentrations of TotHg and MeHg were measured in DI water added to the sampler after cleaning and the concentrations were found satisfactory (TotHg < method detection limit (MDL), MeHg < MDL). Samples collected during ice-free periods were taken at a depth of 1 m. All samples were collected using 250 mL fluoropolymere (FLPE) bottles, following ultraclean sampling procedures to avoid contamination (USEPA, 1996). Unless otherwise specified, all samples were collected at the centre of the lake.

All sampling bottles used throughout this study were previously unused and pre-tested for traces of TotHg (quality tested by Brooks Rand Labs; mean TotHg concentrations = 0.02 ng/L). As discussed in paper 1, TotHg and MeHg were sampled in individual bottles to avoid errors caused by loss of Hg during preservation (Parker and Bloom, 2005, paper 1). Samples were stored cold and kept in double plastic bags. Preservation techniques are based on United States Environmental Protection Agency (USEPA) method 1630 for MeHg (USEPA, 1998) and method 1631 for TotHg (USEPA, 2002). HCl (concentrated trace level grade, 1 mL) was added to yield a 0.4 % solution for the MeHg samples. All samples used for TotHg analysis were oxidized with bromine monochloride (BrCl) within 48 hours after sampling. Samples collected for general water chemistry were collected at the same time and depths as the Hg samples, but in individual bottles (500 – 1000 mL).

2.2.2 *Fish sampling*

Sampling of fish for papers 5 and 6 focused on populations of perch (*Perca fluviatilis*) as this species is of major relevance regarding exceeding the Norwegian recommended human consumption limits (Norwegian Food Safety Authority, 2005). Perch is also common in south east Norway and is thus easily caught in an appropriate sample number. The exception from this is Langtjern, where we collected trout as perch is not present.

We caught fish with series of gill nets (1.5 m x 25 m) of different mesh size (5 – 45 mm), so a broad distribution of fish sizes could be targeted. All fish were frozen immediately after sampling and kept at - 18 °C until analysis. Recording of fish data (length, weight and sex) and sampling of muscle tissue, otoliths and operculum were conducted according to the EMERGE (European mountain lake ecosystems: regionalisation, diagnostic and socio-economic evaluation) manual (Rosseland et al., 2001). For fish age determination we used opercula. Fish maturity stage was determined according to a method modified from (Dahl, 1917) and described in (Jonsson and Matzow, 1979).

2.2.3 *Lower food chain biota sampling*

Zooplankton (littoral and pelagic) was sampled using a 250 µm plankton haul net, towed horizontally through the upper waters (0-2 m). Composition of species was identified in each sample, while chemical analyses were conducted on pooled samples, due to small body size (specific data given in paper 4 and supporting information of paper 6). Littoral zoobenthos were collected by kick sampling, using a hand net with a frame opening of 25 x 25 cm and a mesh size of 0.5 mm that was swept through the water for 20 seconds, while walking slowly backwards and stirring the bottom substrate with the feet. The procedure followed guidelines for sampling and devices for benthic macro-invertebrates in freshwater (EN ISO 10870, 2012). Littoral samples were collected down to 1.0 m water depth, on bottom substrates made up by periphyton, particulate detritus and accumulated flocculated peaty DOM between stones and larger rocks. The zoobenthic communities were species-poor, as is common in humic boreal, and subarctic, lakes of the kind studied in this thesis.

2.3 *Chemical analysis*

2.3.1 *Water sample treatment and analysis*

The analytical method for MeHg in water was based on USEPA Method 1630 (USEPA, 1998) for determining MeHg in water by distillation, aqueous ethylation, purge and trap, and cold vapor atomic fluorescence spectrometry (CVAFS). The method for TotHg followed USEPA Method 1631 for determining Hg in water by oxidation, purge and trap and CVAFS (USEPA, 2002). The MDL was 0.02 ng/L and 0.1 ng/L (3 standard deviations of method blanks) for MeHg and TotHg, respectively. For both species automated systems were used for analysis (Brooks Rand Labs MERX automated systems with Model III AFS Detector). Due to low concentrations of particulate matter all samples were analysed unfiltered.

For every batch of Hg analysis in water (n = 24 individual samples) quality assurance and quality control (QA/QC) measures included method blanks (n = 5), blank spikes (n = 5), sample duplicates (n = 3) and matrix spikes (n = 3). The relative difference of sample duplicates was < 10 % and < 20 % for TotHg and MeHg, respectively. Recovery of blank spikes and matrix spikes were 80 – 120 % for MeHg and 90 – 110 % for TotHg.

Samples for determination of general water chemistry were analysed according to Norwegian Standard (NS) and European Standard (EN-ISO). pH was measured by potentiometry (NS4720); alkalinity was measured by titration (NS-EN-ISO9963); total N (Tot-N; NS4743), total phosphorous (Tot-P; NS-EN1189) and NO₃⁻ (NS4745) was measured by spectrophotometry; and sulphate was measured by liquid chromatography (NS-EN-ISO10304-1). TOC was measured by infrared spectrophotometry after high temperature and catalytic combustion to CO₂ (NS-EN1484).

2.3.2 *Biological analysis*

All lower food chain biota samples (except a few samples from paper 4) were extracted and analysed utilising an acid extraction method described in paper 3. The method is based on Hintelmann and Nguyen (2005). In short, samples (minimum 0.03 g) were weighed out, added 10 mL 30 % nitric acid (HNO₃) and heated at 60 °C overnight (approximately 15 hours). Before analysis the extraction solution was added 10 mL DI water. 0.050 mL extraction solution was neutralized with 0.050 mL 15

% potassium hydroxide (KOH) and ethylated before purge/trap and gas chromatography CVAFS analysis and detection as described below.

The analysis method for MeHg is based on USEPA Method 1630 (USEPA, 1998) for determining MeHg by aqueous ethylation, purge and trap, and CVAFS. As described previously, automated systems were used for analysis (Brooks Rand Labs MERX automated systems with Model III AFS Detector). For every batch of MeHg analysis (n = 30 individual samples) QA/QC measures included method blanks (n = 4), sample duplicates (n = 3), matrix spikes (n = 3) and certified reference materials (CRMs, n = 6).

Concentrations of MeHg in blank extractions were 1.0 ± 0.3 pg/mL (mean \pm 1 standard deviation). This translates to detection limits (DL) of 1.0 pg/mL or better (3 standard deviations of blank concentrations). The actual limit of detection (LOD) varies depending on the weight of sample available for analysis. For sample weights included in this study (0.02 – 0.15 g), the LOD is in the range of 0.1 – 1.0 ng/g (3 standard deviations). No sample concentrations in the present study were found to be below the LOD.

The certified MeHg concentrations of the CRMs used were 0.355 ± 0.056 mg/kg (\pm uncertainty), 0.152 ± 0.013 mg/kg and 28.09 ± 0.31 μ g/kg for DORM-3 fish protein, TORT-2 lobster hepatopancreas and SRM-2976 mussel tissue, respectively. Samples that were analysed in duplicates were also used for matrix spike samples. Samples chosen for matrix spikes were added 1000 pg (0.1 mL of 10.0 ng/mL MeHg hydroxide; MeHgOH). The relative difference of sample duplicates was always < 10 %, recovery of the CRM within 90 – 110 % and matrix spikes recovery within 75 – 125 %.

More than 90 % of Hg in fish is shown to be present as MeHg (Bloom, 1992), and Hg concentrations in fish were therefore determined as TotHg. Wet samples of muscle tissue were analysed by thermal decomposition and direct atomic absorption spectrophotometry (AAS, Lumex Mercury Analyser RA915). For every 10 samples of Hg analysis, QA/QC measures included method blanks sample duplicates (n = 2) and CRM (DORM-3 fish protein; n = 2). The relative difference of sample duplicates was always < 10 % and recovery of the CRM within 90 – 110 %. If QA/QC measures were not met, samples were re-analysed.

2.4 *Data sources*

Catchment area and wetland area were determined using Geographical Information System (GIS) software (ESRI ArcMap 10.0). The GIS software was used in combination with Web Map Services (WMS) available from The Norwegian Geo Network. Background lake data (i.e. lake size, lake identification number and elevation) were gathered from the National Lake Database of The Norwegian Water Resources and Energy Directorate (NVE).

Deposition data for S and N were supplied by The Norwegian Institute for Air Research (NILU). The data set is based on interpolated data from the period 2007 to 2011 (Aas et al., 2012; samples collected on a daily or weekly basis). Top sediment (0 – 0.5 cm) TotHg concentrations were interpolated by kriging, based on measurement of sediment TotHg in Norway during 2006 – 2008 (Skjelkvåle, 2008). Investigations of lake sediments indicated considerable enrichment of Hg in top sediments compared with preindustrial sediments, and good correlations between contents of moss Hg and Hg in top sediments, indicate that the top sediment TotHg concentrations can be used as a proxy for TotHg deposition (Fjeld et al., 1994).

Temperature and precipitation is presented as the yearly average value for each lake between 1961 and 1990, based on procedures described by World Meteorological Organisation (WMO, 1989). We chose data from the last available standard reference period in climatology as it represents the “normal” climate conditions in a specific area. The data is available from Norwegian Meteorological Institute (eKlima, 2013). Run-off was estimated for each lake based on models from NVE (Beldring et al., 2003) and show the annual average between 1961 and 1990 (NVE, 2013).

2.5 *Statistical analysis*

All statistical analyses and calculations were performed in JMP 9.0 or JMP 11.0 with a significance level $\alpha = 0.05$, unless otherwise mentioned. Utilised statistical tests and methods are described in detail in the individual papers, while statistical modelling (paper 2 and 5) are described in the following paragraphs.

2.5.1 *Spatial water data (paper 2)*

To avoid influence from non-normality and reduce heteroscedasticity in the statistical analysis from paper 2, all data variables were tested by the Shapiro-Wilks test. Variables that showed non-normality were transformed to a logarithmic scale and again tested for normality. For variables that did not show normality after a logarithmic transformation the Box and Cox transformation were used to find a power transformation that fitted the response best.

Multivariate correlations between selected variables (predictors) and responses (MeHg concentrations, TotHg concentrations and %MeHg) were explored by Pearson's correlation coefficient, r . To avoid over-fitted models due to multi co-linearity between our predictors we chose partial least squares (PLS) analysis to model and show the predictors that can best describe the spatial variations of our responses in the studied lakes. The PLS method is designed to include co-linear predictors by constructing new variables underlying the observed predictors. By doing this, most variance in the observed predictors is concentrated in the first new variables and the number of dimensions is effectively reduced (Dormann et al., 2013). The final models are represented by the goodness of fit (r^2) and the root mean square error (RMSE) of the linear regression, in addition to individual model coefficients for the selected predictors.

To test for significant differences in lake characteristics between subarctic and boreal catchments, or other groups of lakes/characteristics, Student's t-tests were used.

2.5.2 *Fish data treatment and calculations (paper 5)*

When Hg concentration in fish is to be compared between lakes, years and seasons, a length and/or age adjustment is needed due to the strong co-variation between Hg concentration and fish size (i.e. length and weight; Sonesten, 2003, Chasar et al., 2009) and hence, also age (paper 5). To investigate the Hg concentration variations, we utilised a covariance analysis creating a general linear model. Potential explanatory variables to the model included season and year of sampling, as well as the fish characteristics; length, weight, age, sex, maturity stage and $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$. To evaluate potential changes in the relationship between fish length and Hg concentrations (length*season and

length*year) and between fish age and Hg concentrations over time (age*year), interaction terms were also included in the model (also season*year). Additionally, we included the interaction term evaluating change in relationship between $\delta^{15}\text{N}$ data and Hg concentrations over time ($\delta^{15}\text{N}$ *year). Explanatory variables were chosen, evaluated and included in the model based on significance and the Akaike Information Criterion (AIC). To avoid influence from non-normality and reduce heteroscedasticity in the statistical analysis, the numerical data variables fish Hg concentrations, length, weight and age were transformed to a logarithmic scale.

3 Results and discussion

3.1 Methodological developments

Despite an increasing focus on low level methods for determination of Hg species in general and MeHg specifically in different matrices over the last decades, few studies have paid attention to direct effects and comparisons of different sample preparation methods. Important aspects of this are preservation techniques for water samples and choice of method for biota extraction and digestion.

3.1.1 Water sample preservation techniques

In paper 1 we demonstrate that different preservation techniques give significantly different concentrations of TotHg and MeHg in freshwaters (9 and 14 % on average, respectively, Figure 2). Natural stream samples from a forested lake catchment were studied. Mean stream sample concentrations of TotHg (3.6 ng/L) and MeHg (0.06 ng/L) reflect levels typical for pristine humic boreal catchments.

Two sample preparation techniques were tested, A and B. Technique A involved the use of one bottle (fluorinated ethylene propylene (FEP) 125 mL) for determining both MeHg and TotHg. These samples were preserved with HCl upon arrival at the laboratory (3-5 days after field sampling) and the analysis proceeded by the removal of a sample aliquot (25 mL) for determining MeHg first, before BrCl was added and the remainder of the sample used for determination of TotHg. Technique B involved the determination of MeHg and TotHg in two separate bottles (FLPE, 250 mL). HCl was added to the MeHg bottle just prior to sampling and BrCl to the TotHg bottle upon arrival to the laboratory.

The main causes of the observed differences in TotHg and MeHg concentrations between technique A and B is the use of one instead of two sample bottles and the timing of sample acidification, respectively. Delayed timing of sample acidification (3-5 days after sampling) could possibly cause in-bottle methylation and lead to increased MeHg concentrations (as observed for technique A). For MeHg, the analytical uncertainty value is 10 % (determined as relative percentage difference of sample duplicates), while the mean difference between the two studied sample treatment

methods is 14 %. For 80 % of the analysed samples (32 of 40) the relative percent difference between the two sample treatment methods is larger than the 10 % uncertainty.

By using only one bottle for TotHg and MeHg determination, BrCl has to be added to the sample bottle after removing an aliquot for MeHg analysis. Inorganic Hg adheres to the bottle walls during storage and is removed upon the addition of BrCl. However, if a significant percentage of the sample is removed for MeHg analysis, leaving excess inorganic Hg behind on the bottle walls, the result can be a positive bias in the TotHg concentration measured in the remaining sample (as observed for technique A).

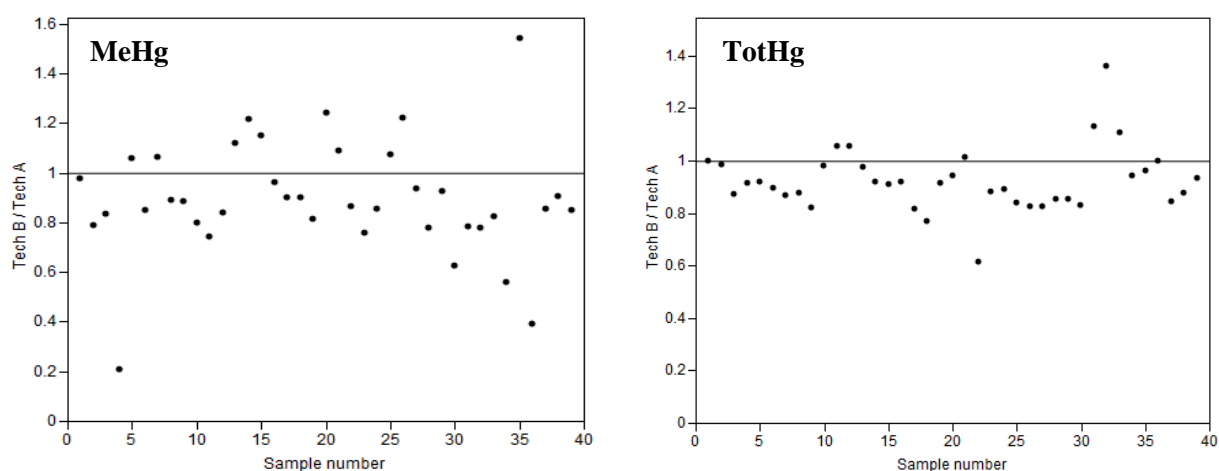


Figure 2 Levels of MeHg (left) and TotHg (right) as concentrations obtained by sample preparation technique B divided by concentrations obtained by technique A. Figure copied from paper 1.

3.1.2 Acid extraction of MeHg in biota

The most widely utilised and accepted technique for preparing biological tissue samples for the analysis of MeHg involves an alkaline digestion of the sample (Bloom, 1992, Liang et al., 1996). Recent studies suggest however, that this technique is inadequate to produce satisfactory recoveries for certain biological samples, including fish, fur, feathers and other “indicator” tissues which contain relatively high levels of MeHg (Hintelmann and Nguyen, 2005, Brooks Rand Labs, 2012). Thus an improved acidic extraction method has been proven to produce more satisfactory results for a wide range of biological tissues (Hintelmann and Nguyen, 2005, Hammerschmidt and Fitzgerald, 2008).

In paper 3 we compare the two methods on real sample material from different organisms of an Arctic marine food chain, and reveal how this could lead to misinterpretation of analytical results. Results show significantly ($p < 0.05$) lower concentrations of MeHg using alkaline digestion for large parts of the food chain, especially in fish and birds (Figure 3). The mean differences in concentrations found between the two different methods were 28, 31 and 25 % for fish (Polar and Atlantic cod), seabird (Little Auk) and seagull (Kittiwake), respectively. For samples lower in the food chain (i.e. zooplankton and krill) no significant differences were found. This leads to a clear underestimation of the levels of MeHg found higher up in these food chains, the ratio of MeHg to Hg in biological samples, and thus potentially erroneous conclusions drawn from these results concerning the biological cycling of mercury species. Specifically, this has implications for studies of MeHg biomagnification, through calculations of TMS.

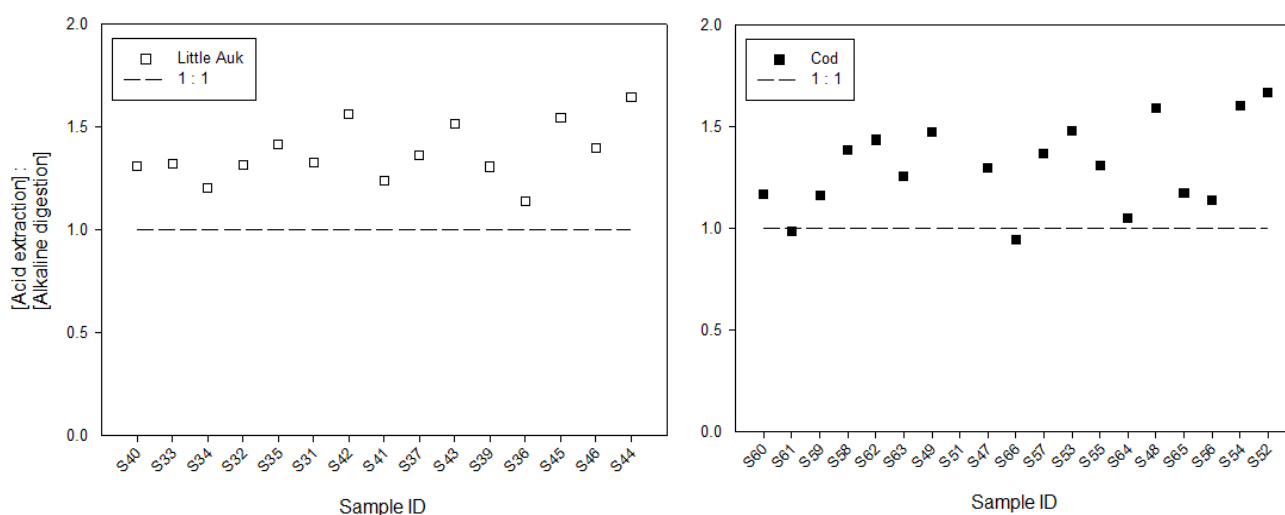


Figure 3 Levels of MeHg in the biological samples as concentration obtained by the acid extraction divided by concentrations obtained by the alkaline digestion. Figure shows Little Auk samples (left) and Polar and Atlantic cod samples (right). The dotted horizontal lines represent the 1:1 relationship between the concentrations obtained by the two sample treatment techniques ([Acid extraction]:[Alkaline digestion]). Samples are sorted by increasing concentrations of MeHg obtained by the acid extraction method from left to right. Figure modified from paper 3.

We hypothesize that the main reasons for the observed differences are poor extraction efficiency and/or matrix effects on the ethylation step prior to analysis. This is the first study to examine the effects of these artefacts on real environmental samples covering a complete food chain. Based on the results we conclude that care must be taken when choosing the sample treatment method for analysis of MeHg in biological samples, and that interpretation of results from alkaline digestions should be carried out with caution.

3.2 *Hg concentration in Norwegian freshwater fish*

Based on the increased fish Hg concentrations documented for perch in southeast Norway between the 1990s and 2008 (Fjeld and Rognerud, 2009), lakes Breidtjern and Tollreien was investigated also for the period 2010 to 2012 (Figure 1, ID 1 and ID11, respectively). Concentrations obtained from 2010, 2011 and 2012 (only autumn data considered, Figure 4) confirm the trend suggested by Fjeld and Rognerud (2009). Although concentrations in 2010 were lower in both Breidtjern (0.31 ± 0.07 mg/kg) and Tollreien (0.42 ± 0.08 mg/kg) compared to concentrations in 2008 (0.39 ± 0.08 and 0.52 ± 0.08 mg/kg, respectively), concentrations in 2011 (0.48 ± 0.09 and 0.37 ± 0.11 mg/kg, respectively) and 2012 (0.44 ± 0.08 and 0.52 ± 0.08 mg/kg, respectively) confirms the trend of increasing concentrations.

Recent literature suggests that concentrations of Hg in fish between the 1970s and today are decreasing or increasing depending on the decades of sampling (Gandhi et al., 2014). Overall (1970-2012) the data from the Great Lakes region in North America shows neutral or declining trends of Hg concentrations (depending of fish species analysed). However, broken down into shorter time periods, the data shows that the trends were decreasing in the early decades (1970-1990), while recent trends are showing increasing concentrations (1985-2005 and 1995-2012). To get a clearer picture of whether the changing concentrations observed in Norway between the 1990s and 2008 (Fjeld and Rognerud, 2009) was because of “outlier-years”, or because it was a general trend of increasing concentrations between the 1990s and today, we studied recent developments in detail.

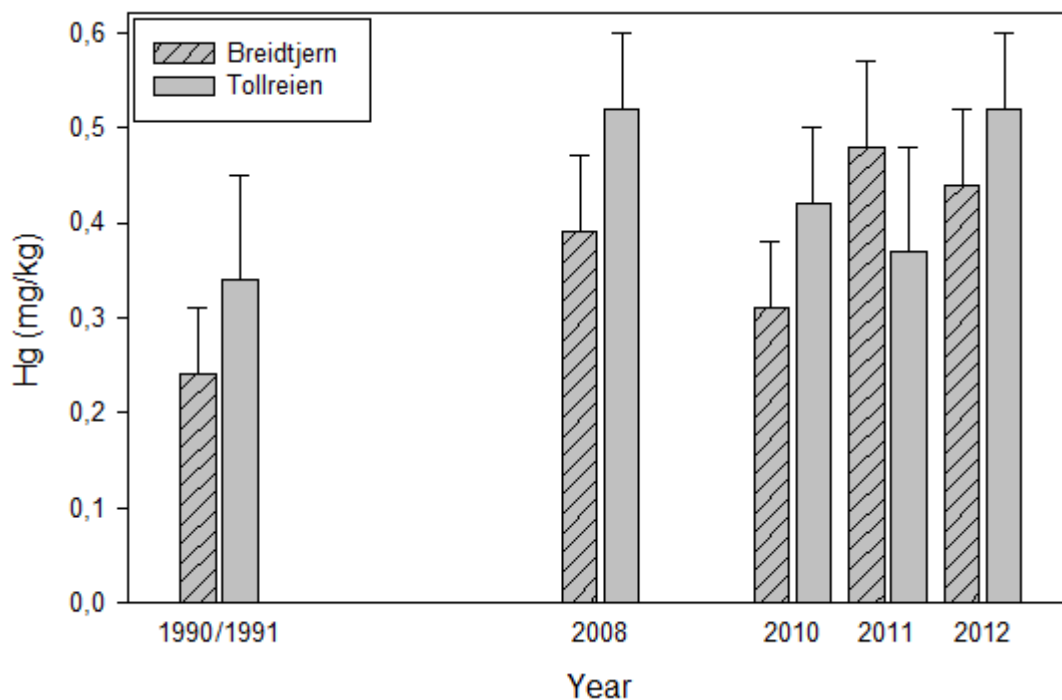


Figure 4 The yearly (autumn data considered) variations of fish Hg concentrations for the Breidtjern and Tollreien perch populations between the 1990s and 2012. Tollreien was sampled in 1990 and Breidtjern in 1991. Error bars represent +/- 95 % confidence interval. Figure is based on data from Fjeld et al., *in prep.*, and because modelling and calculations are based on a different data set (i.e. different fish morphology) than that in paper 5, fish Hg concentrations are not identical with those in Figure 5.

In paper 5 we examined the seasonal and year-to-year variations of Hg concentrations in populations of perch from Breidtjern and Tollreien. Fish Hg concentrations were determined seasonally (spring, summer, autumn) over three years (2010, 2011, 2012), to test the hypothesis that there are substantial changes in fish Hg concentrations throughout the year (seasonal variation) as well as annually. Concentrations were significantly ($p < 0.0001$) different in the two study lakes, with mean seasonal concentrations varying from 0.24 to 0.36 mg/kg (Breidtjern) and from 0.29 to 0.37 mg/kg (Tollreien, Figure 5). The Hg concentrations of both perch populations showed significant year-to-year ($p < 0.0001$) and seasonal variation ($p < 0.01$, see 3.5 for more details on explanatory variables). In both populations concentrations were highest in 2012.

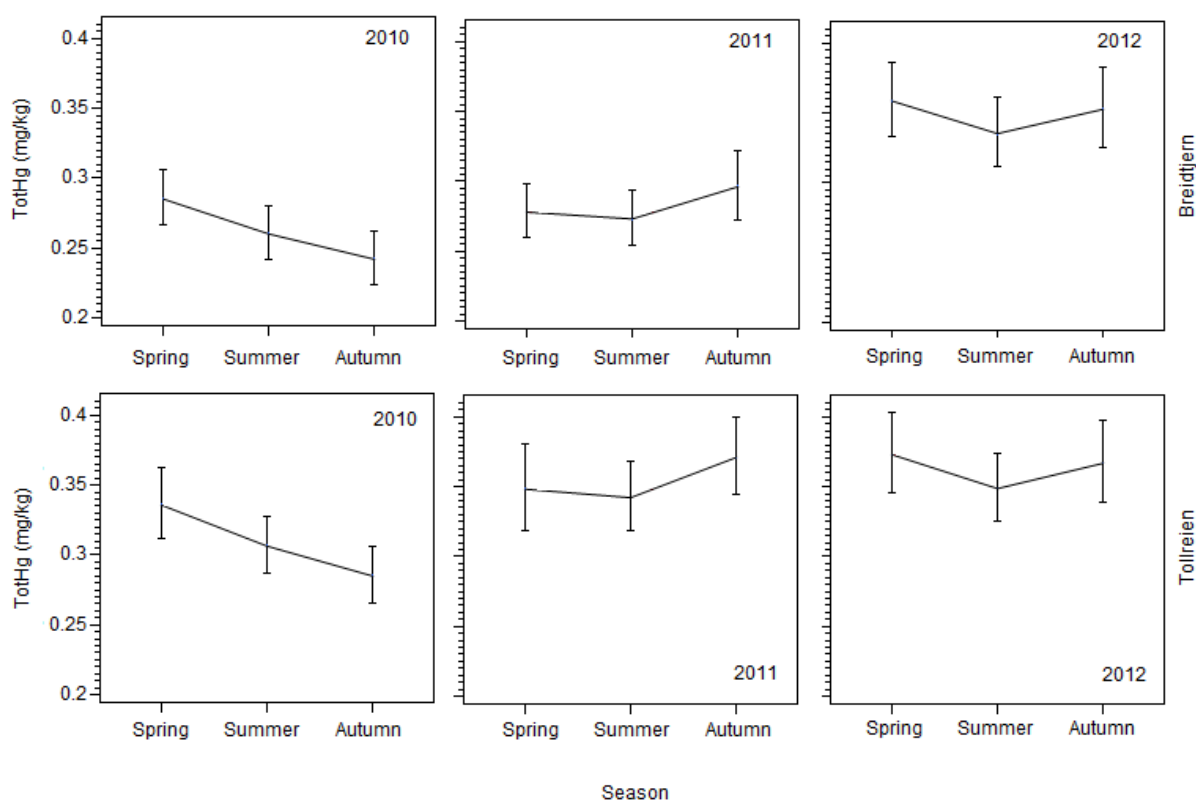


Figure 5 The seasonal (spring, summer, autumn) and year-to-year (2010, 2011, 2012) variations of fish Hg concentrations for the Breidtjern (top panels) and Tollreien (bottom panels) populations. Shown are concentrations for all available data (n = 562). Error bars represent +/- 95 % confidence interval. Figure copied from paper 5.

Paper 5 highlights the clear need for yearly studies of fish Hg concentrations, rather than the three-year cycle suggested in the European Water Framework Directive. Avoiding yearly sampling of fish may result in erroneous conclusions regarding fish Hg concentration time trends.

3.3 Catchment Hg cycling

In paper 2 we assessed the environmental drivers of TotHg concentrations, MeHg concentrations, and %MeHg in the synoptic study of 51 lakes in Norway. Concentrations of TotHg and MeHg ranged between 0.5 – 6.6 ng/L and < 0.02 – 0.70 ng/L respectively. The lakes span wide ranges of explanatory environmental variables including; water chemistry, catchment characteristics, climate

conditions and atmospheric deposition of Hg, sulphur and N. In addition to DOM-vectored transport of Hg species (see 3.3.1) and catchment base cation status (see 3.3.2), a long range of other catchment factors are previously shown to affect methylation of inorganic Hg and the surface water Hg species concentration (Bishop and Lee, 1997). This includes; the size of the catchment area (Grigal, 2002, see 3.3.3), productivity (St. Louis et al., 1996, Tjerngren et al., 2012a, St. Louis et al., 1994), the size of catchment wetlands (Eklof et al., 2012, see 3.3.4), and forestry operations (Bishop et al., 2009, Porvari et al., 2003).

3.3.1 OM as transport vector

Dissolved organic matter (DOM), measured as TOC, was the variable most strongly correlated with TotHg ($r^2 = 0.76$) and MeHg ($r^2 = 0.64$) concentrations in our study lakes (Figure 6). In several boreal lakes, DOC is shown to be the largest pool of organic C (TOC consists of > 90 % DOC; Wetzel, 2001, Hessen, 2005, Kortelainen et al., 2006, de Wit et al., 2012) and we used TOC as a measure of the concentrations of OM in the lake systems. Additionally, the supply of allochthonous DOC in humic lakes is many times higher than the production of autochthonous DOC (Hessen, 1992, Jonsson et al., 2001). Of which, the major component originates from terrestrial catchment primary production (Jansson et al., 2008, Wilkinson et al., 2013).

The relationship observed for both TotHg and MeHg with TOC has also been shown elsewhere; both in Scandinavia (Meili et al., 1991, Skyllberg et al., 2003, Eklof et al., 2012) and North America (Driscoll et al., 1995, Benoit et al., 2003, Shanley et al., 2005). The importance of this correlation is also shown by the PLS analysis (Figure 7), where TOC was the strongest positive explanatory variable for both species. The significant relationship ($p < 0.05$) between TOC concentrations and TotHg and MeHg concentrations indicates that the relationship exists independently of the other explanatory variables included in this study. In other words; independently of location (i.e. climate), deposition patterns and size of the lake-catchment system, Hg species will be transported by OM from the catchment soil to the surface water of the receiving lake.

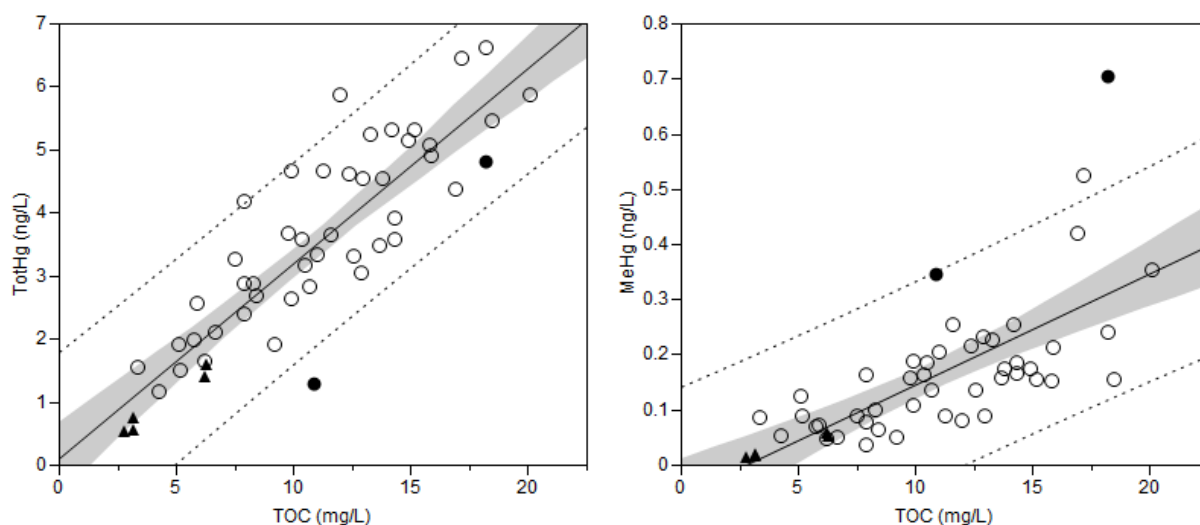


Figure 6 Scatter plots of TotHg (left) and MeHg (right) concentrations versus TOC concentrations in our study lakes (n = 51). Solid lines represent the linear regression models (TotHg = 0.11 + 0.31*TOC, $r^2 = 0.76$; MeHg = -0.06 + 0.02*TOC, $r^2 = 0.50$). The shaded area represent the confidence curve for the linear line and the broken lines the confidence curve for the individual values. Lakes from the subarctic are shown as triangles (n = 5); lakes from the boreal Ecoregion as open circles; and filled data points indicate lakes where MeHg concentrations are > 10% of TotHg (n = 2). Figure copied from paper 2.

3.3.2 Catchment base cation status

After TOC, the most significant explanatory variables in our synoptic lake study were N availability (discussed in detail under 3.3.4), base cation status, lake size and catchment area. Both pH and alkalinity were shown to be significant explanatory variables for TotHg concentrations, MeHg concentrations, and %MeHg in the PLS analysis (Figure 7). TOC, pH and alkalinity are strongly internally correlated (see paper 2 for Pearson's correlations). Such internal correlations could hamper an interpretation of independent effects of these variables on TotHg concentrations, because they have opposite effects on TotHg (consistent with the sign of their internal correlation, Figure 7). However, for MeHg and %MeHg the correlations with TOC, pH and alkalinity are all positive, which could imply that TOC and pH/alkalinity are separate controls for MeHg and %MeHg. With lake water pH and alkalinity and catchment base cation status being correlated (Pennanen et al., 1998), a possible

interpretation is that the microbial activity is stimulated in soils with lower acidity (i.e. higher pH) and higher base cation status (Mulder et al., 2001, Oulehle et al., 2006). Higher MeHg production is a possible side effect of this stimulation, as increased activity of the SRB community has been shown to increase the MeHg production (Ullrich et al., 2001). However, the effect of pH on methylation is debated, with studies showing both increased (Gilmour and Henry, 1991) and decreased (Steffan et al., 1988) methylation rates under low pH conditions.

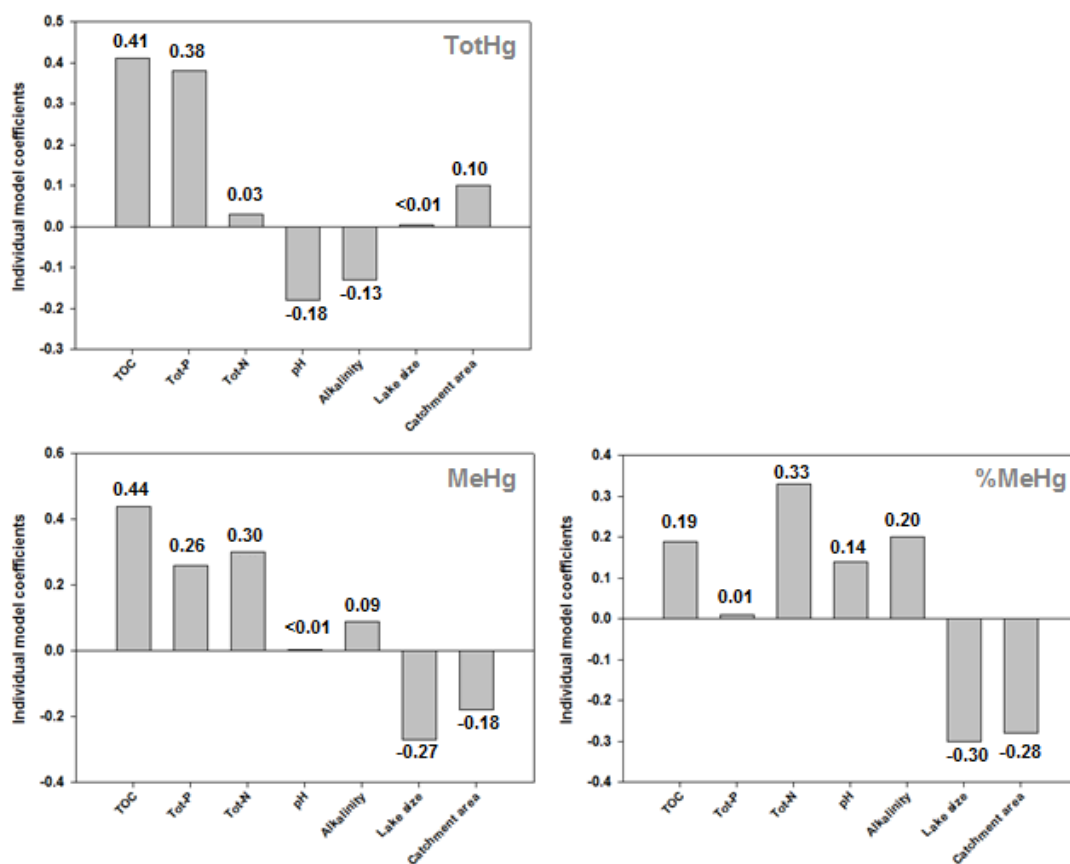


Figure 7 Individual model coefficients from the PLS analysis for each selected predictor for the responses: TotHg concentrations (top left), MeHg concentrations (left) and %MeHg (right). The predictors shown are total organic carbon (TOC), total phosphorous (Tot-P), total nitrogen (Tot-N), pH, alkalinity, lake size and catchment area. Figure copied from paper 2.

3.3.3 Catchment area

In our synoptic study (paper 2), concentrations of TotHg were not significantly related to neither lake size nor catchment area ($p > 0.05$). However, we found a significant negative relationship between TotHg and lake-catchment ratio ($r = -0.41$, $p < 0.01$). This is consistent with the idea that catchment loading of Hg dominates over direct on-lake Hg deposition (Lee et al., 1998, Lee et al., 2000). The larger the catchment compared to the lake area, the larger this effect is. MeHg concentrations and %MeHg were significantly negatively influenced by both lake size ($r = -0.58$ and $r = -0.54$, respectively, both $p < 0.01$) and catchment area ($r = -0.44$ and $r = -0.51$, respectively, both $p < 0.01$, Figure 7), but not by the lake-to-catchment ratio ($r = -0.24$, $p = 0.10$ and $r = -0.01$, $p = 0.92$, respectively). We suggest that the effect of lake size and catchment area could be related to the amount of surface water in the catchment, including both streamwater and the lake itself, where loss of MeHg by PD (Sellers et al., 1996) contributes to decrease MeHg leached from catchment soils and wetlands. The influence of PD on lake MeHg concentrations and future MeHg availability is discussed in detail further on (3.4.3 PD of MeHg).

3.3.4 Nutrient mediated methylation

The main difference between significant explanatory variables for TotHg and MeHg concentrations in paper 2 was Tot-N (Figure 7). While Tot-N concentrations were strongly positively correlated with both concentrations of MeHg ($r = 0.34$, $p < 0.01$) and %MeHg ($r = 0.40$, $p < 0.01$), correlations with TotHg were not significant ($r = -0.03$, $p = 0.83$). We tested other indicators of lake nutrient status (i.e. NO_3^- concentrations, NO_3^- -to-Tot-N ratios and NO_3^- -to-Tot-P ratios; (Bergstrom et al., 2008), in addition to C/N ratios) without finding similar relations with MeHg and %MeHg. Still, Tot-N is an indicator of total N availability and therefore we suggest that methylation is stimulated by N availability. To our knowledge, no previous study has shown a similar influence of N on methylation of Hg in boreal lakes.

A recent study of Hg methylation in wetlands from Sweden (Tjerngren et al., 2012a) indicated that intermediate levels of nutrient status (measured as C/N ratios in the soil and NO_3^- in outlet stream waters) give the highest MeHg production rates. This is consistent with the two lakes that had ratios of

%MeHg outside the 1.5*interquartile range (14.7 and 27.1 %) in the present study (Figure 6). Both lakes had intermediate concentrations of Tot-N (415 and 455 µg/L) and nitrate (43 and 57 µg/L). We did not find support for a relationship between concentrations of NO_3^- and MeHg elsewhere in our dataset however.

In contrast to our study, negative relationships were found between NO_3^- and MeHg concentrations in studies of the water column (Todorova et al., 2009) and sediments (Matthews et al., 2013) of a seasonally stratified, though contaminated lake in North America. The authors show that high concentrations of NO_3^- suppress MeHg accumulation and interpret this as an effect of NO_3^- outcompeting sulphate as electron acceptor for NO_3^- -reducing microorganisms. Further, the authors hypothesize that a negative NO_3^- control of MeHg production could occur in remote areas impacted by atmospheric Hg and N deposition.

In a study from the marine environment, nutrient loading (of mainly N) affected Hg contamination by reducing bioavailability and trophic transfer (Driscoll et al., 2012). Driscoll et al. (2012) conclude that a better understanding of the linkages between nutrient loading and Hg contamination is needed. Another marine study (Zhang et al., 2013) indicates significant relationships of both N and P with MeHg and TotHg concentrations in surface sediments. The authors do not however, provide an explanation other than a link to the OM of the sediments. The relationship between MeHg and N is also demonstrated in pore water and sediments of polluted reservoirs (He et al., 2008). Our study indicates that relationships between methylation and nutrient status are poorly understood and deserve more attention. Based on paper 2 we cannot conclude as to whether the process of nutrient influenced methylation occurs in the catchment or the lake water phase.

3.4 *Aquatic in-lake processes*

In addition to the water chemistry parameters we investigated in paper 2, we also explored aquatic in-lake processes in paper 7 (PD of MeHg), paper 4 (habitat specific methylation) and paper 6 (chlorophyll versus TOC associated transport of MeHg from the water to the food chain).

3.4.1 *OM as methylation substrate*

While the spatial study of Hg speciation in surface waters (paper 2) revealed a relatively strong correlation between TOC and MeHg, this correlation is often not present at a temporal scale (paper 1, data not shown). This is related to the fact that DOM, in addition to being a Hg transport vector, also has an additional influence on MeHg concentrations. This is evident through the fact that the simple linear regression we performed in paper 2 revealed no significant correlation between %MeHg and TOC concentrations ($r = 0.20$, $p > 0.05$). The PLS analysis did however show that TOC had a significant positive influence on the %MeHg, but the relationship was weaker than for both TotHg and MeHg concentrations (Figure 7). The significant positive correlation for both TotHg and MeHg concentrations with TOC concentrations is likely to be related to DOM as a transport vector for Hg species from the catchment to the surface water (see 3.3.1). However, DOM is also a necessary factor in the production of MeHg as a substrate for methylation (Ullrich et al., 2001). Possibly, this explains the weaker, but still positive, influence of TOC concentrations on %MeHg.

3.4.2 *PD of MeHg*

In paper 7 we assessed the importance of PD for the MeHg concentrations in Breidtjern, Tollreien, Sognsvann and Langtjern. We measured light attenuation coefficients for photosynthetically active radiation (PAR) and ultraviolet radiation (UV-A and UV-B), and found that values differed strongly between the study lakes (Langtjern and Sognsvann presented in Figure 8). Much more rapid attenuation of light (for all wavelengths) was documented in the three humic study lakes (Breidtjern, Tollreien and Langtjern) than in clear-water Sognsvann. We found close agreement between measured attenuation coefficients for PAR, UV-A and UV-B and attenuation coefficients calculated based on published relationships between DOC, chlorophyll *a* and attenuation (Morris et al., 1995). Together this highlights the importance of OM in determining the attenuation of light in these boreal lakes. In unproductive lakes with minimal suspended particulate matter, similar to the ones in the present study, DOM can be expected to dominate the light absorption (as previously observed in several studies, including Thrane et al., 2014 and Morris et al., 1995).

We also calculated the whole lake loss of MeHg for Langtjern and Sognsvann (cumulative loss through the water profile shown in Figure 8). Langtjern and Sognsvann are the study lakes with the highest and lowest concentrations of TOC in paper 7 (12.0 and 3.9 mg/L, respectively). Both lakes show highest loss during summer months, due to seasonality in incident PAR flux (data shown in paper 7). Maximum loss of MeHg was observed for June, with losses of 71 and 36 ng/m² for Langtjern and Sognsvann, respectively. The difference in PD loss between the two study sites is in part a reflection of differences in lake water MeHg concentrations, given that Langtjern has 4-fold higher MeHg concentrations than Sognsvann. However, despite this 4-fold difference in MeHg concentrations, there was only a 2-fold difference in MeHg loss between the two lakes, due to higher attenuation of light in Langtjern.

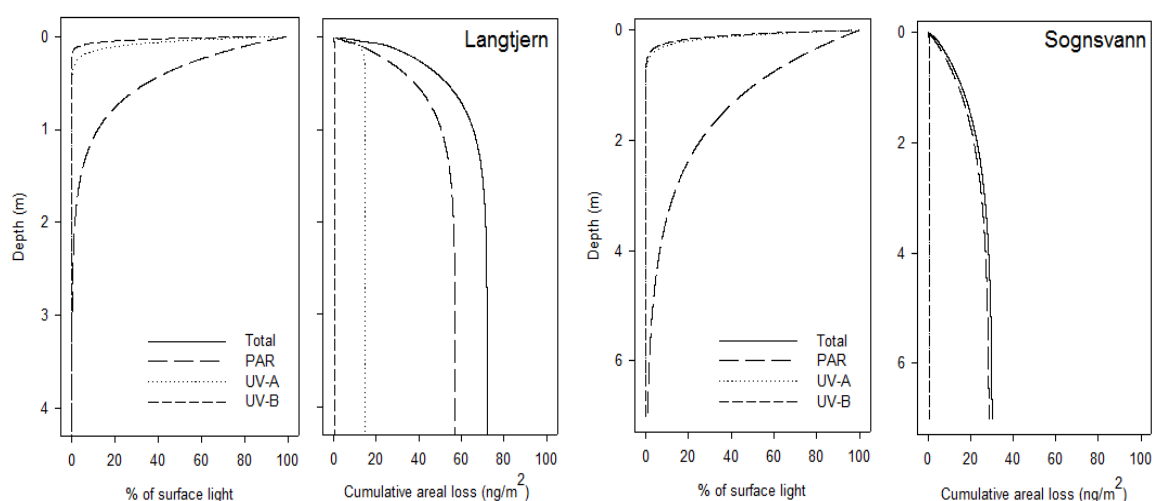


Figure 8 Attenuation (as % of surface light, x-axis) and cumulative areal loss (ng/m², x-axis) of PAR (long dashed lines), UV-A (dotted lines) and UV-B (short dashed lines) versus depth (y-axis, m) for Langtjern (left panels) and Sognsvann (right panels). Shown is also total cumulative areal loss (solid line). Note the different scaling on the y-axis for the two lakes. Figure copied from paper 7.

Our estimated annual whole lake losses of MeHg during the year (excluding periods of ice cover) were, 68 and 63 mg/year for Langtjern and Sognsvann respectively. For Langtjern, detailed information is available regarding Hg inputs, cycling and outputs (see paper 7 for details). Our estimate of annual loss through PD for Langtjern suggests that nearly 27 % of the annual inputs (253

mg/yr) are lost to PD processes during the year (Table 2). Indeed, our estimates indicate that nearly 6 % (16 mg) is lost in June alone. Mean annual losses of MeHg from Langtjern through the lake outflow were estimated to be 265 mg, 5 % higher than the estimated annual inputs. Given the substantial PD losses that we have calculated for Langtjern, the difference in MeHg concentrations between inflow and outflow suggests that there may be substantial methylation of Hg taking place within this lake.

3.4.3 Future PD loss scenarios

In many northern boreal regions, climate change and decreases in atmospheric deposition of sulphate are expected to drive strong increases in DOC export from terrestrial catchments to surface waters (Monteith et al., 2007, Larsen et al., 2011). Additionally, climate change is expected to lead to reduced ice cover duration (ICD) in boreal lakes (Magnuson et al., 2000), with southern Norwegian lakes expected to experience a reduction in ICD of approximately two weeks by 2100. Based on this, we calculated future PD losses for Langtjern and Sognsvann (only Langtjern shown here, Table 2) based on three scenarios: A) a 20 % increase in TOC concentration, B) a 20 % increase in TOC concentration paired with a 20 % increase in MeHg concentrations (since MeHg concentrations are shown to be significantly correlated with TOC concentrations in Norwegian lakes (paper 2)), and C) scenario B along with a reduction in ICD based on estimated ICD for the year 2100.

Due to reduced light penetration, Langtjern PD losses under scenario A (47 mg/yr) were 31 % lower than current values (68 mg/yr), indicating that increases in DOC loading to lakes can be expected to strongly reduce PD of MeHg. In scenario B we tested the cumulative effects of a 20 % increase in DOC paired with a 20 % increase in MeHg concentrations. In this scenario, increased MeHg concentrations led to a 21 % increase (57 mg/yr) in areal PD loss relative to Scenario A. However, if we assume that catchment MeHg inputs have increased 20 %, loss of MeHg through PD relative to total inputs would be 22 % for Langtjern. These proportional losses are lower in Scenario B than for the current situation (27 %), despite higher total inputs, suggesting that under Scenario B, we may expect higher aqueous MeHg concentrations due to a combination of increased inputs and reduced losses. It should also be noted that increased DOC inputs may also lead to increased Hg

methylation due to availability of substrate for methylation and increased anoxia (Ullrich et al., 2001), which could act to further increase aqueous MeHg concentrations.

When future reduced ICD was considered, along with potential increases in DOC and MeHg concentrations (Scenario C), we found that due to changes in ICD, PD losses would increase to 61 mg/yr (7 % increase calculated relative to Scenario B, 30 % increase calculated relative to Scenario A). However, loss of MeHg through PD relative to total estimated inputs (Scenario C: 24 %) were very similar to those observed for Scenario B (22 %), suggesting that reduced ICD will not offset the negative effects of increased DOC loading on PD losses.

Table 2 The scenario parameters DOC concentrations, MeHg concentrations and ICD for study catchment Langtjern with current and future calculated PD loss (both absolute values and relative to the total annual input of 253 mg/yr). Shown are future PD loss scenarios A (20 % increase in DOC concentrations), B (scenario A plus 20 % increase in MeHg concentrations) and C (scenario B and year 2100 ICD estimate). Table modified from paper 7.

Specification (Langtjern)	Units	Current situation	Scenario A	Scenario B	Scenario C
Scenario parameters					
DOC	mg/L	12.0	14.4	14.4	14.4
MeHg	ng/L	0.08	0.08	0.10	0.10
ICD	days	185	185	185	168
Loss (PD)	mg/yr	68	47	57	61
Loss (ratio PD of total input)	%	27	19	23	24

Combined, the future scenarios A, B and C highlight the importance of DOC-related light attenuation in driving losses of MeHg through PD. The data suggest that future increases in DOM loading to boreal aquatic ecosystems may lead to a shift in the balance between inputs and PD-related losses, with the potential for higher aqueous MeHg concentrations. However, the effects of DOC on uptake and trophic transfer of MeHg in aquatic food chains are highly complex, and it is difficult to predict how DOC-related shifts in the aqueous MeHg balance of lakes will affect concentrations in the food chain. There is evidence that DOC can reduce availability and phytoplankton uptake of MeHg

(Luengen et al., 2012). Additionally, French et al. (2014) report a unimodal response of invertebrate MeHg bioaccumulation to DOC concentrations, with increasing MeHg bioaccumulation up to a threshold of ~8.5 mg C/L and decreased bioaccumulation above this threshold. DOC-related ecological changes may also influence MeHg uptake and concentrations at higher trophic levels through changes in primary and bacterial productivity (Vallieres et al., 2008). In particular, increased importance of bacterial food sources has been shown to lead to elevated food chain concentrations of biomagnifying compounds such as MeHg (de Wit et al., 2012).

3.4.4 Habitat specific in-lake methylation

The relation between high Hg concentrations in aquatic organisms and coloured lakes is well acknowledged (Hakanson et al., 1988), but less is known about how in-lake variation in MeHg relates to habitat and dietary uptake to organisms. Concentration of MeHg in different habitats and associated food chains may vary because of habitat characteristics that determine methylation and MeHg transfer. There are strong variations in Hg bioaccumulation rates in lake food chains which are yet poorly explained, but are believed to depend on physical and chemical lake characteristics (Clayden et al., 2013). Few studies have combined focus on Hg concentrations in organisms and sediments in coloured lakes. The three main lake habitats littoral, pelagial and profundal offer contrasting quality and nutrients for their primary consumers (Chetelat et al., 2011, Vadeboncoeur et al., 2002).

The relation between habitat-dependent energy sources and MeHg remains unclear. Profundal sediments are usually viewed as hot spots for MeHg, through methylation of inorganic Hg by SRB under anoxic conditions (Morel et al., 1998, Benoit et al., 2003, Gilmour et al., 1998). In paper 4 we found however, that in Langtjern MeHg in primary consumers increased from profundal to littoral, a pattern reflected by surface sediments concentrations (Figure 9). These findings confirm the studies by Kainz et al. (2003) and Ethier et al. (2010), where higher MeHg concentrations were found in littoral compared to profundal sediments. Moreover, the methylation potential (expressed as the ratio of MeHg to TotHg, %MeHg) was lower in profundal than in littoral sediments, suggesting that littoral sediments have higher net methylation rates.

High MeHg concentrations in littoral primary consumers and sediments suggest that shallow lake sediments are important for MeHg transfer to the aquatic food chain in boreal humic lakes (see 3.4.5 for more on transfer of MeHg to boreal and subarctic food chains). Lake morphometry, specifically the fraction of littoral, is hence likely to add to differences in MeHg bioaccumulation rates in lake food chains. For Langtjern, a small and relatively shallow lake with short water residence time (mean is 72 days for the period 1973-2012, Couture et al. *in review*), there is a high degree of water-sediment contact due to the frequently exchange of water. This is of importance for the methylation of Hg because maximal net methylation is often observed in surface sediments (Ramlal et al., 1993), due to constant input of fresh OM (Benoit et al., 2003). This suggests that elevated concentrations of MeHg in the littoral Langtjern sediments could be exchanged with overlying water and readily become bioavailable for the food chain through biota in the aquatic phase in addition to biota in the top sediments.

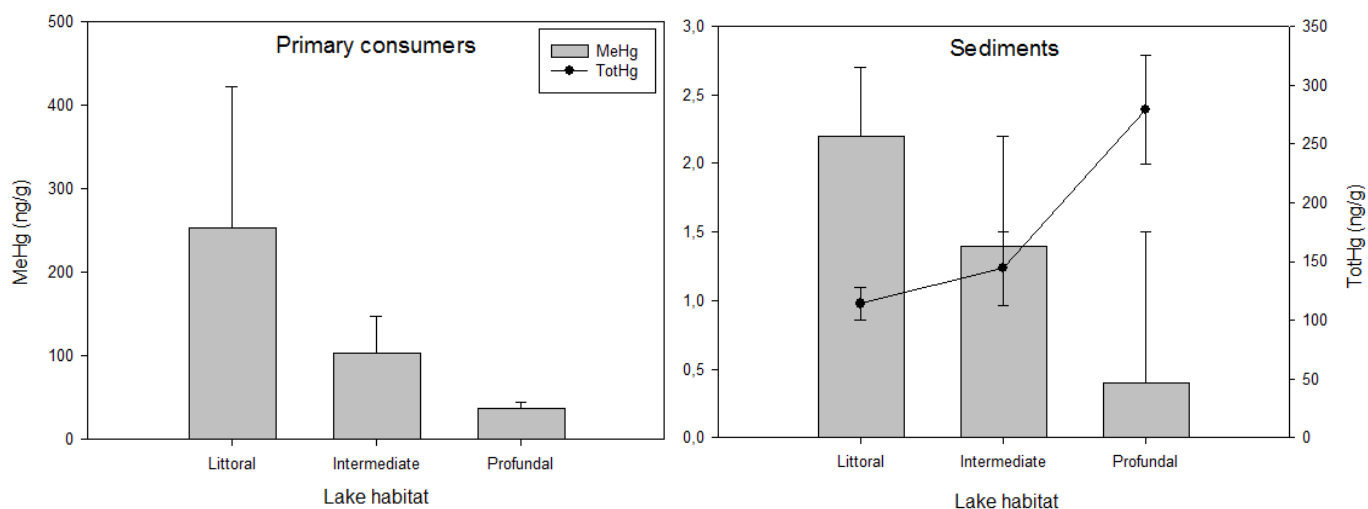


Figure 9 MeHg concentrations in primary consumers (ng MeHg/g, dry weight (dw), left panel) and MeHg and TotHg concentrations in sediments (ng MeHg/g, dw, on x-axes and ng TotHg/g, dw, on y-axes, right panel) of the three ecological main compartments in Langtjern. Error bars represent \pm one standard deviation on mean concentrations. Figure modified from paper 4.

The observed higher MeHg concentrations in littoral consumers relative to the profundal consumers may have a significant impact for the higher trophic level species, particularly relevant for perch and other fish species thought to feed largely on benthic organisms in the littoral zone (Hjelm et al., 2000, paper 5).

3.4.5 Chlorophyll and TOC associated MeHg transport

MeHg concentrations in zooplankton from Breidtjern (ID 1), Tollreien (ID 11), Langtjern (ID 32) and Vuorasjavri (ID 40, paper 6) reflect the aquatic MeHg concentrations in the epilimnion water and thus correspond with the assumption that MeHg concentrations at the bottom of the food chain are determined by the concentrations in the water column (Chasar et al., 2009). In paper 6, we also observed a higher zooplankton bioaccumulation factor (BAF_z , [MeHg in organism, ng/kg] / [MeHg in water, ng/L]) in Vuorasjavri (the subarctic lake, 4.3 L/kg) compared to Breidtjern, Tollreien and Langtjern (the boreal lakes, 0.7 – 1.1 L/kg). This is consistent with the lower aquatic concentrations of TotHg and MeHg in subarctic lakes compared to boreal lakes (paper 2), and more similar MeHg concentrations in the primary and secondary consumers from the two regions (paper 6). However, as fish Hg concentrations also are shown to be lower in the subarctic (Fjeld and Rognerud, 2009), it is evident that also other factors affect accumulation of MeHg in fish in the subarctic (see 3.5 *Biological food chain mechanisms*).

In the literature, bioconcentration of MeHg is shown to be significantly higher in low DOC lake water (< 5 mg/L) compared to those in higher DOC water (Gorski et al., 2008). This is clearly evident also in the present study, where we observe that the linear regression slope of BAF versus baseline corrected $\delta^{15}N$ values are higher (1.6) in the subarctic lake compared to the boreal lakes (all < 0.8, Figure 10). In French et al. (2014), it is shown that MeHg BAF increases with increasing TOC concentrations as long as TOC < 8.5 mg/L. Above this concentration limit, MeHg BAF is decreasing. Since TOC concentrations are low in the subarctic lake of the present study (paper 6), aquatic MeHg transfer to zooplankton is most likely chlorophyll associated rather than the detritus subsidized transport in the boreal lakes (Kainz et al., 2008). This fits well with the observed high BAF_z in Vuorasjavri (4.3 L/kg). Additionally, mean $\delta^{13}C$ levels were significantly higher (less negative) in the

subarctic lake compared to values in the three boreal lakes, indicating differences in energy source quality (see paper 6 for details). This again fits well with the observed food chain structures (i.e. high densities of Lymnaeidae). Although water Hg species concentrations are low, our results show that low TOC concentrations promote relatively high bioaccumulation rates of MeHg also in subarctic lakes (Figure 10).

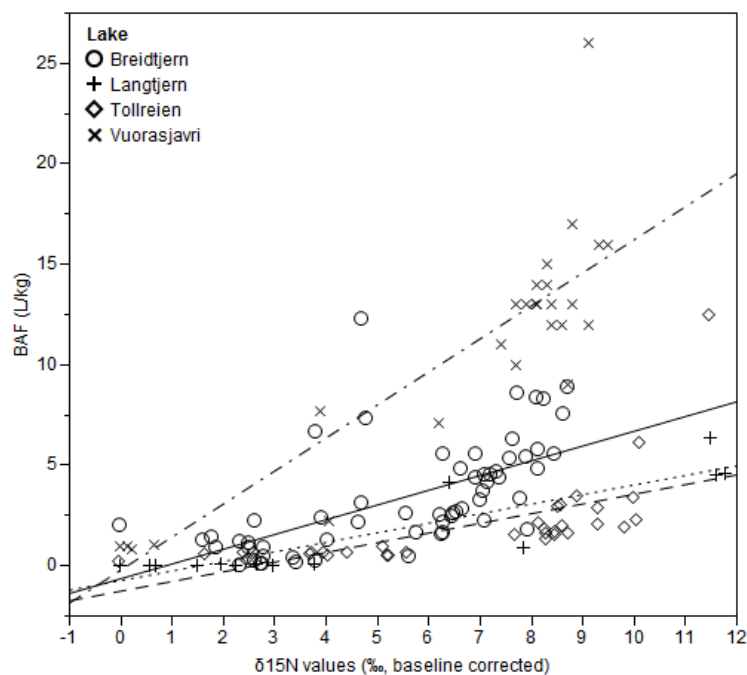


Figure 10 BAF (bioaccumulation factor, x-axis, L/kg) versus $\delta^{15}\text{N}$ values (y-axis, ‰, baseline corrected) for all groups of organisms (included fish) in the four study lakes. Shown are Breidtjern (circles, unbroken line, $y = -0.63 + 0.73x$, $r^2 = 0.38$, $p < 0.0001$), Langtjern (pluses, dotted line, $y = -0.73 + 0.48x$, $r^2 = 0.80$, $p < 0.0001$), Tollreien (diamonds, long dashed line $y = -1.25 + 0.48x$, $r^2 = 0.41$, $p < 0.0001$) and Vuorasjavri (crosses, dashed and dotted line, $y = -0.20 + 1.64x$, $r^2 = 0.75$, $p < 0.0001$). Figure copied from paper 6.

3.5 Biological food chain mechanisms

Biological food chain mechanisms were studied in paper 5 (changing fish Hg concentrations in relation to trophic position and summer growth dilution), paper 4 (habitat use, discussed under 3.4 *Aquatic in-lake processes*) and paper 6 (predation pressure and temperature influence on MeHg biomagnification).

3.5.1 *Changing fish trophic position*

The fish populations from both Breidtjern and Tollreien show a significant increase in Hg concentrations from 2010 to 2012 ($p < 0.0001$), with spring 2012 concentrations being the highest in the data material (Figure 5). Model predicted Hg concentrations in perch from 2012 (least square mean Breidtjern: 0.35 ± 0.03 mg/kg, Tollreien: 0.36 ± 0.03 mg/kg) are higher than any of the previous two years. Based on autumn data, the concentration increase between 2010 and 2012 is 46 % and 28 % in Breidtjern and Tollreien, respectively.

Interestingly, the concentration change in the two lakes is differently distributed between the three years. In Breidtjern the increase was 25 % from 2010 to 2011, and 17 % from 2011 to 2012. Similar numbers for Tollreien are 28 % from 2010 to 2011, while there was no change between 2011 and 2012 (0 %). As both lakes showed an increase from 2010 to 2012, with similar seasonal patterns (Figure 5), governing processes on a regional scale could be suggested to explain this. But, since our two lakes show different increases the three study years, processes on a smaller scale are also likely to occur. The mechanisms controlling these seasonal and yearly variations of fish Hg concentrations are however not clearly defined in the literature. Parameters suggested to influence temporal fish Hg dynamics are water chemistry (OM and pH: Akerblom et al., 2012, Rask et al., 2007 and Porvari, 1998), climate factors (temperature: Rask et al., 2010), dietary patterns and trophic position (Rask et al., 2010).

We investigated the abovementioned parameters, finding that only trophic position influenced the temporal Hg concentration variations in the perch populations. However, the higher fish Hg concentrations found in Tollreien compared to Breidtjern reflects the trends of different surface water chemistry (Table 3), with mean annual concentrations of TotHg, MeHg and %MeHg being significantly higher in Tollreien than Breidtjern for all three study years (t-tests, $p < 0.001$). Tollreien is also more humic than Breidtjern (Table 3). Together this highlights how the availability of aqueous Hg species to freshwater food chains (through catchment TOC transport: Grigal, 2002, and Hg speciation in water: Chasar et al., 2009 and de Wit et al., 2012) controls the general fish Hg

concentration levels. However, as there is no significant yearly variation seen in the water chemistry, this does not explain the short-term (yearly and seasonally) variation of fish Hg concentrations.

Mean seasonal $\delta^{13}\text{C}$ variations show no significant seasonal variations ($p > 0.05$, Figure 11), indicating that the fish populations collected in the present study do not change dietary patterns (within the same year). However, mean yearly $\delta^{13}\text{C}$ levels increase (i.e. less negative values) significantly (comparisons for each pair using Student's t , $p < 0.05$) in Breidtjern from 2010 (-28.7 ‰) to 2011 (-28.4 ‰) and from 2011 to 2012 (-28.1 ‰, Figure 11). This small change in mean $\delta^{13}\text{C}$ signal could indicate a shift in the carbon sources for the food chain, as previously documented for perch in Finland (Rask et al., 2010). This could again lead to changing fish Hg uptake due to habitat specific uptake of MeHg in primary consumers (paper 4).

Table 3 Mean (\pm one standard deviation) annual water Hg speciation and general water chemistry for Breidtjern and Tollreien. Data from 2010, 2011 and 2012 is based on $n = 3$, $n = 5$ and $n = 3$ sampling dates, respectively. Where no standard deviation is indicated, only 1 sample is considered. Table copied from paper 5.

Specification	Unit	Breidtjern			Tollreien		
Year		2010	2011	2012	2010	2011	2012
Hg speciation							
TotHg	ng/L	3.4	3.3 ± 0.8	3.0 ± 0.2	4.5	4.5 ± 1.5	4.6 ± 0.3
MeHg	ng/L	0.07	0.08 ± 0.01	0.07 ± 0.01	0.18	0.16 ± 0.03	0.18 ± 0.01
%MeHg	%	2.0	2.5 ± 0.4	2.4 ± 0.2	4.0	3.7 ± 0.6	4.0 ± 0.4
General water chemistry							
pH	-	4.4 ± 1.0	5.0 ± 0.3	5.0 ± 0.2	5.4 ± 0.1	5.4 ± 0.3	5.4 ± 0.1
Alkalinity	mmol/L	0.02	0.03 ± 0.01	0.03 ± 0.01	0.04 ± 0.01	0.05 ± 0.01	0.04 ± 0.01
TOC	mg/L	9.5 ± 2.1	9.6 ± 2.2	8.1 ± 1.8	13.1 ± 2.5	14.6 ± 4.3	16.0 ± 0.8
Tot-N	$\mu\text{g/L}$	295 ± 56	374 ± 19	358 ± 21	282 ± 32	341 ± 54	358 ± 11
Nitrate	$\mu\text{g/L}$	21 ± 12	40 ± 9	44 ± 8	5 ± 4	22 ± 15	13 ± 5
Tot-P	$\mu\text{g/L}$	3.7 ± 0.6	5.8 ± 1.5	6.3 ± 3.2	8.0 ± 1.0	9.2 ± 1.3	9.5 ± 3.5
Sulphate	mg/L	1.7 ± 0.1	1.7 ± 0.2	1.6 ± 0.1	1.3 ± 0.2	1.3 ± 0.2	1.0 ± 0.1

However, based on present data, changing $\delta^{13}\text{C}$ levels will only influence the fish Hg concentrations seen in Breidtjern, and cannot explain the yearly increase observed in Tollreien. In Tollreien, there is in fact a significant decrease (i.e. more negative value) seen from 2010 (-29.6 ‰) to 2011 (-30.2 ‰),

but an increase to 2012 (-28.1 ‰) in $\delta^{13}\text{C}$ levels (Figure 11). As is also discussed in detail in paper 5, $\delta^{13}\text{C}$ did not contribute to significantly increase the explanatory power of our fish Hg concentrations model, and was hence not included in the model. Based on this we conclude that a possible change in $\delta^{13}\text{C}$ signal is not responsible for the changing seasonal and year-to-year variation of perch Hg concentrations documented in the present study.

In Breidtjern there is a significant increase in $\delta^{15}\text{N}$ levels from 2011 to 2012 ($p < 0.01$, Figure 11), but no difference between 2010 and 2011. Since the fish Hg concentration increase in the lake is relatively large in both years (25 and 17 %, respectively), it is clear that $\delta^{15}\text{N}$ patterns cannot explain the increase alone. However, the fish caught in 2012 have a mean trophic position higher than the fish caught in 2010 and 2011, and could at least explain parts of the increasing fish Hg concentrations in Breidtjern.

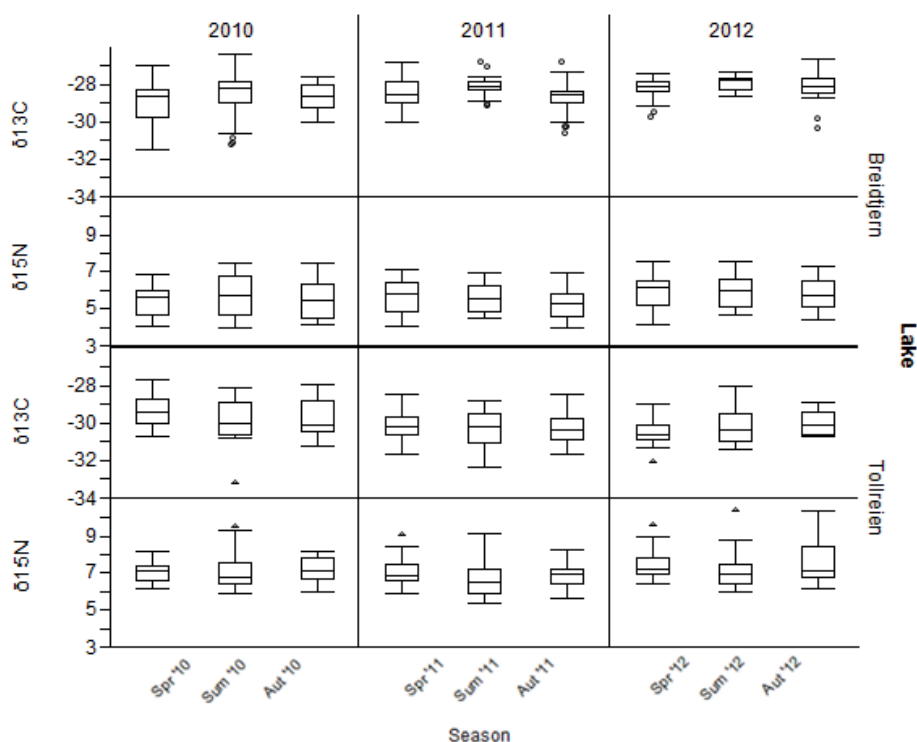


Figure 11 Box and whisker plots of seasonal $\delta^{13}\text{C}$ (top) and $\delta^{15}\text{N}$ (bottom, not adjusted) data in Breidtjern (above bold horizontal line) and Tollreien (below bold horizontal line). The horizontal line inside the box represent seasonal median value, the ends of the box represent 75th and 25th quantiles, and the end of the lines represent $\pm 1.5 \times$ interquartile range. Values outside this range are shown as circles (Breidtjern) and triangles (Tollreien). Figure copied from paper 5.

In Tollreien it is a significant ($p < 0.05$) decrease in $\delta^{15}\text{N}$ levels from 2010 to 2011 (Figure 11), while the fish Hg concentrations increase with 28 %. From 2011 to 2012, $\delta^{15}\text{N}$ levels increase significantly ($p < 0.0001$), while fish Hg concentrations show no increase (0 %). But, since it significantly increased the explanatory power, data of $\delta^{15}\text{N}$ were added to our fish Hg concentrations model. This implies, as for Breidtjern, that $\delta^{15}\text{N}$ can, if not alone, at least partly, explain the year-to-year increase in mean fish Hg concentration from 2010 to 2012 (Figure 5).

A point that could clarify the observed relationship between year-to-year increase in fish Hg concentrations and $\delta^{15}\text{N}$ levels is the bioaccumulation rates in the two fish populations (Figure 12). As is discussed in paper 5, the mean perch collected from Tollreien is larger than the mean perch collected from Breidtjern. However, Hg (i.e. MeHg) accumulates at a slower rate in Tollreien (TMS = 0.43, all data) compared to Breidtjern (TMS = 0.50). We hypothesise that this is related to the group of large predatory fish in Tollreien that feed on smaller fish. This will lead to shortened life history for the smaller fish due to stress, and they will not accumulate as much MeHg as fish without this top predator pressure (as for Breidtjern perch). Hence, the TMS will be less steep than what is present in Breidtjern where the smaller fish live longer.

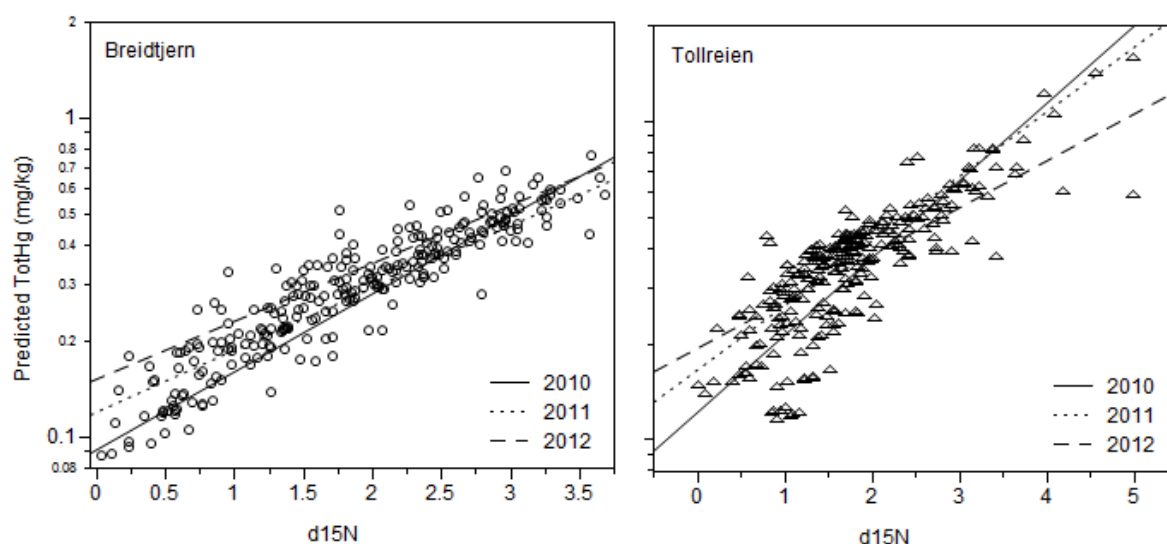


Figure 12 Log Hg concentrations (y-axis, mg/kg) versus $\delta^{15}\text{N}$ values (x-axis, ‰, adjusted) for all fish included in the present study. Shown are Breidtjern (left panel, circles) and Langtjern (right panel, diamonds). Trophic

magnification slopes for the years 2010 (unbroken lines), 2011 (dotted line) and 2012 (dashed line) is 0.56, 0.45 and 0.42 for Breidjtjern. Similar numbers for Tollreien are 0.56, 0.47 and 0.34. Figure copied from paper 5.

The TMS are decreasing in both our lakes from 2010 through 2011 to 2012 (Figure 12). In 2010 TMS values are 0.56 in both Breidjtjern and Tollreien, while TMS in 2011 and 2012 are 0.45 and 0.47, and 0.42 and 0.34 in the two lakes, respectively. This indicates that MeHg is accumulating slower in the fish populations every year, suggesting that biological mechanisms are responsible for the changing fish Hg concentrations. An explanation for the reduced TMS could be increased pressure on the fish population, for example from exterior factors we have been unable to access in the present data set, leading to shorter life histories and reduced MeHg accumulation in the fish populations. This reflects again the significant contribution from $\delta^{15}\text{N}$ levels on fish Hg concentrations, and hence explains the increasing fish Hg concentrations observed.

3.5.2 Variation of MeHg biomagnification

In paper 6 we studied biomagnification of MeHg through documentation of lake-specific TMS and BAF in four of our study lakes: Breidjtjern, Tollreien, Langtjern and Vuorasjavri. Due to the long list of variables suggested to affect MeHg bioaccumulation rates, TMS are shown to vary significantly both within individual food chains and across systems with different physicochemical characteristics (Lavoie et al., 2013, Kidd et al., 2012). Therefore, in paper 6, our main goal was to use the principles of TMS and BAF for MeHg to determine the factors significantly controlling MeHg biomagnification in boreal and subarctic lakes. We included a wide range of possible explanatory environmental parameters, including different food chain characteristics in addition to physical and chemical lake features. Our main hypothesis was that for lakes where physicochemical features and food chain carbon sources are similar, biomagnification of MeHg are significantly affected by the top predator.

We found that mean concentrations of MeHg in organisms (excluding fish) from Vuorasjavri (44 ± 59 ng/g) were significantly lower than the mean concentrations from Breidjtjern (157 ± 201 ng/g, Tukey-Kramer, $p < 0.01$). However, there were no significant difference between mean MeHg concentrations in Vuorasjavri, Langtjern (63 ± 94 ng/g) or Tollreien (90 ± 32 ng/g, Kruskal-Wallis

test, $p = 0.70$). This is somewhat surprising, as concentrations of aqueous Hg species (both TotHg and MeHg) in lakes from subarctic Norway are shown to be significantly lower than concentrations from boreal lakes (paper 2). It is also documented a positive correlation between MeHg concentrations in invertebrates and stream water concentrations of MeHg, TotHg and DOC (Chasar et al., 2009). A previous study also indicates how an acidic lake food chain (pH 5.2 – 5.6) accumulates more Hg than in a less acidic lake (pH 6.3 – 6.9), when concentrations of DOC (2.7 mg/L) are similar (Scheuhammer and Graham, 1999).

MeHg concentrations increased significantly with trophic position in Breidtjern, Tollreien, Langtjern and Vuorasjavri, as indicated by increasing $\delta^{15}\text{N}$ signatures ($r^2 = 0.43, 0.82, 0.91$ and 0.93 , respectively, all $p < 0.0001$, lower food chain and fish data included, Figure 13). For three of the lakes, ANCOVA revealed no significant differences in TMS ($p = 0.33$): Breidtjern (TMS \pm 95 % confidence interval: 0.34 ± 0.10), Tollreien (0.26 ± 0.04) and Vuorasjavri (0.31 ± 0.03). A previous review of TMS values for MeHg biomagnification from food chain studies shows a mean TMS of 0.24 ± 0.08 ($n = 106$) from freshwater when the complete food chain is considered (Lavoie et al., 2013). This is in approximate agreement with what we found when including fish in our calculations from Tollreien, Breidtjern and Vuorasjavri.

Other studies show that the variation can be even larger than what is documented in Lavoie et al. (2013), e.g. Clayden et al. (2013): 0.13 – 0.23, Clayden et al. (2014): 0.46, and we found in fact a higher TMS value in Langtjern (0.51 ± 0.09). Langtjern TMS was significantly higher than in the other three lakes (ANCOVA, $p < 0.001$). Since water chemistry and climate is similar (not significantly different) in Langtjern and the other boreal lakes (Breidtjern and Tollreien, Kruskal-Wallis test, $p > 0.05$), the explanation for the high TMS probably relates to biological factors (see discussion on biological influence on MeHg biomagnification further on).

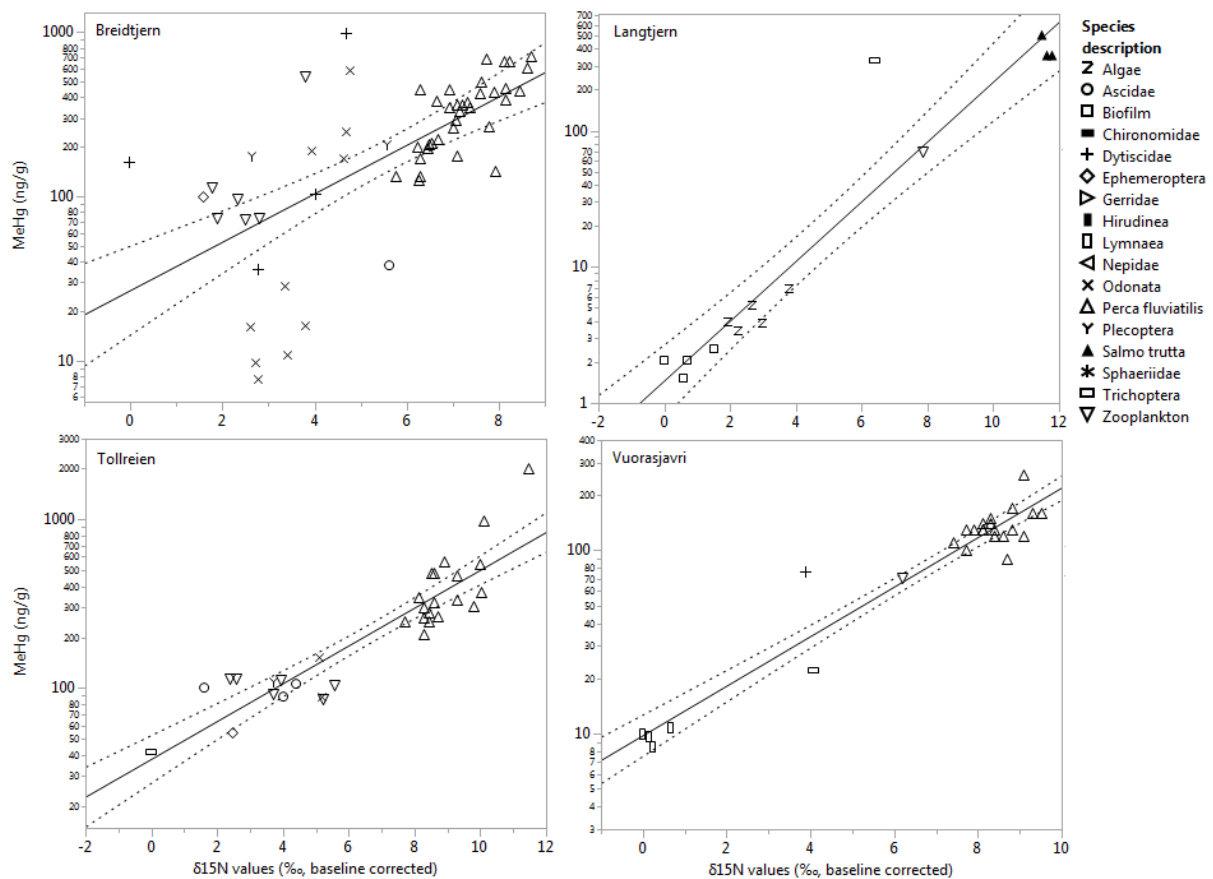


Figure 13 Log MeHg (Hg for fish) concentrations (x-axes, ng/g dw) versus $\delta^{15}\text{N}$ values (y-axes, ‰, baseline corrected) for all groups of organisms (included fish) in the four study lakes. Shown are Breidtjern (top left, $y = 3.32 + 0.34x$, $r^2 = 0.44$, $p < 0.0001$), Langtjern (top right, $y = 0.39 + 0.51x$, $r^2 = 0.91$, $p < 0.0001$), Tollreien (bottom left, $y = 3.64 + 0.26x$, $r^2 = 0.82$, $p < 0.0001$) and Vuorasjavri (bottom right, $y = 2.29 + 0.31x$, $r^2 = 0.93$, $p < 0.0001$). TMS are shown with confidence curves (broken lines). Figure copied from paper 6.

3.5.3 Temperature dependent MeHg biomagnification

In the Lavoie et al. (2013) review, the authors found that MeHg biomagnification in aquatic food chains on a global scale is positively related to latitude (MeHg TMS versus latitude: $r^2 = 0.10$, $p < 0.001$). TMS values are shown to be higher in polar (0.28 ± 0.09) compared to temperate regions (0.24 ± 0.07). In our study, we did not find support for this to be true, as TMS values for the subarctic lake Vuorasjavri (with and without fish: 0.31 ± 0.03 and 0.35 ± 0.17) did not significantly differ from TMS values for the two boreal lakes Tollreien (0.26 ± 0.04 and 0.12 ± 0.10) and Breidtjern (0.34 ± 0.10 and 0.24 ± 0.20). The mechanisms thought to be responsible for a possible south-north gradient mainly

relate to temperature (discussed in Lavoie et al., 2013) and include growth dilution (increased in warmer regions, Simoneau et al., 2005, trophic transfer efficiency, which is reduced in warmer regions, and excretion rates of MeHg, which is reduced in colder regions, Trudel and Rasmussen, 1997). Mean annual temperature is lower in the subarctic region (-3.0 °C for the area of Vuorasjavri) compared to the boreal areas where the three other lakes are located (1.3 – 5.8 °C, Table 1, also discussed in paper 2), but still no significant TMS value difference was observed.

3.5.4 Biological influence on MeHg biomagnification

To assess the impact of top predators on MeHg biomagnification, we calculated TMS when fish samples were removed from the data material (Figure 14). TMS values in Breidtjern and Tollreien decreased from 0.34 ± 0.10 to 0.24 ± 0.20 (ANCOVA, $p < 0.001$) and from 0.26 ± 0.04 to 0.12 ± 0.10 (ANCOVA, $p = 0.06$), respectively. This is contrary to what Lavoie et al. (2013) reported, where TMS values were found to increase to 0.31 ± 0.10 ($n = 3$), similar to what we observed in Langtjern (from 0.51 ± 0.09 to 0.62 ± 0.21 , ANCOVA, $p < 0.001$) and Vuorasjavri (from 0.31 ± 0.03 to 0.35 ± 0.17 , ANCOVA, $p < 0.01$). In the Lavoie et al. (2013) study, differences in TMS values were hypothesised to be related to different energy requirements of chosen organisms. We suggest that additional factors may deserve consideration. When fish was excluded from the TMS calculations, no significant differences was evident in the data material (ANCOVA, $p = 0.08$).

When only fish was included in the calculations, values were 0.46 ± 0.16 ($p < 0.0001$), 0.46 ± 0.18 ($p < 0.0001$) and 0.18 ± 0.16 ($p < 0.05$) for Breidtjern, Tollreien and Vuorasjavri, respectively (data not shown). In the three lakes, no significant difference in TMS was evident when only fish was included (ANCOVA, $p = 0.14$). TMS for fish only were not calculated for Langtjern due to the small number of samples available ($n = 3$).

Breidtjern and Tollreien both have abundant perch populations, with no (perch only fish species) predator pressure on the perch (paper 5). Perch undergo life history dependent dietary shifts (Collette et al., 1977), but in Breidtjern and Tollreien very few fish reach the sufficient size to be piscivorous (paper 5). Hence, we assume that all fish in the two lakes prey on primary and secondary invertebrates. Fish predation may principally induce various shifts in prey population dynamics

(Gilljam et al., 2011, Wooster and Sih, 1995), depending on predation intensity, prey behaviour, life history and reproductive habits (Henderson et al., 2012). Increased stress (from predator pressure) may cause prolonged development and could in principle favour MeHg accumulation. But high predator pressure on the primary and secondary consumers could lead to shortened life history, as well. Following this last argument, low MeHg concentrations relative to trophic level seems likely, as the prey species do not live long enough to accumulate substantial amounts of MeHg. Hence, the process counteracts optimal somatic growth in the prey community and lead to low MeHg magnification rates (i.e. low TMS) for the food chain. This explains the patterns of significantly lower TMS observed in Breidtjern (0.34 ± 0.10) and Tollreien (0.26 ± 0.04) compared to Langtjern (0.51 ± 0.09) when all data is included (Figure 13).

The high TMS for Langtjern appears to be a function of lower MeHg concentrations in low $\delta^{15}\text{N}$ species rather than higher MeHg concentrations in high $\delta^{15}\text{N}$ consumers (Figure 13), which, following the arguments above, indicates low predation pressure. This is consistent with the low number of fish caught in this lake. The weak top down signal from fish in Langtjern probably increases the abundance of prey items compared to the other boreal lakes, as discussed by Gliwicz (2002).

Le Jeune et al. (2012) found that in lakes inhabited by fish, the *Chaoborus* larvae were unable to biomagnify MeHg, while in fishless lakes, *Chaoborus* larvae biomagnified MeHg. The authors states that growth dilution, amount and type of prey items or trophic position could not explain the different MeHg biomagnification patterns. In stead Le Jeune et al. (2012) points to other possible biological explanations, specifically that the biomagnification capacities of *Chaoborus* larvae are affected by diel vertical migration. The migration is affected by the fish patterns in the lakes (i.e. whether fish is present or not), supporting the findings of our paper 6.

Breidtjern and Tollreien show the same pattern of higher TMS when fish is included in the MeHg concentrations versus $\delta^{15}\text{N}$ levels plot (Figure 13 and 14), contrary to the patterns reported by Lavoie et al. (2013). Based on this we would like to draw attention to other possible mechanisms influencing MeHg biomagnification. With the high predator pressure from perch in these two lakes (see paper 5 and 6 for details), we suggest that primary and secondary consumers do not grow old to

accumulate substantial amounts of MeHg. This will lead to low biomagnification rates of MeHg (i.e. low TMS) through the lower parts of the food chain, compared to when the full food chain is considered (higher TMS). This fits well with previous findings showing that biomass, individual body size and population density are top-down controlled (Gliwicz, 2002).

Langtjern and Vuorasjavri show an opposite pattern. In Langtjern, the TMS increase from 0.51 ± 0.09 to 0.62 ± 0.21 when fish is excluded from the calculations (Figure 13 and 14). We hypothesise that this is related to a weak predator control in the Langtjern food chain, due to the sparse and artificially stocked population of brown trout. Lower trophic levels in Langtjern remain relatively undisturbed from predatory fish and may have accordingly a prolonged life history compared to the primary and secondary consumers in Breidtjern and Tollreien. Following this argument, MeHg in Langtjern will show higher biomagnification rates in invertebrates (0.62 ± 0.21) compared to when the full food chain is considered (0.52 ± 0.09), opposite of what is observed in Breidtjern and Tollreien.

In Vuorasjavri we see a similar pattern to that of Langtjern (Figure 13 and 14), and a low TMS when only fish is considered (0.18 ± 0.16). This is again opposite of the patterns in Breidtjern and Tollreien, even though predator pressure is similar to Tollreien (see paper 5 and 6 for details). We believe this to be an indication of the Vuorasjavri perch to be more stressed by other predators than what we see in the two boreal lakes. Since TMS is higher in the subarctic lake when the fish is excluded (0.35 ± 0.17), we suggest that the lower trophic levels in Vuorasjavri are less affected by fish predation, probably because the perch population is under a considerable top down pressure from large piscivorous perch and pike (and possibly other species as well). These conditions will then lead to a TMS pattern similar to that of Langtjern.

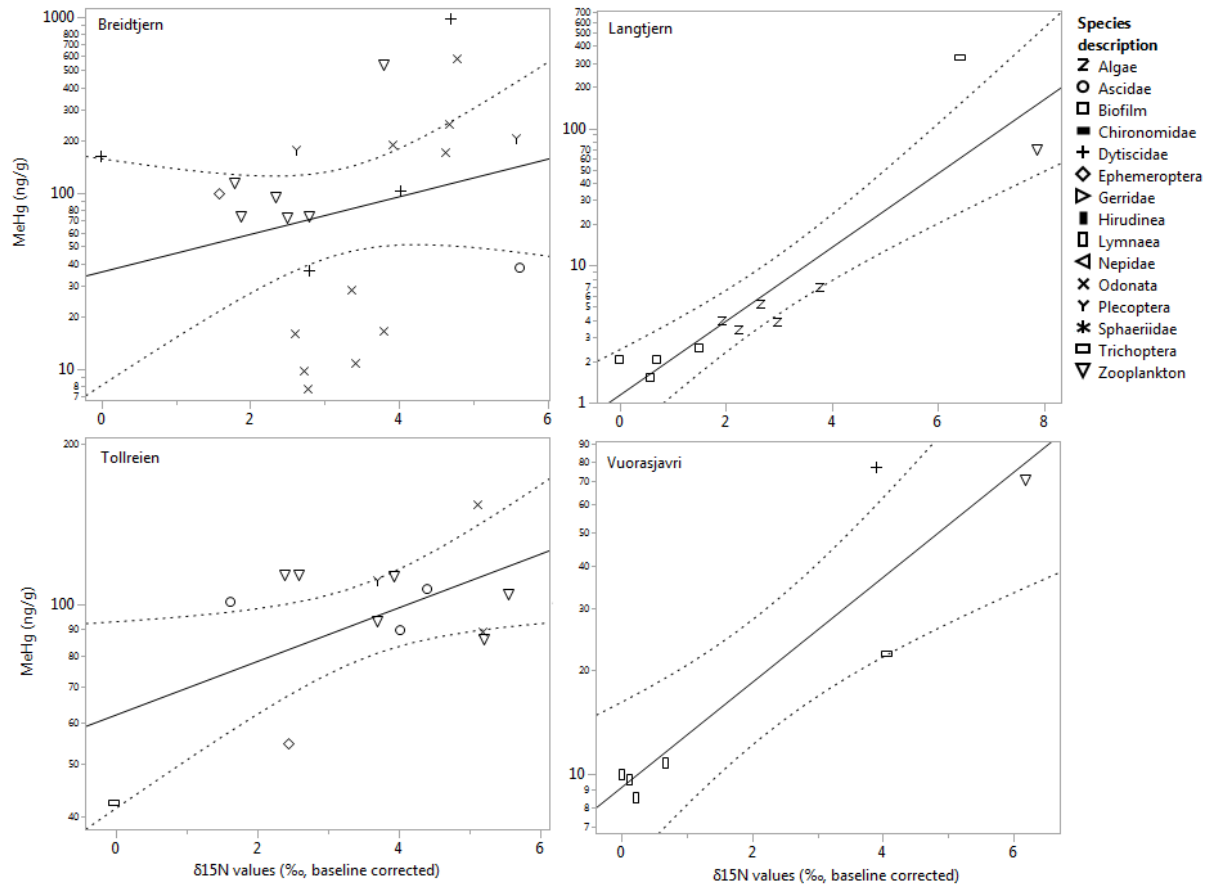


Figure 14 Log MeHg concentrations (y-axes, ng/g dw) versus $\delta^{15}\text{N}$ values (x-axes, ‰, baseline corrected) for all groups of organisms (excluded fish) in the four study lakes. Shown are Breidtjern (top left, $y = 3.62 + 0.24x$, $r^2 = 0.06$, $p = 0.23$), Langtjern (top right, $y = 0.14 + 0.62x$, $r^2 = 0.83$, $p < 0.0001$), Tollreien (bottom left, $y = 4.13 + 0.12x$, $r^2 = 0.33$, $p < 0.05$) and Vuorasjavri (bottom right, $y = 2.22 + 0.35x$, $r^2 = 0.83$, $p < 0.01$). TMS are shown with confidence curves (broken lines) for the slopes. Figure copied from paper 6.

4 Conclusions

In pristine areas of Norway, where no local emission of Hg exists, concentrations of Hg in fish are not only high, but also increasing. The explanatory factors and their processes that are directly and indirectly governing Hg concentrations in freshwater fish are abundant and diverse. This thesis shows that significant processes are occurring in the catchment, in the lake itself and also in the food chain (see *Illustration*). Processes of particular importance highlighted in the present thesis are:

- *Catchment Hg cycling:*

OM as the transport vector from stored Hg in soil to the lake; OM as methylation substrate in the catchment; and catchment nutrient mediated methylation.

- *Aquatic in-lake processes:*

OM as methylation substrate and nutrient mediated methylation in the aquatic phase or the sediments; PD of aquatic MeHg; littoral sediment methylation; and chlorophyll versus DOM associated MeHg transport from the aquatic phase to the food chain.

- *Biological food chain processes:*

Changing fish trophic position; predator related MeHg biomagnification variation.

Based on the observations documented in the present thesis, our four main objectives have given the following conclusions:

1. *Are concentrations of Hg in freshwater fish in Norway still increasing (after 2008), and what are the potential drivers behind such a possible increase?*

In both study lakes Tollreien and Breidtjern fish Hg concentrations are higher in all study years (2008, 2010, 2011 and 2012) documented after the 1990s. So, although concentrations are varying significantly from year-to-year, this suggests that the concentrations of Hg in perch in these lakes are still increasing. In both Tollreien and Breidtjern perch Hg concentrations show a significantly increase also from 2010 to 2012. Although the lakes are both located in south eastern Norway, the increases of Hg concentrations are differently distributed between the study years 2010, 2011 and

2012. Together this suggests that both regional and local processes should be considered as primary drivers for the increasing concentrations. However, with the present data set we can only conclude that biological factors related to change in fish trophic position and changing MeHg bioaccumulation are significant.

2. *What are the key variables explaining the spatial concentration levels of Hg and MeHg, in addition to methylation potential, in Norwegian surface waters?*

TOC was the variable most strongly correlated with TotHg and MeHg concentrations in our spatial data set. Statistical modelling revealed that, after TOC, the most significant explanatory variables were N availability, base cation status, and lake and catchment size. A key process driving TotHg concentrations is DOM as a transport vector, while the role of DOM for MeHg and %MeHg is likely related to a combination of transport and DOM as a substrate for methylation. The observed negative correlations between MeHg and catchment and lake size are consistent with in-lake and in-stream de-methylation processes. Statistical modelling suggests that N availability exerts a positive contribution on concentrations of MeHg and %MeHg.

3. *What are the main biological and physicochemical lake features affecting the bioaccumulation and biomagnification of MeHg through boreal and subarctic lake food chains?*

Data from four lakes in boreal and subarctic Norway suggests that inter-lake differences in pressure from predatory fish may significantly affect bioaccumulation and biomagnification of MeHg through the food chains. Low predator pressure lead to prolonged life history for primary and secondary consumers, producing higher TMS as a result of increased MeHg biomagnification. In the subarctic lake we also show how aquatic MeHg transfer to zooplankton is most likely chlorophyll associated, rather than the detritus associated transport in the boreal lakes. As %MeHg is documented to be lower in profundal relative to littoral sediments and MeHg concentrations in primary consumers follow this pattern, we also suggest that shallow lake sediments are important for MeHg transfer to the aquatic food chains in boreal humic lakes.

4. *How will photochemical degradation affect concentration levels of MeHg in Norwegian surface waters today and in terms of different future DOC concentration scenarios?*

For the study at Langtjern, losses of MeHg through PD equalled almost 1/3 of total annual inputs. This clearly highlights the importance of PD in the MeHg budget of boreal lakes. Future scenario calculations showed how changes in catchment DOC export to freshwaters may lead to higher aqueous MeHg concentrations due to increased DOC-associated MeHg inputs paired with strong decreases in losses of MeHg through PD due to increased light attenuation. The data also suggests that future climate driven reduced ICD will not offset the negative effects of increased DOC loading on PD losses.

5 Future work

With atmospheric deposition of Hg showing decreasing or unchanged patterns in the period where concentrations in fish are increasing, the historically accumulated Hg in the catchment soil are likely to affect the adjoining lake for decades and centuries. Policy on Hg in the environment must acknowledge the large Hg stores in the environment, accumulated from centuries of anthropogenic and natural Hg emissions, that may be mobilized and contaminate aquatic food chains. Climate change, together with other factors and drivers, may enhance both mobility of recently and historically deposited Hg and production of MeHg, as highlighted through the processes studied in the present thesis.

However, to be able to increase our understanding of what controls the Hg concentrations in fish in northern ecosystems, research needs to be focused on future combined effects of climate and pollution (i.e. atmospheric deposition), as well as transport and accumulation processes of MeHg. Climate factors such as precipitation and temperature can enhance the transport of Hg and production of MeHg in the studied ecosystems, which will influence aqueous MeHg concentrations and bioavailability. A detailed study of these factors under different pollution scenarios, e.g. atmospheric deposition of Hg, S and N, is critical for improving predictions of bioaccumulation of Hg in these food chains.

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