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▶ To cite this version:

Stephen Kohler, Lars-Eric Heimbürger-Boavida, Philipp Assmy, Oliver Müller, Stefan Thiele, et al.. Biotic transformation of methylmercury at the onset of the Arctic spring bloom. Progress in Oceanography, In press, 10.1016/j.pocean.2024.103224. hal-04470915

HAL Id: hal-04470915 https://hal.science/hal-04470915v1

Submitted on 21 Feb 2024

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Biotic transformation of methylmercury at the onset of the Arctic spring bloom

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16 Graphical Abstract



19 Keywords

20 Barents Sea, mercury, seasonality, Arctic Ocean, spring bloom

21 Abstract

22 Despite the lack of local anthropogenic mercury sources, methylated mercury (MeHg) 23 concentrations in Arctic biota are higher than in biota from lower latitudes. The main entry route 24 occurs during the bioconcentration of seawater monomethylmercury (MMHg) into phytoplankton. 25 Despite the known seasonal changes in biological activity in the region, little is known about the 26 seasonal cycling of total mercury (THg) and MeHg in the Arctic Ocean. Here, we report the 27 concentrations of THg and MeHg in seawater sampled from the northwestern Barents Sea water 28 column during late winter and spring. In the upper 500 m, the THg concentrations are significantly higher in spring $(0.64 \pm 0.09 \text{ pmol } \text{L}^{-1})$ compared to late winter $(0.53 \pm 0.07 \text{ pmol } \text{L}^{-1})$, driven by 29 30 seasonal inputs to surface waters from atmospheric deposition and the dynamics of changing sea 31 ice conditions. Contrastingly, the MeHg concentrations in spring were significantly lower (41 \pm 39 fmol L⁻¹) compared to late winter (85 ± 42 fmol L⁻¹). We suggest that most MeHg is biotically 32 demethylated by both phytoplankton and bacteria, with additional losses from photodemethylation 33 34 and evasion. Our observations highlight the importance of demethylation during potential uptake 35 of methylmercury coinciding with the Arctic spring bloom. Lastly, we use our new data together 36 with previously published seasonal data in the region to construct a simplified seasonal mercury 37 cycle in an Arctic marginal ice zone.

38 1 Introduction

39

40 The bioaccumulation of mercury (Hg) in the Arctic Ocean continues to remain an issue of concern 41 for the marine ecosystem and indigenous peoples (Basu et al., 2022). Specifically, the 42 bioconcentration of toxic monomethylmercury (MMHg) into phytoplankton and its subsequent 43 bioaccumulation in the marine food web represents a risk to higher trophic level organisms (Dietz 44 et al., 2022). The development and peak of primary productivity in the Arctic Ocean is restricted 45 to the spring and summer (April to August), when light is available. In the spring, the 46 phytoplankton bloom develops in the marginal ice zone (Strass and Nöthig, 1996; Reigstad et al., 47 2002). Several studies have highlighted phytoplankton blooms below fully consolidated sea ice as 48 a result of the thinning and more dynamic icescape (Assmy et al., 2017, Ardyna et al., 2020).

49 Concurrently, the Arctic Ocean receives seasonal inputs of mostly inorganic Hg (iHg) via atmospheric deposition, riverine input, erosion, snow, glacial and sea ice melt (Dastoor et al., 50 51 2022). However, seasonal cycling of Hg in the water column in the Arctic region is still poorly 52 understood (Kohler et al., 2022a). As the Arctic spring bloom provides a crucial transfer of energy 53 and potentially Hg into higher trophic levels, more knowledge on the amount of Hg available for bioconcentration prior and during the Arctic spring bloom is needed. Seasonal inputs from thawing 54 55 permafrost, glacial melt, and erosion are expected to deliver greater amounts of iHg to the 56 biologically productive Arctic Ocean shelf seas in the future (Chételat et al., 2022) with most Hg expected to remain in the marine environment (Araujo et al., 2022). Hg can be scavenged to shelf 57 58 sediments, advected with the halocline, cascaded into the deep basin (Dastoor et al., 2022) or 59 potentially transported into the Central Arctic Ocean via the transpolar drift (Charette et al., 2020). 60

61 Several major Hg inputs to the Arctic region have strong seasonal patterns. Atmospheric mercury 62 depletion events (AMDEs) (Schroeder et al., 1998) occur during polar sunrise in March and 63 frequently occur throughout April, depositing iHg to snow, sea ice, and surface waters. Although 64 a substantial fraction is photochemically reduced and reemitted back to the atmosphere as gaseous Hg⁰ (Durnford and Dastoor 2011, Araujo et al., 2022), high concentrations of total Hg (THg) are 65 66 seen in snow, sea ice and frost flowers (Dommergue et al., 2010, Chaulk et al., 2011). Sea ice, 67 melting snow and glacial runoff can also provide additional Hg to the coastal and open ocean 68 (Schartup et al., 2020, Kim et al., 2020a, Dastoor et al., 2022). Rivers also deliver large amounts 69 of iHg from the snowpack, surface soils, thawing permafrost, and erosion to the Arctic shelf seas 70 during a freshet in springtime (Sonke et al., 2018, Zolkos et al., 2020). While new evidence 71 suggests seasonal cycling of iHg in the upper water column during autumn and early winter 72 (Kohler et al., 2022a) with suspected surface losses to deeper waters and sediments with sinking 73 particles, water column measurements of Hg species from late winter are scarce (Wang et al., 74 2012).

75

Arctic surface waters are elevated in THg during the summer (Heimbürger et al., 2015) from
seasonal iHg inputs. Light reduces iHg to Hg⁰ and photolytically degrades MeHg (Sellers et al.,
1996). Gaseous Hg species can evade from the marginal ice zones (Yue et al., 2023) while ionic
Hg species can be transported downward with sinking particles (Tesán Onrubia et al., 2020). Ionic

80 Hg is also often complexed to dissolved organic matter (DOM) and reduced sulfur compounds, 81 which can affect Hg uptake and biogeochemical transformations between iHg and MeHg 82 (Ravichandran 2004). During and after the phytoplankton bloom, bacterial remineralization of sinking particulate organic matter fuels in situ MeHg production, leading to elevated subsurface 83 concentrations of MeHg in the marginal ice zones and the permanently ice-covered central Arctic 84 Ocean (Heimbürger et al., 2015). However, sea ice hinders both photodemethylation (Point et al., 85 2009) and the evasion of gaseous species (Hg⁰ and dimethylmercury (DMHg)) (DiMento et al., 86 2019). Gaseous Hg species potentially build up under and within the ice throughout winter 87 (Andersson et al., 2008, Schartup et al., 2020). Hg that is removed from surface waters via sinking 88 particles and delivered to bottom sediments may ultimately be methylated to MeHg (Kim et al., 89 90 2020b). Resuspension of fine-grained sediment and diffusion of porewater can deliver both iHg and MeHg back to subsurface Arctic waters (Kim et al., 2020b). While in situ incubations indicate 91 92 Arctic waters can be in net states of methylation or demethylation (Lehnherr et al., 2011), the advection of outflowing Arctic waters indicate that the Arctic Ocean is a net producer of MeHg 93 94 (Petrova et al., 2020, Jonsson et al., 2022a). Consequently, understanding the timing and 95 magnitude of both Hg methylation and demethylation is important. Due to a limited number of observations, the fate of marine MeHg throughout the Arctic winter and during the transition to 96 97 the spring bloom is unknown.

98

99 We hypothesized the potential seasonality of pelagic Hg species in the northern Barents Sea in the 100 late winter to spring transition due to seasonal Hg inputs and biological development. We 101 measured unfiltered THg (the sum of all mercury species) and unfiltered MeHg (the sum of MMHg 102 and DMHg) in seawater, under ice water, and sea ice on two seasonal cruises in late winter (March) 103 and spring (end of April-May). We report Hg concentrations from seven stations along a south-104 north transect and explore how hydrography, biological parameters, and seasonal processes affect 105 our observations using literature and statistical relationships. We combine our Hg and environmental data with that of two other seasonal cruises in the region (Kohler et al., 2022a) to 106 107 propose a simple seasonal model of the pelagic Hg cycle in surface waters of an Arctic marginal 108 ice zone.

110 **2 Materials & Methods**

111 **2.1 Oceanographic Conditions**

112 The northern Barents Sea consists of water masses with both Atlantic Water (AW) and Polar Water 113 (PW) origins (Sundfjord et al., 2020). AW, characteristic of warmer water with higher salinity, 114 flows into the northern Barents Sea from the south through the Barents Sea Opening and through 115 troughs in the north (Lundesgaard et al., 2022). This AW sinks below colder and fresher PW which 116 arrives from the central Arctic Ocean and the Kara Sea in the east (Loeng 1991). The boundary 117 between these two water masses is often referred to as the polar front, and generally delineates the 118 extent of sea ice in the region. Surface PW with colder temperatures entering the northern Barents 119 Sea leads to sea ice formation and can advect additional sea ice into the region. In the summer, sea 120 ice melt and surface heating form a stratified meltwater layer in the upper water column. By 121 autumn, the beginnings of sea ice formation, cooling, and turbulence lead to water column mixing. 122 This overturning of the water column continues throughout the polar night and into spring. Dense 123 water formation on the Barents Sea shelf enters the Arctic Basin through deep water plumes along 124 the continental shelf (Rudels et al., 2000).

125 On the western coast of Svalbard, the West Spitsbergen Current brings AW into the Eurasian 126 Basin. This current continues parallel along the Barents Sea shelf break and past Franz Josef Land 127 in the east. This AW shows a distinct seasonality, with warmer temperatures in autumn and colder 128 temperatures in early spring (Pérez-Hernández et al., 2019). As AW enters the Arctic Basin, it 129 begins to cool and sinks below PW, forming an AW core at around 300 m depth in the water 130 column north of Svalbard. As mixing occurs between AW and PW, continued temperature losses 131 lead to the formation of modified Atlantic Water (mAW) below the pycnocline and AW core. As 132 this water continues to cool below 0°C and sinks, typically below 1000 m, it forms Eurasian Basin 133 Deep Water (EBDW). In addition, extensions of the West Spitsbergen Current flowing onto the 134 northern Barents Sea shelf (Lundesgaard et al., 2022) can form warm Polar Water (wPW). This wPW is not as salty as AW yet warmer than PW, and can be found in deeper depths of the northern 135 136 Barents Sea shelf or at the surface during summer stratification.





Figure 1: Map of sampling locations (stations P1-P7) in the northern Barents Sea region. A solid
black line indicates the approximate location of the Polar Front, while the dashed colored arrows
show a simplified water mass scheme and current direction, with Atlantic Water (AW) in red and
Polar Water (PW) in blue. The color bar indicates water depth. Map created with Ocean Data View
(Schlitzer 2021).

145 All samples were collected on a south-north transect in the northern Barents Sea and Nansen Basin 146 (76 – 83°N) on two cruises, Q1 from March 2-24, 2021 (hereafter referred to as late winter), and 147 Q2 from April 27 - May 20, 2021 (hereafter referred to as spring), on R/V Kronprins Haakon as 148 part of The Nansen Legacy project (Fig. 1). Seawater samples for Hg were taken at seven stations 149 at 8 depths down to 500 m depth using Teflon-coated GOFLO bottles attached sequentially to a 150 600 m nonmetallic wire. At depths deeper than 500 m, Hg samples were collected with 10 L Niskin 151 bottles mounted to a rosette. Upon retrieval, GOFLO bottles were transported immediately to a 152 custom-made clean lab onboard. THg samples were collected unfiltered into pre-cleaned (Cutter 153 et al., 2017) 40 mL borosilicate glass bottles with PTFE-lined caps with no headspace and kept 154 unpreserved. MeHg samples were collected unfiltered into brand new 125 or 250 mL PET bottles 155 and were acidified under a Class 100 clean air laminar flow hood to 0.4% v/v with double-distilled 156 acidification, dimethylmercury HCl. Upon (DMHg) converts quantitatively to

157 monomethylmercury (MMHg) (Black et al., 2009); as a result, the sum of MMHg and DMHg is 158 measured as total methylated mercury (MeHg). All Hg samples were double-bagged and stored in 159 the dark at 4°C until analysis. Field blanks for THg were collected by bringing laboratory MilliQ 160 onboard and subsampling at the time of field sampling into precleaned borosilicate bottles and 161 treated identically to seawater samples. However, as we could not guarantee trace metal clean 162 Niskin bottles for samples collected below 500 m, these data are reported (SI Fig. 1) but not 163 included in subsequent statistical processing when noted.

164

165 Under ice water (UIW) was collected using a handheld 10 L Niskin bottle lowered into a hole cut 166 into the sea ice at stations P4, P6, and P7 on both cruises. UIW was collected directly from the 167 Niskin similar to the other seawater samples. Sea ice was collected at two stations, P4 and P7, 168 using a 9 cm diameter ice corer (Kovacs Enterprises, Inc., USA), bagged whole and taken directly 169 onboard into a -20°C cold lab. The bottom 10 cm of the ice core was cut using a handheld carbon 170 steel saw. Approximately 3 mm of ice was shaved off from all sides of the ice section with a 171 ceramic knife to remove any potential contamination before being double bagged and melted at 172 room temperature. One frost flower sample was collected in late winter using plastic equipment 173 and melted at 4°C. After melting, all samples were decanted into precleaned 40 mL borosilicate 174 glass bottles and 125 or 250 mL PET bottles and stored in the dark at 4°C until analysis.

175 **2.3 Mercury Analysis**

176 The concentration of THg in the samples was measured by cold vapor atomic fluorescence 177 spectroscopy (CVAFS) according to USEPA Method 1631 at the Mediterranean Institute of 178 Oceanography (MIO) using a custom-made single gold trap setup described elsewhere 179 (Heimbürger et al., 2015, Kohler et al., 2022a), within one year after sampling. As samples were 180 unfiltered to reduce contamination risk from filtration equipment, THg concentration includes both 181 particulate and dissolved species of Hg. The THg concentration in all samples was calculated from 182 a 6-point calibration curve traceable to NIST 3133 (National Institute of Standards and 183 Technology) standard reference material in previously purged, Hg-free seawater. Standard 184 bracketing of samples with a NIST 3133 matrix spike were used to correct for eventual instrument 185 signal drift and ensure ongoing precision and recovery. Aliquots of certified reference materials

ORMS-5 (Research Council of Canada) and ERM-CA400 (European Reference Materials) were spiked into previously purged Hg-free seawater matrix with recoveries of $93 \pm 2\%$ and $125 \pm 4\%$, respectively. Selected seawater samples run in triplicate from both cruises had an average standard deviation of 0.05 pmol L⁻¹. Field blanks of MilliQ water were measured for THg for both late winter (0.31 ± 0.02 pmol L⁻¹, n = 5) and spring (0.23 ± 0.02 pmol L⁻¹, n = 3) cruises. As MilliQ water can potentially contain traces of Hg, we concluded that there was no significant contamination affecting our THg measurements.

193 The MeHg concentrations in seawater samples was measured using the isotope dilution (ID) technique via gas chromatography (GC) – sector field ICP-MS (GC-SF-ICP-MS) at MIO within 194 one year after sampling (Heimbürger et al., 2015). Briefly, approximately 125 mL of seawater 195 sample was spiked with enriched isotopes of i¹⁹⁹Hg and MM²⁰¹Hg (ISC Science) and equilibrated 196 197 for 15 minutes. Samples were then adjusted to pH 3.9 using a sodium acetate buffer solution 198 (ULTREX® II Ultrapure Reagent, J.T. Baker, USA) and NH₃ (ULTREX® II Ultrapure Reagent, 199 J.T. Baker, USA). A fresh aliquot of 1% sodium tetrapropylborate solution (Merseburger Spezialchemikalien) and 150 µL of isooctane (Sigma Aldrich) were added to derivatize and 200 201 preconcentrate Hg species. Samples were sealed and shaken for 15 minutes before the organic 202 extract was transferred to GC vials for analysis on a coupled GC (THERMO GC 1300 with GC220 203 transfer module) SF-ICP-MS (Thermo Element XR) system. Ultra-trace clean techniques and coupling to a high resolution ICP-MS allows a detection limit of 2 fmol L⁻¹ and a limit of 204 quantitation of 7 fmol L^{-1} (LOQ = 3.3*LOD). Selected seawater samples run in duplicate from 205 both cruises had an average standard deviation of 7 fmol L⁻¹. The MM²⁰¹Hg spike concentration 206 207 used to calculate the sample MeHg concentrations had an accuracy of $94.4 \pm 8.4\%$. The accuracy 208 of our spike was determined using reverse isotope dilution using the Brooks Rand MMHgCl 209 standard, traceable to NIST1641E.

210 2.4 Additional Data and Parameters

Water temperature and salinity were recorded on a Seabird SBE 9 CTD unit attached to a rosette. Quality controlled temperature, salinity, sea ice concentration, nutrients, chlorophyll *a*, particulate organic carbon, nutrients, and phytoplankton abundance data for all seasonal cruises were retrieved from the Norwegian Marine Data Centre and the Norwegian Polar Institute (Gerland 2022, 215 Ludvigsen 2022, Reigstad 2022, Søreide 2022, Steer and Divine 2022, Marquardt 2022b, 2022c, 216 2022d, 2022e, Müller 2023a, 2023b, 2023c, 2023d, Chierici et al. 2021a, 2021b, Jones et al., 217 2022a, 2022b, Vader 2022, Assmy 2022a, 2022b). Part of the late winter MeHg data (station 218 average, upper 400 m only) presented in this analysis was previously published in the bottom 219 image in Fig. 6 in Dietz et al., 2022. Flow cytometry, bacterial community analysis, and dissolved 220 organic matter analysis details are presented in the Supplementary Information. Latitude and 221 longitude coordinates for CTD stations differ slightly from GOFLO casts due to ship drift. 222 Conservative temperature (Θ), absolute salinity (S_A), and potential density anomaly (σ_{θ}) were 223 calculated in Ocean Data View. Water mass definitions were assigned based on the most recent 224 report from the Nansen Legacy project for the northern Barents Sea (Sundfjord et al., 2020).

225

Statistical processing was performed for data above 500 m depth in Minitab software. Spearman 226 227 rank correlation coefficients were calculated using a 95% CI with significant p-value < 0.05. Data 228 selected for Student's t-test were log transformed for non-normal data distributions prior to 229 evaluation using a 95% CI with significant p-value < 0.05. Log-log regression was performed on 230 selected environmental variables with significance evaluated by F-test with a 95% CI and 231 significant p-value < 0.05. Selected environmental parameters were chosen for Principal Component Analysis, where data is standardized prior to dimension reduction and plotted 232 233 orthogonally as loadings plots for the first two principal components.

235 **3 Results & Discussion**

236 **3.1 Hydrography & sea ice**

In general, all stations in the northern Barents Sea showed both similarities and differences in 237 238 water mass distributions (SI Figs. 2 & 3) in both late winter and spring, with changes influenced 239 by sea ice coverage or temperature shifts (Jones et al., 2023, Van Engeland et al., 2023). In the late 240 winter and spring sampling periods, station P1 south of the polar front had a mAW signature, with S_A greater than 35.06 g kg⁻¹ and $\Theta > 0^{\circ}$ C. Stations P2, P3, and P4, all north of the polar front, had 241 PW ($\Theta < 0^{\circ}$ C, $\sigma_{\theta} < 27.97$ kg m⁻³) in surface waters and wPW ($\Theta > 0^{\circ}$ C, S_A < 35.06 g kg⁻¹) in deeper 242 243 waters extending to the bottom. A deepening of PW in spring at stations P2-P5 could suggest 244 rapidly changing sea ice conditions (SI Fig. 4), with small increases in salinity indicating refreezing and brine rejection. Station P5 had low temperatures, with PW throughout the entire 245 246 water column in both seasons. Stations P6 and P7 intersect the West Spitsbergen Current, with PW 247 and wPW in surface waters extending until approximately 200 m depth, where both mAW and AW ($\Theta > 2^{\circ}$ C, S_A > 35.06 g kg⁻¹) extends down to about 500 m. In waters below 500 m, mAW 248 persists until about 800 m depth and EBDW ($\Theta < 0^{\circ}$ C, S_A > 35.06 g kg⁻¹) fills the Nansen Basin. 249 250 At the time of sampling in late winter, first-year sea ice covered all stations except for P1 251 (Marquardt et al., 2023a). Some sea ice changes were observed during sampling (SI Fig. 4), but in 252 general most stations were covered by close drift ice. Minor changes in ice coverage occurred 253 before sampling for shelf stations, while P6 and P7 showed significant variations in ice coverage. 254 However, by the end of the late winter campaign, P6 and P7 were also covered by close drift ice. 255 Shelf stations P2-P5 showed dynamic ice conditions in between the late winter and spring cruises, 256 while P6 and P7 were covered by slightly older sea ice in spring (Marquardt et al., 2023a).

257 **3.2** THg and MeHg distributions in the water column during late winter and spring

THg concentrations in late winter (Fig. 2, top left) in the northern Barents Sea and Nansen Basin were on average 0.56 ± 0.17 pmol L⁻¹ (Table 1). In spring (Fig. 2, top right), THg concentrations in the northern Barents Sea and Nansen Basin were on average 0.69 ± 0.24 pmol L⁻¹ (n = 67, ± 1SD, range: 0.46 - 2.37 pmol L⁻¹). In the upper 500 m, our average THg concentrations on the spring cruise (0.64 ± 0.09 pmol L⁻¹, n = 56) were significantly higher (log transformed, Student's t-test, p < 0.05) compared to late winter (0.53 ± 0.07 pmol L⁻¹, n = 51). On the shelf, our THg concentrations in both late winter and spring (Table 1) are similar to THg concentrations in the southern Barents Sea Opening $(0.43 \pm 0.14 \text{ pmol } \text{L}^{-1})$ (Petrova et al., 2020) and comparable to previous expeditions in this area $(0.63 \pm 0.19 \text{ pmol } \text{L}^{-1}$ in summer, and $0.46 \pm 0.07 \text{ pmol } \text{L}^{-1}$ in early winter) (Kohler et al., 2022a). In late winter, shelf stations P1 - P5 exhibited relatively constant THg concentration profiles with depth, similar to early winter 2019 (Kohler et al., 2022a) while spring vertical THg profiles are similar to observed THg profiles in summer 2019 (Kohler et al., 2022a), with enrichments in surface waters < 50 m.



Figure 2: THg concentrations (top, pmol L⁻¹), MeHg concentrations (middle, fmol L⁻¹), and % MeHg as THg (bottom) along the shelf-basin transect in the northern Barents Sea during late winter (left) and spring (right) in 2021. The maximum plotted THg concentration range is 1.25 pmol L⁻¹. Depths where MeHg concentration < LOQ (7 fmol L⁻¹) have been removed from the visualization. The x-axis displays latitude and the nonlinear y-axis is in units of depth (m). Transect created with Ocean Data View (Schlitzer 2021).

278 MeHg concentrations (Table 1, Fig. 2 middle left) along the entire transect during late winter were 279 112 ± 76 fmol L⁻¹ (n = 58, ± 1 SD). At station P6 and P7, MeHg concentrations were lowest near

the surface and directly below the ice (UIW = $59 \pm 11 \text{ fmol } \text{L}^{-1}$, n = 2). At P7, MeHg concentrations 280 281 greater than 250 fmol L⁻¹ began at 300 m depth in the AW core of the West Spitsbergen Current. 282 By the spring cruise, MeHg concentrations along the entire transect were 69 ± 75 fmol L⁻¹ (n = 60, \pm 1SD) (Fig. 2, middle right) and covered a relatively large range (Table 1). In the upper 500 m, 283 284 our average MeHg concentrations on the spring cruise $(41 \pm 39 \text{ fmol } \text{L}^{-1}, n = 49)$ were significantly lower (log transformed, Student's t-test, p < 0.05) compared to late winter (85 ± 42 fmol L⁻¹, n =285 286 51). Shelf stations in spring exhibited low MeHg concentrations in general (Table 1), with the 287 lowest MeHg concentrations observed at P1 in mAW and some measurements below the LOQ of 7 fmol L⁻¹. Previous seasonal expeditions on the northern Barents Sea shelf showed similarly 288 289 constant MeHg distributions with depth, albeit with slightly higher average MeHg concentrations 290 (Kohler et al., 2022a). Our spring MeHg concentrations on the shelf however, are also lower than both the Barents Sea Opening (~ 60 ± 30 fmol L⁻¹) (Petrova et al., 2020) and among the lowest 291 reported for the Arctic region (Jonsson et al., 2022a). Percent MeHg (%MeHg) in both seasons 292 293 follows a similar distribution pattern as MeHg concentrations (Figure 2, bottom), with higher %MeHg observed in subsurface waters. In late winter, %MeHg is $20 \pm 13\%$ (n = 58) while spring 294 295 % MeHg is $10 \pm 10\%$ (n = 60). Similar to MeHg, there is a significant difference in % MeHg in the upper 500 m (log transformed, Student's t-test, p < 0.05) between late winter ($16 \pm 9\%$, n = 51) 296 297 and spring cruises $(7 \pm 8\%, n = 49)$.

298 The highest THg concentrations for both late winter and spring were observed in deep waters of 299 P6 and P7 between 500 - 1500 m depth and both maxima are observed 10 m from the bottom of 300 the Nansen Basin (SI Fig. 1). Our unfiltered THg observations in basin waters in both late winter 301 and spring are within the ranges reported in the central Arctic Ocean (Heimbürger et al., 2015, 302 Wang et al., 2018, Agather et al., 2019, Jonsson et al., 2022b) and the Fram Strait (Petrova et al., 303 2020). Similarly, relatively high MeHg concentrations were observed in basin waters > 500 m (SI 304 Fig. 1). These findings in late winter are similar to deep water MeHg concentrations observed in 305 the Fram Strait (Petrova et al., 2020) and the central Arctic Ocean (Jonsson et al., 2022b). 306 However, as we could not guarantee trace metal clean Niskin bottles for our deep water sampling 307 (> 500 m), some caution is needed when interpreting these results.

- **Table 1:** Average (± one standard deviation) and range of THg, MeHg, concentrations and %
- 310 MeHg on both late winter and spring cruises for all water column samples, shelf stations (P1 P5),
- 311 basin stations (P6, P7), under ice water (UIW), ice cores, and frost flowers (FF).

		Total Hg	(pmol L ⁻¹)			Methylated Hg (fmol L ⁻¹)					
	Late	winter	Spi	ring		Late winter			Spring		
	Average ± 1SD	Range	Average ± 1SD	Range		Average ± 1SD	Range	% MeHg as THg	Average ± 1SD	Range	% MeHg as THg
All water column data	0.56 ± 0.17 (n = 60)	0.40 - 1.61	0.69 ± 0.24	0.46 - 2.37 (n = 67)	All water column data	112 ± 76 (n = 58)	20 - 340	20 ± 13	69 ± 75 (n = 60)	< 7 - 288	10 ± 10
Upper 500 m only	0.53 ± 0.07 (n = 51)	0.40 - 0.77	0.64 ± 0.09	0.46 - 0.94 (n = 56)	Upper 500 m only	85 ± 42 (n = 51)	20 - 254	16 ± 9	41 ± 39 (n = 49)	<7 - 204	7 ± 8
Shelf (P1-P5) all depths	0.54 ± 0.07 (n = 34)	0.47 - 0.77	0.64 ± 0.08	0.46 - 0.84 (n = 40)	Shelf (P1-P5) all depths	71 ± 32 (n = 34)	20 - 161	13 ± 5	23 ± 13 (n = 33)	< 7 - 60	4 ± 3
Basin (P6, P7) all depths	0.59 ± 0.24 (n = 26)	0.40 - 1.61	0.76 ± 0.36	0.47 - 2.37 (n = 27)	Basin (P6, P7) all depths	171 ± 82 (n = 24)	72 - 340	30 ± 15	127 ± 79 (n = 27)	21 - 288	18 ± 10
UIW	0.58 ± 0.02 (n = 3)	0.56 - 0.60	$\begin{array}{c} 1.00 \pm \\ 0.30 \end{array}$	0.73 - 1.32 (n = 3)	UIW	59 ± 11 (n =2)	50 - 67	10 ± 2	36 ± 17 (n = 3)	26 - 55	4 ± 1
Ice core	0.27 ± 0.10 (n = 2)	0.20 - 0.34	1.83 ± 0.40	1.55 - 2.12 (n = 2)	Ice core	n/a			n/a		
FF	24.03 (n = 1)		n/a		FF	n/a			n/a		

To understand how changes in physical oceanography may affect our distributions of both THg and MeHg species, we averaged Hg concentrations by water mass in the upper 500 m, visualized in Fig. 3 and summarized in Table S1. Significant differences (log transformed, Student's t-test, p < 0.05) for both THg and MeHg concentrations in late winter and spring were observed for PW. A significant difference in THg concentrations was also observed for mAW, which was mainly observed in waters at station P1. No difference was observed for MeHg in mAW, although we assumed this is mainly due to local MeHg concentration differences in surface mAW at station P1 319 and subsurface mAW at stations P6 and P7. No statistically significant differences were observed 320 for THg nor MeHg concentrations in wPW, which comprised only subsurface and bottom waters 321 at our stations (Van Engeland et al., 2023). As PW comprises most of our shelf stations and surface 322 waters at P6 and P7, our observed changes in THg concentrations are best explained by external 323 inputs to surface waters such as atmospheric deposition or ice melt instead of changes in water 324 mass. In addition, our general transect (Fig. 2) and insignificant differences in wPW (Table 2) do 325 not suggest major inputs of benthic THg nor MeHg during this time period. We also cannot rule 326 out potential concentration differences in individual water parcels between sampling periods, but 327 we infer that our observed THg profiles and differences in THg concentrations in spring reflect 328 seasonal THg inputs to surface waters. Similarly, we hypothesize our lower MeHg concentrations 329 in spring are driven by internal biogeochemical processes rather than external inputs and/or water 330 masses.



332 Figure 3: THg (top) and MeHg (bottom) concentrations averaged for water masses in both late

- 333 winter (LW) and spring (SP) in the northern Barents Sea in 2021. Water masses are Atlantic Water
- 334 (AW), modified Atlantic Water (mAW), Polar Water (PW), and warm Polar Water (wPW). Box
- 335 plots display median (\pm IQR) and individual data points for each water mass. For averages and
- 336 sample sizes, see Table S1.

337 **3.3 Seasonal inputs of THg to surface waters**

338 Elevated THg in surface waters in spring could suggest inputs from AMDEs or sea ice processes, 339 where changing sea ice conditions facilitate atmospheric deposition to surface waters. Although 340 much of the AMDE-derived Hg is photochemically re-emitted back to the atmosphere from the 341 cryosphere during the summer (Dastoor et al., 2022), a temporary enrichment could exist during 342 spring. Under ice water (UIW) taken at stations P4, P6 and P7, in late winter was 0.58 ± 0.02 pmol L^{-1} (n = 3), similar to the remaining THg levels in the water column. However, the bottom 10 cm 343 344 of ice cores taken at P4 and P7 had low THg concentrations, with values of only 0.34 pmol L⁻¹ and 345 0.20 pmol L⁻¹, respectively. As sea ice appeared to both melt and freeze in between our two 346 sampling periods (SI Fig. 4), it is plausible that dynamic melt and freeze processes facilitated 347 additional THg to surface waters. However, changes in integrated THg in seawater (upper 100 m) 348 do not show a correlation with the average fraction of open water (SI Fig. 5). We suspect that the 349 heterogeneity in sea ice and snow may deliver variable amounts of Hg to surface waters, as the sea 350 ice covering our stations during the two seasonal cruises have different origins and drift patterns 351 (Marquardt et al., 2023a). Previous studies in the Arctic found higher concentrations and a wide 352 range of THg (range: 0.089 - 60.8 pmol L⁻¹) in both first year ice and multiyear ice compared to 353 seawater (Chaulk et al., 2011, Beattie et al., 2014, Schartup et al., 2020, Jonsson et al., 2022b). 354 Although we measured low THg concentrations in the bottom sea ice in late winter, THg can vary 355 by up to an order of magnitude throughout the core, with highest concentrations near the sea ice-356 atmosphere interface (Beattie et al., 2014). As sea ice drifts and snow and ice melts between cruise 357 samplings (SI Fig. 4), these reservoirs could supply additional Hg to surface seawater (Huang et 358 al., 2023). Furthermore, any leads in the ice pack during these changing ice conditions could also 359 facilitate direct atmospheric deposition of AMDEs to surface seawater or deposit additional Hg to 360 the cryosphere (Moore et al., 2014).

Refreezing also results in the rejection of brine between sampling periods (SI Fig. 4), with brine 361 THg concentrations potentially up to 355 pmol L⁻¹ (Chaulk et al., 2011). Although we did not 362 363 measure brine directly, a frost flower formed from brine expulsion in late winter had a concentration of 24 pmol L⁻¹ (n = 1). By spring, UIW concentrations were 1.00 ± 0.30 pmol L⁻¹ (n 364 365 = 3), and higher than the average concentration for the remaining water column. Likewise, in sea 366 ice, the THg concentration in the bottom 10 cm of ice cores taken at P4 and P7 were higher in spring, with values of 1.55 pmol L⁻¹ and 2.12 pmol L⁻¹, respectively. Spring THg concentrations 367 in the bottom of ice cores are higher compared to our late winter observations, which contrasts 368 369 with a seasonal study in Antarctica (Nerentorp Mastromonaco et al., 2016). Local melt and freeze 370 processes, downward brine movement, and varying sea ice advection are most likely responsible 371 for the differences observed in the two studies. However, seasonal increases in THg in the Antarctic water column from winter to spring were suggested to be a result of both atmospheric 372 373 and sea ice processes (Nerentorp Mastromonaco et al., 2017). Consequently, we conclude that our 374 observed seasonal THg water column concentrations most likely reflect similar seasonal inputs 375 and processes.

376 **3.4 Potential uptake of THg by the spring bloom**

377 Increased THg concentrations in surface waters and sea ice in spring could also be driven by the 378 onset of primary productivity (SI Fig. 6). Development of ice algae during the spring bloom can 379 take up THg from brine and seawater (Burt et al., 2013), explaining our increased THg 380 concentrations in the bottom of sea ice. In the water column, Spearman's rank correlation analysis 381 in spring indicated significant positive relationships between THg, particulate organic carbon 382 (POC) and chlorophyll a (Table S3). Additionally, elevated chlorophyll a, pico-, nano-, and 383 microphytoplankton in surface waters were observed in spring, indicating recent development of 384 the phytoplankton spring bloom (SI Figs. 6 & 9). Log-log regression models between THg, POC, 385 and chlorophyll a concentrations in the upper 500 m for all stations in spring (Fig. 4A, 4C) showed 386 a positive and statistically significant relationship (F-test, p < 0.05). Compared to late winter, this 387 analysis suggests that THg is more associated with the particulate phase. If we assume POC as a proxy for total particulate matter and a constant partition coefficient K_d to calculate dissolved Hg 388 389 (dHg) and particulate Hg (pHg) (SI Equations 3 and 4), this also indicates a greater percentage of 390 pHg in spring when compared to late winter. A strong positive Spearman rank correlation also

exists between POC and chlorophyll *a* (Table S3, p < 0.05), suggesting that the positive correlation of THg with both of these biological indicators may suggest adsorption and/or uptake of Hg species to primary producers during the spring bloom (Burt et al., 2014, Luengen and Flegal 2009).



Figure 4: Log scale scatterplots with log-log regressions of THg (left) and MeHg (right) concentrations against POC (top) and chlorophyll *a* (bottom) concentrations in the upper 500 m for both late winter (blue) and spring (orange) in the northern Barents Sea in 2021.

394

398 However, it is important to note that higher THg concentrations in spring suggest that seasonal 399 inputs increase both pHg and dHg concentrations, with a significant fraction of THg in the 400 dissolved phase. While our THg relationships to biological parameters during the spring bloom 401 suggests that THg is either taken up into primary producers by diffusion or adsorbed to cell 402 surfaces, the remaining dHg in the water column can also be complexed by extracellular algal 403 exudates (Cossart et al., 2022). In the dissolved phase, Hg shows strong affinity to DOM and 404 especially to reduced sulfur ligands (Ravichandran 2004). Indeed, we saw increased relative 405 intensities of dissolved organic sulfur compounds (CHOS) in surface waters in both sampling 406 periods compared to deeper waters (SI Fig. 7), most likely released by phytoplankton. These

biologically driven processes, such as uptake by phytoplankton and organic matter complexation,
could keep THg sourced from the atmosphere or sea ice in surface waters as well as facilitate
transformations of Hg species.

410 **3.5 MeHg dynamics in late winter and spring**

411 Significant differences in MeHg concentrations from late winter to spring can be driven by several 412 biogeochemical processes. As our spring MeHg concentrations showed a significant decline 413 compared to late winter, the collective sinks of MeHg in the water column outweigh MeHg 414 sources. Potential MeHg sinks to consider are photodemethylation, particle scavenging of MMHg, 415 atmospheric evasion of DMHg, and dark demethylation processes. To systematically evaluate the 416 contribution of each of these potential sinks, we used both literature and field data to calculate the 417 percentage contribution to the observed difference in the upper 100 m. It is important to note that 418 our observed differences in MeHg are net values, which can lead to high uncertainty when 419 estimating the contribution from various sources and sinks.

420 We integrated the water column for MeHg for each station to calculate the rate of change per day, 421 called Δ MeHg (Kohler et al., 2022a) for each station using the following equation:

422
$$\Delta MeHg = \left(\int_{0}^{z} [MeHg]_{SP}dz - \int_{0}^{z} [MeHg]_{LW}dz\right) / dt$$

Here, we used the trapezoidal integration method to calculate from 0 to 100 m depth, where z = depth, SP = spring, and LW = late winter, and t = time in days between sampling of each station. The total integrated pools are summarized in Table S4. All stations show a net decline, or loss, in the integrated pool from late winter to spring with an average of -81 ± 36 pmol m⁻² d⁻¹ (n = 7), demonstrating that sinks outweigh sources for observed change in MeHg concentrations.

428 Our measured declines of MeHg pools by spring could be due to increased photodemethylation 429 during the transition to polar day. However, both lower water temperatures and the presence and 430 thickness of snow and sea ice hinder MeHg photodemethylation (Point et al., 2009). Previous 431 estimates in the Arctic Ocean suggest photodemethylation rates of 0.04 d⁻¹ (Lehnherr et al., 2011) 432 and 0.0082 d⁻¹ (Zhang et al., 2020) at the sea surface. Assuming first-order kinetics and using 433 Δ MeHg, our calculated rate constant for our seven stations in the upper 100 m between late winter

to spring is an average MeHg loss of $0.021 \pm 0.013 \text{ d}^{-1}$ (n = 7). This loss, although a net loss and 434 not directly comparable, agrees well with the estimated photodemethylation rate constants; 435 436 however, the photodemethylation rate would be attenuated with depth and dependent on available 437 light. To estimate the percentage of the MeHg pool potentially degraded due to 438 photodemethylation at each station, we used photosynthetically active radiation (PAR) approximated to 15 E m⁻² d⁻¹ between the months of March and May 2021 (SI Fig. 8) and rate 439 constant from Arctic waters of 0.001 m² E⁻¹ when expressed to solar irradiance (Lehnherr et al., 440 2011). After accounting for the fraction of open water (with a simple assumption of zero 441 442 transmittance through snow-covered sea ice) and attenuation of PAR with depth (Table S6), our 443 calculations suggest that photodemethylation can only account for $5 \pm 4\%$ of our observed Δ MeHg 444 in the upper 100 m (Fig. 5). The highest percentage of photodemethylation (11%) is observed at station P1 due to no sea ice coverage. This calculation carries a high uncertainty due to additional 445 446 variables we could not account for, such as the partial transmittance of light through sea ice, 447 enhanced or inhibited photodemethylation in the presence of DOM, and scattering and absorption of light by both particles and DOM. Nevertheless, potential photodemethylation plays a relatively 448 449 small role compared to our observed changes in MeHg pools.



Figure 5: Percent contribution of selected processes influencing the change in MeHg pools, Δ MeHg, between late winter and spring. Photodemethylation is in yellow, MMHg particle scavenging in red, DMHg evasion in green, and dark demethylation (all other unaccounted for processes) in dark blue.

455 Particulate scavenging could also potentially serve to transport particle-bound MMHg (pMMHg) to deeper waters. As the total amount of sinking particles, including POC, are low in late winter 456 457 waters due to the lack of biological productivity (Dybwad et al., 2022), we could assume that scavenging of pMMHg is negligible. Furthermore, a previous study has estimated that only 4% of 458 459 pHg is pMMHg, which might be overestimated due to several samples below the detection limit 460 (Agather et al., 2019). Additionally, observed POC fluxes across our transect in late winter were low at 27 ± 9 mg C m⁻² d⁻¹ (Bodur et al., 2023) with no flux attenuation, suggesting minimal 461 vertical pHg transport. For comparison, POC fluxes in spring were 178 ± 199 mg C m⁻² d⁻¹ (Bodur 462 et al., 2023). While we only measured MeHg and do not know exactly when primary production 463 464 began to develop, we made several assumptions to attempt to determine the contribution of pMMHg scavenging to Δ MeHg (Table S6), with an estimate of $1 \pm 1\%$ of Δ MeHg (Fig. 5). 465

466 In surface waters, shifts in ice coverage could lead to evasion of DMHg trapped below the ice and help explain lower MeHg in surface waters during spring. In UIW, MeHg concentrations were 59 467 \pm 11 fmol L⁻¹ (n = 3) in late winter and 36 \pm 16 fmol L⁻¹ (n = 3) in spring. As our average UIW 468 469 concentrations of MeHg in spring were lower than in late winter, it is possible DMHg has evaded 470 to the atmosphere during melting and freezing, or DMHg has built up in the overlying sea ice 471 (Schartup et al., 2020). Previous studies have emphasized the importance of the evasion of DMHg from Arctic waters (Baya et al., 2015, Soerensen et al., 2016) but recent studies of DMHg in sea 472 473 ice and surface seawater (Agather et al., 2019, Jonsson et al., 2022) suggest a relatively modest 474 evasion rate. Although we did not measure DMHg, we expect a large proportion of our surface 475 MeHg concentrations to consist of MMHg, similar to other Arctic Ocean studies (Baya et al., 2015, 476 Agather et al., 2019, Jonsson et al., 2022b). We attempted to quantify the amount of DMHg evasion 477 during our transition to spring with additional limitations and assumptions (Table S6), with 478 atmospheric evasion making up only $3 \pm 3\%$ of Δ MeHg (Fig. 5). As this estimate is a gross value 479 and does not account for atmospheric re-deposition of MMHg, it may be overestimated.

480 Water mass distribution along our transect did not change drastically from late winter to spring 481 (Van Engeland et al., 2023), suggesting relatively minimal importance of advection of lower 482 concentration MeHg waters in the upper 100 m. However, station P6 and P7 do intercept the West 483 Spitsbergen Current, with slight shifts from wPW to PW in spring (Van Engeland et al., 2023). 484 MeHg concentrations in PW are lower than wPW in late winter (One-way ANOVA with Tukey's test, p < 0.05), and could contribute to the observed Δ MeHg. Nevertheless, PW is significantly 485 486 different between the two cruises, so for the sake of simplicity we have assumed advection plays 487 a negligible role in explaining our observed Δ MeHg.

The importance of dark demethylation processes in unfiltered waters has been highlighted in previous studies using spiked stable Hg isotopes to measure demethylation rates (Lehnherr et al., 2011, Monperrus et al., 2007). In addition, the recent MeHg annual budget for the East Siberian Sea attributed approximately 50% of potential MeHg sinks to gross dark demethylation (Kim et al., 2020b). Consequently, we attribute the remaining Δ MeHg observed (90 ± 8%, Fig. 5) as dark demethylation processes and hypothesize that they are mainly driven by biological activity.

494 **3.6 Biotic demethylation of MeHg**

495 As POC fluxes were low in late winter (Bodur et al., 2023), it is likely that there is no substantial 496 in situ biotic Hg methylation occurring throughout the water column. Additionally, despite a higher 497 concentration of THg by spring, both MeHg concentrations and % MeHg have decreased (Fig. 2), 498 suggesting methylation processes were dominated by demethylation processes. However, the 499 increase in phytoplankton abundance by orders of magnitude in surface waters along the transect 500 in spring could support biotic transformations of Hg species (SI Fig. 6). Upon development of the 501 phytoplankton bloom, we observed that diatoms, particularly species of the genus *Thalassiosira*, 502 had the highest relative abundance in the upper 90 m at five out of seven stations along our transect 503 (SI Fig. 9). Our regression analysis (Fig. 4) supports the hypothesis that THg is indeed potentially 504 taken up by primary producers. Unlike THg, however, MeHg has significant negative relationships 505 with both POC and chlorophyll a in spring (Fig. 4B, 4D, F-test, p < 0.05), supporting declining 506 MeHg concentrations in association with biological particles. As diatoms and other marine 507 phytoplankton passively take up MMHg (Mason et al., 1996, Lee and Fisher 2016), we 508 hypothesize that our observed MeHg losses in surface waters could be due to phytoplankton uptake 509 and subsequent demethylation. This hypothesis is further supported by relatively low contributions 510 of the other MeHg sinks in surface waters (Fig. 5) and possible biological demethylation mechanisms identified in literature. Previous research has observed reduction of inorganic Hg^{II} to 511 Hg⁰ in the diatom Thalassiosira weissflogii (Morelli et al., 2009) while the green alga 512 513 Chlamydomonas reinhardii was demonstrated to demethylate MMHg (Bravo et al., 2014). 514 Another study in a freshwater lake has shown that demethylation of intracellular MMHg correlated 515 positively with diatom abundance (Cossart et al., 2021). Recently, a study suggested that 516 phytoplankton can intracellularly demethylate MeHg through the production of reactive oxygen 517 species (Liang et al., 2023).

518 This MMHg uptake during the onset of the Arctic spring bloom could potentially facilitate the 519 seasonal bioaccumulation of MMHg by higher trophic level organisms throughout the summer. A 520 previous study in an Arctic lake observed decreasing MeHg levels in water from winter to spring, 521 yet saw a seasonal increase in MeHg concentrations in freshwater zooplankton (Varty et al., 2021). 522 Furthermore, another winter to spring increase of MeHg concentrations in zooplankton was 523 observed in waters of the Canadian Arctic Archipelago (Pućko et al., 2014). While other 524 mechanisms like zooplankton grazing activity or direct methylation in the gut could explain their observed seasonal Hg biota concentrations (Pućko et al., 2014), we speculate that our seasonal 525 526 changes of water column MeHg partly reflect a MMHg biomagnification pathway for higher 527 trophic level organisms in the Arctic's productive spring and summer seasons.

528 Further studies on Hg methylation activity in specific marine phytoplankton species demonstrated 529 that MeHg demethylation positively correlated with thiol concentrations in algal exudates (Li et 530 al., 2022). We observed more elevated relative intensities of CHOS compounds in surface waters 531 compared to deeper waters (SI Fig. 7). Dissolved CHOS compounds could facilitate the 532 transformation of MMHg to DMHg via localized reduced thiol groups (Jonsson et al., 2016), with a proposed disproportionation reaction mechanism of two MMHg molecules to form DMHg and 533 iHg. Inorganic Hg could be reduced to Hg⁰ or be scavenged, while DMHg could evade to the 534 535 atmosphere as previously highlighted. Complexation of MMHg to dissolved thiols in DOM could 536 also reduce uptake of MMHg by phytoplankton (Lee and Fisher 2017) reducing MMHg 537 bioavailability. MMHg-DOM complexes can also undergo enhanced photodemethylation (Qian et 538 al., 2014) in surface waters, although the specific mechanisms remain debated (Barkay and Gu 2021). Additional Arctic seasonal studies are needed to characterize the types of DOM and its role
in driving MeHg biogeochemical processes in the water column. Collectively though, our data and
literature support biotically mediated MeHg demethylation during the development of the spring
bloom.

543 Interestingly, deeper waters (100 to 500 m) also show a net loss of MeHg concentrations, with an average loss rate of 0.015 ± 0.011 d⁻¹ (n = 7) in the 100 - 200 m depth interval. While potential 544 phytoplankton demethylation could explain MeHg losses in surface waters, we expect different 545 546 biotic processes to affect MeHg losses in subsurface and bottom waters. Recent research has 547 demonstrated that the presence of the *merB* gene, found in certain bacteria and responsible for the demethylation of MeHg to Hg^{II}, is present in deep waters of the Arctic Ocean (Bowman et al., 548 2020). In bacteria, the mer operon, encoding for merA and merB genes, exists primarily as a 549 550 detoxifying mechanism for high concentrations of Hg, in a process called reductive demethylation. 551 Lineages of a particular bacterial strain genome containing merA and merB, with high resistance 552 and strong degradation capacity of MeHg, are present in polar regions (Sanz-Sáez et al., 2022). 553 On our transect, potential reductive demethylators (*Alteromonas, Marinobacter*, and *Oleibacter*) 554 are present, but in low relative abundance during both sampling periods compared to the entire 555 bacterial and archaeal community (SI Text). However, bottom waters at shelf stations had higher 556 total abundances of bacteria and higher relative abundances of potential demethylators than surface 557 waters in late winter and could support MeHg degradation during the winter-spring transition (SI 558 Text). This includes methanotrophic bacteria and sulfate reducing bacteria, which can also 559 participate in oxidative demethylation of MeHg under anoxic conditions (Barkay and Gu, 2021).

560 While dark abiotic demethylation could influence our losses of MeHg in deep waters, more 561 research is needed to uncover these mechanisms. Dissolved CHOS compounds, seen in both 562 surface and deeper waters in both late winter and spring (SI Fig. 6), could facilitate transformation 563 and demethylation of MMHg as previously discussed. Additionally, resuspension of sediments 564 may supply manganese oxide to bottom waters, resulting in both the demethylation and scavenging 565 of MMHg and iHg (Vlassopolous et al., 2018, Murphy et al., 2021, Kohler et al., 2022a). However, 566 due to the uncertainty of these potential abiotic processes, we hypothesize much of our MeHg in 567 deeper waters is also biotically demethylated from late winter into spring.

568 **3.7** Hg in basin deep waters (> 500 m)

569 Bottom waters below 500 m at stations P6 and P7 are elevated in THg compared to the shelf 570 stations (SI Fig. 1). These elevated concentrations in EBDW could contain additional Hg from 571 resuspended sediment or sinking particles, as suggested in the Canadian Basin (Agather et al., 572 2019). As these basin stations are close to the shelf break, turbidity along the continental slope 573 caused by the West Spitsbergen Current may resuspend and deliver Hg-rich particles into the basin. 574 Throughout the Arctic winter (December to March), overturning on the Barents Sea shelf generates 575 a nepheloid layer of suspended sediment which can sink as dense water plumes into the basin 576 (Rudels et al., 2004, Gardner et al., 2022, Rogge et al., 2022). High concentrations of Hg and other 577 trace elements in surface sediments in the Eurasian Basin have suggested hydrodynamic transport 578 and scavenging of particles sourced from the shelf (Kohler et al., 2022b). Beam transmittance data 579 (SI Fig. 10), however does not support significant resuspension of sediments in basin waters. For 580 MeHg, stations P6 and P7 show similar patterns in late winter and spring, with the highest MeHg 581 concentrations observed at 500 m and below. In EBDW, MeHg concentrations remain elevated above 170 fmol L⁻¹ at P7 until 2500 m depth. Compared to the shelf, EBDW and deeper waters 582 583 remain enriched in MeHg. These observations are in line with previous measurements in the Fram 584 Strait with maximum MeHg between 400 – 1000 m (Petrova et al., 2020). Diffusion of porewater-585 derived MeHg could help explain elevated MeHg concentrations along the continental slope by 586 station P6 (Kim et al., 2020b). Although both MeHg and THg concentrations in the deep basin 587 agree with previous literature (Wang et al., 2012, Heimbürger et al., 2015), our sampling with 588 standard Niskin bottles lends a degree of caution.

589 **3.8** Seasonal cycle of Hg in the northern Barents Sea

590 Our observations for both THg and MeHg suggest a distinct seasonality of Hg in the northern 591 Barents Sea marginal ice zone from late winter to spring. Based on our analysis, we conclude that 592 the main drivers of both THg and MeHg changes in surface waters are ultimately driven by 593 seasonal Hg inputs and the biotically driven processes of the phytoplankton spring bloom. In 594 deeper waters, biotic demethylation is likely to play a more important role than abiotic 595 demethylation mechanisms. A previous study suggested iHg seasonality in this region during the 596 transition from summer to winter, with no significant change in MeHg concentrations (Kohler et 597 al., 2022a). By assuming Hg follows a distinct seasonal cycle in the Arctic Ocean every year, we

- 598 combined our observations from late winter and spring 2021 with that of summer and early winter
- 599 2019 (Kohler et al., 2022c,d) to illustrate a simple seasonal cycle of Hg over the course of a year
- 600 in the northern Barents Sea surface waters (Fig. 6). To best illustrate the seasonal dynamics of THg
- and MeHg, we integrated our results from the surface to 100 m depth for each station (Tables S4
- 602 & S5), where most biological productivity and Hg input occurs.



Seasonal Cycle of Hg in the Northern Barents Sea

●--P1 -●--P2 --●--P3 -●--P4 -●--P5 --●--P6 -●--P7



Figure 6: Integrated water column (0 to 100 m) observations of THg (top) and MeHg (bottom) from seasonal cruises (this study) and a previous study (Kohler et al., 2022a) in the northern Barents Sea. Graphs are superimposed on a simplified Arctic seasonal cycle in the marginal ice zone with polar day occurring between April and August and polar night beginning November until February. Green particles represent the development of the ice algal and phytoplankton spring blooms. Integrated pools for THg and MeHg are found in Tables S4 and S5. The x-axis is in months. Note the difference in scales on the y-axis.

611 Seasonal patterns of both THg and MeHg emerge and appear to be driven by the development of 612 biological activity. The THg pools are higher during the polar day, while MeHg pools are at the 613 minimum at the onset of the phytoplankton bloom and maximum in the second half of the year. 614 As hypothesized, seasonal inputs and biological activity increase THg concentrations in surface 615 waters. To better assess how relevant environmental drivers impact seasonal THg and MeHg 616 concentrations, a PCA was conducted for each seasonal cruise (Fig. 7), with the first two principal 617 components explaining > 60% of the variance for each cruise (Table S7). Clustering of THg with 618 biological parameters POC and chlorophyll *a* in spring and summer suggest seasonal changes in 619 THg concentrations are driven by biological activity and its partitioning to the particulate phase. Notably, this includes potential uptake of Hg species, transformation to MeHg during 620 621 remineralization of POC, and downward transport with sinking particles. Sinking POC supports 622 the production of subsurface in situ MeHg throughout polar day (Heimbürger et al., 2015), 623 reflected in increasing MeHg pools progressing into summer (Fig. 6) and the general trend of 624 positive MeHg loadings on the first component with depth, nutrients, and salinity for the spring 625 and summer cruises. Late summer POC fluxes (Bodur et al., 2023), and increased proportions of 626 manganese oxides leading into the polar night (Xiang and Lam 2020) lower THg pools by early 627 winter (Kohler et al., 2022a).

Less clear is the behavior of both THg and MeHg throughout the polar night into late winter. It was proposed that potential scavenging by manganese oxides could be a driver of THg during Arctic winter (Kohler et al., 2022a), although a negative trend in THg pools from early winter to late winter is not observed (Fig. 6). We speculate that changes in Hg speciation, such as increased proportions of gaseous Hg⁰ and DMHg could protect against scavenging losses. A recent experimental study argued that gross dark reduction rates of Hg^{II} to Hg⁰ are significant in both 634 photic and aphotic zones, with dark reduction most likely dominant in the Arctic except during 635 polar day (Lamborg et al., 2021). Reduction could be driven by bacteria with the merA gene, and 636 potentially include bacteria with both merA and merB, facilitating declining MeHg pools throughout the polar night (Fig. 6). The bacterial community in this region shifts from carbon 637 cycling in summer to nitrogen cycling in winter (Thiele et al., 2023), suggesting a possible link 638 between MeHg and the nitrogen cycle (Despins et al., 2023) during the polar night. In our PCA 639 640 analysis, we observed that MeHg and nitrite (NO₂) exhibited opposite loadings on PC2 for all cruises (Fig. 7). This observation could link nitrite-oxidizing *Nitrospina*, known to harbor and 641 642 express hgcAB-like proteins for Hg methylation (Gionfriddo et al., 2016, Villar et al., 2020), as a 643 driver for in situ seasonal MeHg concentrations in the Arctic. Our in situ concentration data 644 however, show a net loss of the MeHg pool from winter to late winter (Fig. 6). As our studies comprise two different sampling years (2019 vs. 2021), more seasonal and interannual studies are 645 646 needed to fully understand the drivers of MeHg concentrations in the Arctic water column.



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Figure 7: Principal component analysis (PCA) loadings plot for each seasonal cruise in the
 northern Barents Sea from this study and previous cruises where Hg observations are available (<

500 m only) (Kohler et al., 2022a). Rotation is orthogonal with the first two principal components
plotted. Selected environmental parameters included depth, conservative temperature
(ConsvTemp), absolute salinity (AbsSal), bacterial abundance (bacteria), chlorophyll *a* (Chla),
particulate organic carbon (POC), nitrate (NO3), nitrite (NO2), phosphate (PO4), and silicate
(SiOH4), THg, MeHg, and %MeHg.

655 To the best of our knowledge, only one model has attempted to represent seasonal cycling of Hg 656 in the Arctic Ocean (Fisher et al., 2012). In this model, sources and sinks of Hg are categorized by 657 month, with subsurface waters resupplying Hg to surface waters through vertical mixing. However, the latest Arctic budget highlights that export of Hg to deeper waters is most likely 658 659 underestimated (Dastoor et al., 2022), further supporting our hypothesis that seasonal changes in 660 Hg may be driven by spring Hg inputs and changes in the particulate phase. Our new seasonal data 661 help fill in this knowledge gap but continue to remain a "seasonal snapshot" from both a seasonal 662 and interannual perspective. Recent studies on seasonality in this region highlight large seasonal 663 and interannual variability in the northern Barents Sea regarding water masses and sea ice 664 dynamics (Jones et al., 2023, Kohlbach et al., 2023, Van Engeland et al., 2023). Specifically, 665 interannual variability may exert a greater impact on perceived seasonality, as our four cruises were sampled in two different years (2019 vs. 2021). As a result, our interpretations between the 666 667 spring to summer transition and the early winter to late winter transition are subject to both 668 seasonal and interannual variability, and thus more speculation. Nevertheless, we propose that our 669 seasonal cycle of Hg in the northern Barents Sea is supported by our observational data and is 670 reflected in current literature on Arctic Hg cycling.

671 **4 Conclusion**

We show that both THg and MeHg concentrations in the water column of the northern Barents Sea exhibit dynamic changes throughout the year. Variations in THg and MeHg are driven by seasonal processes during the early spring, such as Hg input to surface waters and biological activity. Notably, we observed a significant decline in MeHg concentrations during the development of the phytoplankton spring bloom. We suggest biotic demethylation mediated by phytoplankton and bacteria as the main mechanism underlying our observations. This demethylation by phytoplankton and bacteria during the spring bloom may mitigate the bioaccumulation of Hg up the food web. Although it is difficult to completely disentangle these
losses and transformations of MeHg, our data suggests that potential seasonal uptake of MeHg
during the spring bloom represents an important entry point of Hg into the Arctic marine
ecosystem.

683 Using our current data, we constructed a simplified seasonal Hg cycle in the northern Barents Sea 684 marginal ice zone and conducted additional analysis to try and identify major environmental 685 parameters driving both THg and MeHg concentrations. THg was clustered with chlorophyll a and 686 POC in spring and summer, suggesting seasonal changes of THg are modulated by changes in 687 primary productivity, particles, and food web processes. MeHg concentrations also exhibited a 688 seasonal pattern, with potential uptake and demethylation during spring, followed by hypothesized 689 in situ production throughout the summer. Our simple seasonal cycle also encompasses potential 690 interannual variability, which may underestimate additional drivers of Hg concentrations such as 691 sea ice coverage and changing water masses. In addition, the northern Barents Sea has limited 692 riverine inputs, highly variable sea ice coverage and a unique water mass distribution compared to 693 the other Arctic shelf seas. Thus, the applicability of our proposed seasonal cycle may have limited 694 use in other Arctic regions, like the river-influenced Siberian shelf seas or less biologically 695 productive Central Arctic Ocean. Nevertheless, our study represents an important first step in 696 identifying Hg seasonal processes and drivers in Arctic waters, with additional seasonal 697 environmental data from the Nansen Legacy project available for use in budgets and models. We 698 suggest future Arctic Hg budgets and coupled ocean-atmosphere/earth system models include an 699 element of seasonality.

700 The projected climate driven inputs of increasing amounts of Hg from melting glaciers and 701 thawing permafrost to Arctic shelf seas during the onset of the productive season may facilitate 702 greater uptake of MMHg in the marine food web (Chételat et al., 2022). As ongoing climate change 703 continues to transform the Arctic region, more research is needed to elucidate how region-specific 704 seasonal processes affect Hg distributions in the Arctic Ocean and the surrounding shelf seas. 705 Specifically, we suggest more studies examining the connection between demethylation and 706 biological uptake in the marginal ice zones as well as seasonal measurements of gaseous Hg 707 species. We suspect that these transformation processes and rates that control MeHg distributions 708 in the water column, and subsequently uptake in biota, follow a seasonal pattern.

709 CRediT authorship contribution statement

710 Stephen G. Kohler: Conceptualization, methodology, validation, formal analysis, investigation, 711 writing - original draft, visualization, funding acquisition. Lars-Eric Heimbürger-Boavida: conceptualization, methodology, validation, investigation, writing - review and editing, 712 713 visualization, supervision. Philipp Assmy: investigation, formal analysis, writing - review and 714 editing, visualization. Oliver Müller: investigation, formal analysis, writing - review and editing, 715 visualization. Stefan Thiele: investigation, formal analysis, writing - review and editing. Maria 716 G. Digernes: Investigation, formal analysis, visualization. Kuria Ndungu: writing - review and 717 editing, supervision. Murat V. Ardelan: Conceptualization, methodology, writing - review and 718 editing, funding acquisition, supervision.

719

720 Acknowledgements

721 Funding for this study was provided by The Nansen Legacy project (RCN# 276730) and the 722 Anders Jahres fond til vitenskapens fremme. The authors would like to thank the captain, crew 723 members, cruise leaders, and fellow scientists onboard R/V Kronprins Haakon for cruises Q1 and Q2. Specifically, the authors would like to thank L. Kull and M. Adams for assistance in collecting 724 725 mercury samples onboard and N. Torres-Rodriguez and A. Dufour for assistance in analysis. 726 Additional thanks are to the sea ice coring teams and under ice water sampling teams for help in 727 collecting ice samples. We also thank N. Sanchez and M. Petrova for their insightful comments on 728 this manuscript. Lastly, we thank F. A. Cnossen for her artistic design of the graphical abstract.

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1007 Flow cytometry measurements (abundance of virus, bacteria and small protists (primarily <20μm))
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1009 Sea (2023c) https://doi.org/10.21335/NMDC-39569968

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1139	Supplementary Material
1140	
1141	Biotic transformation of methylmercury at the onset of the Arctic spring bloom
1142	
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1154 Integration of THg and MeHg

1155 We used trapezoidal integration for the upper 100 m in the water column for both THg and MeHg 1156 concentrations (Kohler et al., 2022). We assumed that the first available sampling depth (10 m or 1157 UIW depending on station) represented the sea ice-ocean interface (0 m) or the atmosphere-ocean 1158 interface (0 m). For 100 - 200 m integration, stations P2 and P5 were integrated to bottom sampling 1159 depths less than 200 m. Stations were linearly interpolated to 100 m using the closest available sampling depths. The exact sampling dates and days between for each station were used to 1160 1161 calculate the rate constants and Δ MeHg values. Where MeHg samples were less than the LOD or LOQ, the LOD (2 fmol L⁻¹) and LOQ (7 fmol L⁻¹) were used as the sample depth's concentration 1162 for integration, respectively. As most samples are run in singles, uncertainty in Δ (±1SD) for late 1163 winter and spring Δ MeHg and Δ THg was calculated using the absolute THg reproducibility 1164 reported (± 0.05 pmol L⁻¹) and absolute MeHg reproducibility reported (± 7 fmol L⁻¹) for all depths 1165 1166 and propagated using the sum of squares method for integration, and reported as ± 1 SD.

1167 Depth matching for environmental parameters

1168 THg and MeHg sampling geographic coordinates and depths differ slightly from environmental1169 parameter sampling geographic coordinates due to ship drift during multiple samplings. Station

- 1170 samplings were assumed to be identical despite multiple sampling CTD casts and deviance from
- 1171 exact coordinates. For Spearman rank analysis, regression analysis, and PCA, environmental data
- 1172 were linearly interpolated to sampling depths of THg and MeHg for each cruise using the closest
- 1173 sampling depths when necessary. Otherwise, matching depths were assumed to be from the same
- 1174 water parcel for data analysis.

1175 Percentage of ΔMeHg calculations

1176 *Photodemethylation*

Delta photodemethylation (Table S6, Δ MeHg_{pd}, pmol m⁻² d⁻¹) was estimated using a similar 1177 equation (1) as Kim et al., 2020. MeHg concentrations (C_{MeHg}, pmol L⁻¹) at each station were 1178 1179 linearly interpolated for each depth in the surface 100 m in late winter using our observed 1180 concentrations at discrete depths. The photodemethylation rate was estimated based on satellite PAR (I₀, 15 E m⁻² d⁻¹) in the region in April 2021 multiplied by the photodemethylation rate 1181 constant (k_{pd}, 0.001 m² E⁻¹) (Lehnherr et al., 2011). The k_{pd} was assumed constant over 100 m, 1182 while PAR (I) was attenuated by depth (z) using equation (2) found in Kim et al., 2020, with 1183 1184 original reference Paulson and Simpson 1977, type I (clear water). We accounted for the sea ice coverage by multiplying by the average fraction of open water (f_{ow}) at each station in the previous 1185 1186 60 days from sampling in May, assuming zero transmittance through snow-covered sea ice.

1187
$$\Delta MeHg_{pd} = \int_0^{100} C_{MeHg} * (f_{OW} * PAR * k_{pd})$$
(1)

1188
$$\frac{I}{I_0} = Re^{\frac{Z}{\xi_1}} + (1-R)e^{\frac{Z}{\xi_2}}$$
(2)

1189 Scavenging of MMHg

1190 Delta MMHg scavenging (Table S5, Δ MeHg_{scav}, pmol m⁻² d⁻¹) was estimated using several 1191 assumptions. First we calculated pHg in the upper 100 m using our THg concentrations at each 1192 discrete depth in late winter. To do this, we used in situ POC concentrations (Marquardt 2022a, 1193 2022b) at each sampling depth as a proxy for total particulate matter and K_d for Hg in the Arctic 1194 shelf (1.38*10⁶ L kg⁻¹, Tesán Onrubia et al., 2020) to calculate dHg (3) and pHg (4). We also 1195 assumed that K_d does not change seasonally nor with depth. Then, we assumed that pMMHg makes 1196 up approximately 4% of pHg (Agather et al., 2019). We used our estimated pMMHg and POC concentrations to estimate a pMMHg/TPM ratio in the upper 100 m, by assuming that POC in late 1197 1198 winter makes up approximately 25% of TPM (Dybwad et al., 2022). We used the average late 1199 winter POC fluxes reported in Bodur et al., 2023 using a short term sediment trap method (27 mg C m⁻² d⁻¹) and assumed it represented 25% TPM flux of 108 mg m⁻² d⁻¹, which is in agreement 1200 with previous TPM fluxes in the area determined by long term moorings (Dybwad et al., 2022). 1201 1202 As POC fluxes were not attenuated with depth in late winter (Bodur et al., 2023), we multiplied our average pMMHg/TPM ratio by our estimated TPM flux to calculate a pMMHg flux at 100 m 1203 depth (5), for an average of 0.76 ± 0.06 pmol m⁻² d⁻¹ for Δ MeHg_{scav}. 1204

1205
$$dHg = \frac{THg}{((K_d * POC) + 1)}$$
(3)

$$pHg = THg - dHg \tag{4}$$

1207
$$\Delta MeHg_{scav} = pMMHg_{TPM} * TPM flux$$
(5)

1208 Evasion of DMHg

We followed a similar approach to Kim et al., 2020 (and references therein) for calculating delta 1209 1210 DMHg evasion (Table S6, Δ MeHg_{ev}) (6). This included using estimated windspeeds from 1211 Longyearbyen airport in March and April 2021 ($\sim 8 \text{ m s}^{-1}$), surface water temperatures ($\sim 10 \text{ m}$ depth), at each station in late winter, temperature dependent Schmidt numbers for CO₂ and DMHg 1212 1213 to calculate the DMHg transfer coefficient (K_w), and temperature dependent Henry's law constant 1214 for DMHg (H). We estimated the DMHg (C_{DMHg water}) concentration in surface waters using the % DMHg as MeHg as reported by Jonsson et al., 2022 (~ 6% in primary mixed layer) at shallowest 1215 sample depth for MeHg in late winter, and a literature value of 0.023 pmol m⁻³ for DMHg in air 1216 1217 (C_{DMHg air}, Baya et al., 2015). We multiplied our estimated DMHg evasion flux by the average 1218 fraction of open water (fow) in the previous 60 days at each station from sampling in May, 1219 assuming sea ice completely restricts atmospheric evasion (DiMento et al., 2019).

1220
$$\Delta MeHg_{ev} = K_w * \left[C_{DMHg_{water}} - \left(\frac{C_{DMHg_{air}}}{H}\right)\right] * f_{OW}$$
(6)

1222 Dark demethylation

1223 We calculated percentage dark methylation (Δ MeHg_{dark}) as the proportion of Δ MeHg unaccounted 1224 for from the sum of Δ MeHg_{pd}, Δ MeHg_{scav} and Δ MeHg_{ev} for each station.

1225 Flow cytometry (FCM) Analysis

1226 To measure the abundance of microorganisms (prokaryotes [bacteria and archaea], pico- and nanophytoplankton), four replicate water samples (1.8 mL) were taken, fixed with glutaraldehyde 1227 1228 (0.5 % final concentration) at 4 °C for a minimum of 2 h, flash-frozen in liquid nitrogen and stored 1229 at -80 °C until analysis. Samples for the analysis of prokaryote abundance were first diluted 10 1230 times with 0.2-µm-filtered TE buffer (Tris 10 mM and EDTA 1 mM, pH 8), then stained with a 1231 green fluorescent nucleic acid dye (SYBR Green I; Molecular Probes, Eugene, OR, USA) and 1232 incubated for 10 min at 80 °C and measured on a FACS Calibur (Becton Dickinson, Oxford, UK) 1233 flow cytometer (Marie et al., 1999). Samples were counted at a low flow rate of around 60 µL 1234 min-1 and groups discriminated on a biparametric plot of green fluorescence (BL1) vs. side scatter 1235 (SSC). Pico- and nanosized phytoplankton abundance was measured using an Attune® Acoustic 1236 Focusing Flow cytometer (Applied Biosystems by Thermo Fisher Scientific, Waltham, MA, USA) 1237 and the different phytoplankton groups were discriminated based on their red fluorescence (BL3) 1238 vs. orange fluorescence (BL2) (Paulsen et al., 2016). The data for all cruises can be found at the 1239 Norwegian Marine Data Centre (Müller et al., 2023a, 2023b, 2023c, 2023d).

1240 Bacterial Community Analysis

1241 We investigated the bacterial and archaeal community with respect to potential demethylators 1242 (Christakis et al., 2018), methanotrophs, and sulfate reducing bacteria (SRB) using 16S rRNA gene 1243 amplicon sequences collected on our transect to help explain the MeHg concentrations at different depths. Using PICRUSt2 with standard parameters in bioconda, the metabolic potential of the 1244 1245 different amplicon sequence variants was inferred based on their location in a phylogenetic tree of 1246 fully sequenced organisms and the genomic assets of the closest relative in this tree (Grüning et 1247 al., 2018, Douglas et al., 2020). Using KEGG-numbers, the marker genes merA and merB 1248 (reductive demethylation), dsrA (sulfate reduction), and mmoX (methanotrophy) were extracted 1249 and used to identify SRB, methanogens, and methanotrophs. For merA and merB, only Oleibacter

1250 was identified from the samples and was complemented with manual searches for *Alteromonas* 1251 and Marinobacter (Sanz-Sáez et al., 2022). Similarly, no mmoX genes were predicted and known 1252 methanotrophs were extracted based on recent literature (Guerrero-Cruz et al., 2021). SRB were well covered by the PICRUSt2 results. In the water column, the relative abundance of potential 1253 1254 reductive demethylators and SRB was highest in the deepest sample of station P7 with 1.9% and 0.7% (potential reductive demethylators) and 4.8% and 3.9% (SRB) in late winter and spring. 1255 1256 Potential reductive demethylators were otherwise low in relative abundance, rarely exceeding 1257 0.1% of the total bacterial and archaeal community. Higher abundance was found in most samples 1258 at 10 m and below 200 m depth as compared to the samples in between, regardless of bottom depth for both seasons, although on average the abundance was higher in late winter ($\sim 0.1\%$) than in 1259 spring (~0.06%). SRB were similarly abundant in both seasons with 0.39% in late winter and 1260 0.37% in spring but were lower in surface samples (10 and 20 m) as compared to deeper samples. 1261 1262 Methanotrophs were also low in abundance with $\sim 0.15\%$ both in late winter and spring. The highest abundance was found in both seasons at P1 (near bottom depth) with 1.3% in spring as 1263 compared to 0.7% in late winter. Generally, the abundance was higher in samples > 20 m. 1264

1265 Dissolved Organic Matter (DOM) Analysis

1266 DOM seawater and under ice water samples were collected at stations P1, P4, P6, and P7 on both late winter and spring sampling expeditions. Water samples were immediately extracted after 1267 collection using solid phase extraction (Dittmar et al., 2008) and stored at -20°C until analysis. 1268 1269 Extracts were analyzed by LTQ-Velos-Pro Orbitrap mass spectrometer (Thermo Scientific, 1270 Germany) with an electrospray ionization source (ESI) in negative mode. Each sample spectrum was calibrated internally in mass range 200-800 m/z with data resolution of 100 000. Additional 1271 1272 detailed instrument parameters are noted in Hawkes et al., 2016. Dissolved organic sulfur compounds (CHOS) formulas are determined based on the following elemental criteria: C₄₋₅₀ H₄₋ 1273 $_{100}$ O 2-40N 0-2S 0-1, hydrogen to carbon ratio (H/C) = 0.3 - 2.2, oxygen to carbon ratio (O/C) = 0-1, 1274 1275 double bond equivalent-oxygen (DBE-O) = -10 - 10, valence electron as even number, mass to charge ratio (m/z) = 150-800 and mass defect (mass after nominal mass) = -0.1 - 0.3. 1276

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SI Figure 1: Complete depth profiles of THg (left, pmol L⁻¹) and MeHg (right, fmol L⁻¹) for stations P6 (top) and P7 (bottom) in late winter (blue) and spring (orange) in the northern Barents Sea in 2021.



1388 SI Figure 2: Conservative temperature (Θ , °C) along the shelf-basin transect in the northern 1389 Barents Sea during late winter (top) and spring (bottom) in 2021. The x-axis displays latitude and 1390 the nonlinear y-axis is in units of depth (m). Transect created with Ocean Data View (Schlitzer 1391 2021).



SI Figure 3: Absolute salinity (S_A, g kg⁻¹) along the shelf-basin transect in the northern Barents
Sea during late winter (top) and spring (bottom) in 2021. The x-axis displays latitude and the
nonlinear y-axis is in units of depth (m). Transect created with Ocean Data View (Schlitzer 2021).



1396

SI Figure 4: Fraction of sea ice coverage in the northern Barents Sea for stations P2 - P7 during
January 1 - May 30, 2021. P1 had no ice coverage. An ice fraction of 1 indicates complete sea ice
coverage and 0 indicates open water. Sampling periods are outlined in rectangles, with late winter
(blue) and spring (orange). The y-axis is fraction of sea ice coverage (0 - 1) and the x-axis is in
days starting from January 1, 2021.



1403 **SI Figure 5**: Linear regression of Δ THg (pmol m⁻² d⁻¹, left) and Δ MeHg (pmol m⁻² d⁻¹, right) in 1404 the upper 100 m against the average fraction of open water in between sampling periods (60 day-1405 average) in the northern Barents Sea in 2021. No significance (F-test, p > 0.05) was observed for 1406 either regression.



1408

SI Figure 6: Bacterial abundance (cells mL⁻¹), chlorophyll *a* (μ g L⁻¹), picophytoplankton abundance (cells mL⁻¹), nanophytoplankton abundance (cells mL⁻¹), and POC (mg C m⁻³) for all seven stations (horizontal rows). Sampling periods are blue (late winter) and orange (spring). Each y-axis is in units of depth (m). Note the different scales in both x and y axes. Data from Marquardt 2022a, 2022b, Müller 2023a, 2023b, and Vader 2022.



1415

1416 SI Figure 7: Relative intensities of dissolved organic sulfur compounds (CHOS) at selected 1417 stations on our transect in the northern Barents Sea for late winter (blue) and spring (orange). The 1418 y-axis is in units of depth and the x-axis is relative intensity. Note the different scales in both x 1419 and y axes.



SI Figure 8: Satellite PAR (E m⁻² d⁻¹) during the late winter and spring sampling periods (April 2021). PAR in the transect east of Svalbard (in red box), estimated to be approximately 15 E m⁻² d⁻¹ for all stations, despite significant ice coverage. Retrieved from GlobColour, Hermes, ocean surface optical PAR, © ACRI-ST – GlobColour (https://hermes.acri.fr/index.php?class=archive).



SI Figure 9: Integrated standing stocks (upper 90 m) of major protist groups (g C m⁻²) during the
late winter and spring sampling periods in the northern Barents Sea. The colors illustrate diatoms
(dark green), dinoflagellates (orange), other flagellates (yellow) and ciliates (dark red). The x-axis
shows each individual station. Data from Assmy 2022a, 2022b.



1435 SI Figure 10: Raw beam transmittance (%), the fraction of light that reaches the detector, along 1436 the shelf-basin transect in the northern Barents Sea during late winter (top) and spring (bottom). A 1437 lower beam transmittance can indicate greater scattering or absorption by particles. The x-axis 1438 displays latitude and the nonlinear y-axis is in units of depth (m). Transect created with Ocean 1439 Data View (Schlitzer 2021).

Table S1: Average (\pm one standard deviation) of THg and MeHg concentrations on both late1442winter and spring cruises for water column samples above 500 m, categorized by water masses;1443Atlantic Water (AW), modified Atlantic Water (mAW), Polar Water (PW), warm Polar Water1444(wPW). Significant differences (log-transformed, Student's t-test, p < 0.05) between THg and</td>1445MeHg in each respective season when categorized by water mass have each seasonal average1446marked with an asterisk (*).

	Total Hg	(pmol L ⁻¹)	Methylated]	Hg (fmol L ⁻¹)
Water mass	Late winter	Spring	Late winter	Spring
AW	0.47 ± 0.01 (n = 2)	N/A	200 ± 76 (n = 2)	N/A
mAW	$0.53 \pm 0.04*$	$0.64 \pm 0.09*$	63 ± 19	57 ± 77
	(n = 7)	(n = 10)	(n = 7)	(n = 6)
PW	$0.55 \pm 0.07*$	$0.66 \pm 0.08*$	$72 \pm 30*$	$25 \pm 14*$
	(n = 26)	(n = 34)	(n = 26)	(n = 31)
wPW	0.52 ± 0.07	0.56 ± 0.07	105 ± 37	75 ± 36
	(n = 14)	(n = 12)	(n = 14)	(n = 12)

1448 Table S2: Spearman rank correlation analysis (95% CI) for select environmental parameters for all stations and depths less than 500 m

1449 on the late winter cruise, with depth, total mercury (THg) methylated mercury (MeHg) particulate organic carbon (POC), chlorophyll a

1450 (Chla), nitrite (NO₂), nitrate (NO₃), phosphate (PO₄), silicate (SiOH₄), conservative temperature (Θ), absolute salinity (S_A), and bacteria.

1451 A significant p-value (p < 0.05) marked in bold.

	Depth	THg	MeHg	POC	Chla	NO ₂	NO ₃	PO ₄	SiOH ₄	Θ	SA
THg	-0.097										
MeHg	0.364	0.03									
POC	0.037	0.201	0,095								
Chla	-0.545	0.459	-0.117	0,2							
NO_2	-0.077	0.317	-0.315	0.213	0.217						
NO ₃	0.623	-0.053	0.497	0.42	-0.323	0.153					
PO ₄	0.545	-0.025	0.417	0.523	-0.277	0.237	0,957				
SiOH ₄	0.588	0.061	0.565	0.455	-0.204	0.135	0,912	0.867			
Θ	0.63	-0.243	0.475	0.38	-0.429	0	0,889	0.833	0.817		
S_A	0.559	-0.241	0.205	0.294	-0.329	0.082	0,785	0.757	0.654	0.8	
Bacteria	0.105	0.298	-0.418	0.267	0.133	0.618	0,113	0.249	0.037	-0.082	0.077

Table S3: Spearman rank correlation analysis (95% CI) for select environmental parameters for all stations and depths less than 500 m1463on the spring cruise, with depth, total mercury (THg) methylated mercury (MeHg) particulate organic carbon (POC), chlorophyll a1464(Chla), nitrite (NO₂), nitrate (NO₃), phosphate (PO₄), silicate (SiOH₄), conservative temperature (Θ), absolute salinity (S_A), and bacteria.1465A significant p-value (p < 0.05) marked in bold.</td>1466

	Depth	THg	MeHg	POC	Chla	NO_2	NO ₃	PO ₄	SiOH ₄	Θ	SA
THg	-0.609										
MeHg	0.394	-0.42									
POC	-0.622	0.714	-0.582								
Chla	-0.707	0.708	-0.467	0.895							
NO_2	-0.05	0.045	-0.308	0.212	0.251						
NO ₃	0.66	-0.417	0.51	-0.489	-0.534	-0.22					
PO ₄	0.724	-0.484	0.542	-0.59	-0.64	-0.235	0.937				
SiOH ₄	0.668	-0.324	0.438	-0.389	-0.472	-0.162	0.938	0.897			
Θ	0.603	-0.414	0.378	-0.311	-0.315	-0.081	0.802	0.674	0.823		
S_A	0.649	-0.329	0.319	-0.155	-0.255	-0.057	0.723	0.621	0.724	0.856	
Bacteria	-0.249	0.469	-0.389	0.468	0.438	-0.061	0.107	0.021	0.188	0.166	0.249

	Integrated Total Hg pool (pmol m ⁻²)							
0 - 100 m	Late winter (March 2021)	Spring (May 2021)	Summer (August 2019)	Early winter (December 2019)				
P1	51647 ± 1696	68280 ± 1620	64033 ± 869	37050 ± 1736				
P2	51197 ± 1620	67652 ± 1541	63050 ± 865	40050 ± 1800				
Р3	55570 ± 1620	63074 ± 1620	64483 ± 922	40267 ± 1829				
P4	49786 ± 1601	67616 ± 1601	73635 ± 838	51400 ± 1866				
Р5	53409 ± 1677	63737 ± 1541	78773 ± 886	50136 ± 1753				
P6	52700 ± 1994	72807 ± 1743	63300 ± 1115	43955 ± 2049				
P7	52244 ± 2016	63251 ± 1989	55300 ± 1089	58200 ± 2200				

Table S4: Integrated THg pools (pmol m⁻²) from 0 -100 m from all seasonal cruises completed in
the area. We used data from Kohler et al. 2022a for summer and early winter.

1471 Table S5: Integrated MeHg pools (pmol m⁻²) from 0 -100 m from all seasonal cruises completed
1472 in the area. We used data from Kohler et al. 2022 for summer and early winter.

	Integrated Methylated Hg pool (pmol m ⁻²)							
0 - 100 m	Late winter (March 2021)	Spring (May 2021)	Summer (August 2019)	Early winter (December 2019)				
P1	5486 ± 237	710 ± 293	10133 ± 583	9350 ± 648				
P2	3999 ± 227	2187 ± 216	10000 ± 604	9513 ± 731				
P3	6912 ± 227	1341 ± 260	10317 ± 583	9450 ± 555				
P4	7339 ± 224	1378 ± 245	9444 ± 548	9433 ± 583				
P5	6611 ± 235	2640 ± 216	9550 ± 547	11000 ± 547				
P6	8297 ± 279	5878 ± 244	13000 ± 679	10290 ± 624				
P7	13381 ± 282	5485 ± 278	12850 ± 679	14350 ± 679				

1473 **Table S6:** Δ MeHg (pmol m⁻² d⁻¹) for 0-100 m, calculated from 0 – 100 m integrated pools for late 1474 winter and spring. A negative value assigned in front of each Δ MeHg indicates in a negative 1475 change in the MeHg pool from late winter to spring, or a loss of MeHg.

Station	Days between	ΔMeHg	ΔMeHg _{pd}	ΔMeHg _{scav}	ΔMeHgev
	sampning	pmorm u	pmorm u	pmorm u	pmorm u
P1	56	-85 ± 7	-10	-1	-7
P2	56	-32 ± 6	-3	-1	-2
P3	57	-98 ± 6	-3	-1	-2
P4	57	-105 ± 6	-1	-1	-1
P5	56	-71 ± 6	-1	-1	0
P6	58	-42 ± 6	-3	-1	-2
P7	58	-136 ± 7	-3	-1	-2

1476 Table S7: Percent of variance explained for the first two plotted principal components for each1477 cruise.

	% of Variance			
Cruise	PC 1	PC 2		
Late Winter 2021 (LW)	43.3	20.2		
Spring 2021 (SP)	52.8	17.0		
Summer 2019 (SU)	56.5	12.1		
Winter 2019 (W)	46.8	16.2		

1478