1	<u>Contrasting spatial and seasonal trends of methylmercury exposure</u>
2	pathways of Arctic seabirds: combination of large-scale tracking and
3	stable isotopic approaches
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29 Abstract

30 Despite the limited direct anthropogenic mercury (Hg) inputs in the circumpolar Arctic, 31 elevated concentrations of methylmercury (MeHg) are accumulated in Arctic marine 32 biota. However, the MeHg production and bioaccumulation pathways in these ecosystems 33 are not completely unravelled. We measured Hg concentrations and stable isotope ratios 34 of Hg, carbon and nitrogen in feathers and blood of geolocator-tracked little auk Alle alle 35 from five Arctic breeding colonies. The wide-range spatial mobility and tissue-specific 36 Hg integration times of this planktivorous seabird allowed the exploration of their spatial 37 (wintering quarters/breeding grounds) and seasonal (non-breeding/breeding periods) 38 MeHg exposure. An east-to-west increase of head feather Hg concentrations (1.74-3.48 $\mu g \cdot g^{-1}$) was accompanied by significant spatial trends of Hg isotope (particularly Δ^{199} Hg: 39 0.96 to 1.13‰) and carbon isotope (δ^{13} C: -20.6 to -19.4‰) ratios. These trends suggest 40 41 distinct mixing/proportion of MeHg sources between western North Atlantic and eastern Arctic regions. Higher Δ^{199} Hg values (+0.4‰) in northern colonies indicate an 42 43 accumulation of more photochemically impacted MeHg, supporting shallow MeHg 44 production and bioaccumulation in High Arctic waters. The combination of seabird tissue 45 isotopic analysis and spatial-tracking helps tracing the MeHg sources at various spatio-46 temporal scales.

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51 **TOCArt- graphical abstract**



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54 **1 Introduction**

55 Mercury (Hg) induces major risks for wildlife and human health, especially under its methylated form (methylmercury, MeHg), a potent bioaccumulative neurotoxin¹, which 56 57 is mainly assimilated via fish and seafood consumption. In the ocean, MeHg production mainly occurs by biotic in situ methylation of inorganic Hg^{2,3}. Once formed, MeHg 58 59 incorporates into the food webs and biomagnifies its concentrations leading to lifeimpacting levels in top predators and humans. Despite little direct anthropogenic pressure 60 61 in the Arctic region, Arctic ecosystems are subject to contamination by Hg transported 62 from lower latitudes. Indeed, total Hg concentrations measured in the Arctic surface seawater are up to 2-fold higher compared to other oceanic regions ^{4,5}. Sea-ice melting, 63 64 direct atmospheric deposition and continental inputs originating from soil erosion and riverine circulation are considered major drivers of the high Hg levels in the Arctic ^{6–10}. 65 66 However, the MeHg production pathways and zones in the Arctic Ocean are still not 67 completely identified. Several studies demonstrated that Hg in Arctic marine 68 environments may be methylated in the water column or sediments ^{2,11}. Potential Hg methylating bacteria were also identified in Antarctic sea ice ¹². Recent findings and 69

70 modelling studies evidenced that the largest net MeHg production in Arctic water columns may occur in oxic waters at the subsurface layer (20–200 m) ^{6,13}. A new study 71 72 also reported the high abundance of Hg methylating gens in the oxic subsurface waters 73 of the global ocean ¹⁴, where the highest MeHg concentrations are typically observed ⁴. 74 All these findings suggest that Hg methylation in oxic waters could be a significant source 75 of MeHg towards Arctic marine food webs. Although policy implementations for the 76 reduction of anthropogenic Hg emissions were achieved over the last 30 years in some 77 parts of the world, Hg levels continue to increase in biota from several regions of the Arctic¹⁵. Medium to high predators such as seabirds are exposed to significant 78 environmental MeHg concentrations through their diet ^{15,16} and have been extensively 79 studied as bioindicators of Hg exposure in marine food webs (e.g ^{17,18}), including the 80 Arctic ^{19–21}. Specific foraging habitats and migratory movements of Arctic seabirds 81 82 strongly determine their exposure to distinct environmental MeHg sources in marine ecosystems ^{22,23}. However, studies on Hg exposure in Arctic seabirds have commonly put 83 84 the focus towards the breeding season (summer) when seabirds are more accessible for 85 researchers. Consequently, the investigation of Hg exposure during the non-breeding 86 season is still scarce due to sampling logistical difficulties.

The combination of carbon and nitrogen stable isotopes with Hg stable isotopes has demonstrated its suitability for the identification of Hg sources and the associated geochemical processes in the different marine compartments $^{24-26}$. Therefore, its use can help understanding Hg exposure pathways of seabirds according to their migratory behaviour. Hg has seven stable isotopes (196 to 204) and fractionates dependently and independently of the isotopic masses. The combined use of Hg isotopic mass-dependent (MDF, e.g. δ^{202} Hg) and mass-independent (MIF, e.g. Δ^{199} Hg) fractionation enables the

94 quantification of processes and the identification of sources and pathways of Hg in the environment²⁷, including marine ecosystems^{25,26,28,29}. MDF of Hg isotopes occurs during 95 many physical, chemical or biological processes ^{30–33}. However, large Hg MIF of odd 96 isotopes (Δ^{199} Hg and Δ^{201} Hg) is observed during light-induced reactions, such as 97 98 inorganic Hg photoreduction and MeHg photodemethylation. Hg MIF signature is not affected by biological or trophic processes, so it is preserved up to the food webs ³⁴, then 99 100 presenting a significant advantage to trace Hg marine sources. For instance, Arctic marine 101 top predators reported much higher Hg odd MIF values (more photochemically impacted 102 Hg) in non-ice covered regions, relating the importance of the accelerated melting of sea ice on the Hg polar cycle ^{25,35}. Also, consistent decrease of Hg odd MIF (and MDF) in 103 104 pelagic fish according to their foraging depth in the North Pacific Ocean demonstrated 105 the dilution of surface MeHg by *in situ* methylated Hg at depth³. More recently 106 discovered, Hg MIF of even Hg isotopes (reported as Δ^{200} Hg) seems to occur during complex atmospheric mechanisms such as photo-oxidation in the tropopause ³⁶. Even-107 108 MIF is not induced during any biogeochemical nor photochemical processes in the lower troposphere or the photic zone ^{36–38}, therefore the signature is preserved and useful to 109 identify major potential Hg sources of atmospheric origin ^{10,39,40}. Due to the different 110 111 combinations of the processes involving Hg transformations in the environment, Hg 112 isotopes fractionate differently and with different degrees of magnitude in every specific 113 environmental compartment. Thus, the analysis of Hg stable isotopes of mobile predators 114 such as Arctic seabirds can give access to interesting information about MeHg trophic 115 sources at large scales of the Arctic Ocean and neighbouring water bodies. 116 Here we propose an original approach consisting in the combination of isotopic analyses

117 (Hg, C and N) and wildlife tracking to provide new information about MeHg exposure

118 pathways of seabirds at both temporal and spatial scales. For this purpose, we focused on 119 the little auk (or dovekie, Alle alle), the most numerous seabird species breeding in the 120 High Arctic (between 37 to 40 million breeding pairs estimated ^{41,42}). Little auks have 121 several ecological advantages for their use as a bioindicator models. 1) They are 122 zooplanktivorous and mainly feed on copepods belonging to two *Calanus* species (i.e., C. glacialis and C. hyperboreus) during the breeding period ⁴³. Therefore, they reflect 123 124 MeHg accumulation in a short food chain that is strongly dependent on sea ice abundance 125 and seawater temperature ⁴⁴. 2) They exhibit colony specific wintering areas ⁴⁵, then reflecting wide-ranging spatial variability of Hg exposure ⁴⁶. 3) Little auks moult their 126 127 feathers twice during their annual cycle: a partial moult (head, neck and throat feathers, 128 hereafter 'head feathers') during the pre-breeding period (in ca. April) and a complete post-breeding moult in September¹³. During moult, seabirds excrete the Hg accumulated 129 130 in their body tissues into feathers ⁴⁷. Thus, feather Hg reflects blood Hg levels at the time of feather growth which occurred at the last moulting sequence ⁴⁸, then integrating Hg 131 132 from current diet and/or the remobilization of Hg from tissues during moult. Thereby, the 133 different Hg integration times between the types of feathers allow studying Hg exposure 134 during both the non-breeding (head feathers) and the breeding (body feathers) periods in a same individual ⁴⁶. Besides, C and N isotopic ratios of blood sampled during the 135 136 breeding – chick rearing – period can provide information about summer diet and then be 137 compared to Hg levels and isotopic composition in body feathers of little auks ^{49,50}. We 138 hypothesized that variations of tissue-specific Hg isotopic signatures (body vs head 139 feathers) will allow reflecting the seasonal variability (summer vs winter) on the Hg 140 cycling. Besides, the exploration of both spatial grounds and isotopic information (Hg, C

and N) would help tracking distinct sources of Hg contamination along with seabirdmigratory circulation.

143 2 Material and methods

144 2.1 <u>Sampling sites and description of sample collection</u>

145 This study was conducted during the seabird breeding seasons of 2015 and 2016 at five 146 colonies of the Arctic Ocean: Franz Josef Land (FJL) (Hooker Island; 80.23°N, 53.01° 147 E), Bear Island (Bjørnøya; 74.45°N, 19.04° E), East Greenland (Kap Høegh; 70.72°N, 148 21.55°W), Spitsbergen (Hornsund; 76.97°N, 15.78 °E) and North West Greenland 149 (Thule; 77.47°N, 69.22° W). Blood and feathers were sampled from ten individuals per 150 colony (n=50 for the 5 colonies). Individuals from all sites, but Thule, were equipped with 151 a miniature geolocator data-logger (GLS, Biotrack MK4083 or Migrate Technology C65) 152 to track their non-breeding movements and distribution, as described in previous works ^{45,51,52}. We treated GLS tracking data from 1st December to 30th January (period when all 153 154 little auks were at their winter grounds) and calculated the median individual winter 155 latitude and longitude for each individual separately.

156 2.2 Description of analytical methods

157 Sample preparation, analyses of total Hg and Hg species concentrations

Body and head feathers were cleaned in a 2:1 chloroform:methanol solution for 5 min in an ultrasonic bath, followed by two methanol rinses to remove surface impurities, and then oven dried at 50 °C during 48 h and homogenized to powder ⁴⁶. Since fluctuations of Hg concentrations have been observed among and within individual feathers from the same bird ^{53,54}, we pooled a representative number of feathers of each individual (5-8 163 feathers) to limit the variability and provide results as accurately as possible. Blood 164 samples were dried, lyophilized and ground to powder as described in a previous work ⁴⁶. 165 Feather and blood total Hg concentration (hereafter expressed as $\mu g \cdot g^{-1}$, dry weight) was 166 quantified by using an advanced Hg analyser (AMA-254, Altec).

167 Prior to Hg speciation analyses, blood and feathers (0.01–0.05 g) were digested following 168 a previously developed method by microwave-assisted extraction (using a Discover SP-D microwave, CEM Corporation) 55,56. We used 5 mL of tetramethylammonium 169 170 hydroxide (25% TMAH in H2O, Sigma Aldrich) for blood samples and 5 mL nitric acid 171 (HNO₃·6N, INSTRA quality) for feather Hg extraction. The extraction was carried out 172 in CEM Pyrex vessels by 1 min of warming up to 75 °C and 3 min at 75 °C with magnetic 173 agitation to homogenise the samples. Quantification of Hg species was carried out by isotope dilution analysis (details in ⁵⁵), using a GC-ICP-MS Trace Ultra GC equipped 174 175 with a Triplus RSH autosampler coupled to an ICP-MS XSeries II (Thermo Scientific, 176 USA). We performed Hg speciation analyses of certified reference materials (CRM) for 177 QA/QC purposes (Table S1). Human hair reference material (NIES-13) and feather 178 internal reference material (F-KP, king penguin feathers) were used for validation of 179 feather analyses (keratin-based matrixes). Blood analyses were validated with dogfish 180 liver reference material (Dolt-5) and with internal reference material (RBC-KP, king 181 penguins red blood cells). The reported results total Hg concentrations obtained by Hg 182 speciation analyses (i.e., the sum of inorganic and organic Hg) were compared to total Hg 183 concentrations obtained by AMA-254 to verify the recovery of the extraction. Recoveries 184 of Hg and MeHg concentrations with respect to the reference values for each material 185 varied between 92 and 108% (Table S1).

186 Total Hg isotope analyses

187 Feather (and blood) samples (0.05–0.10 g) were digested with 3 or 5 mL of HNO₃ acid 188 (65%, INSTRA quality) after a predigestion step overnight at room temperature. Hg 189 extraction was carried out by Hotblock heating at 75 °C during 8 h (6 h in HNO₃ and 2 h 190 more after addition of 1/3 of the total volume of H₂O₂ 30%, ULTREX quality). The digest 191 mixtures were finally diluted in an acidic matrix (10% HNO₃ and 2% HCl) with final Hg 192 concentrations ranging from 0.5 to 1 ng \cdot mL⁻¹. Hg isotopic composition was determined 193 using cold-vapor generator (CVG)-MC-ICPMS (Nu Instruments), detailed in previous 194 work ⁵⁶. Hg isotopic values were reported as delta notation, calculated relative to the 195 bracketing standard NIST SRM-3133 reference material to allow interlaboratory 196 comparisons, as described in the SI. NIST SRM-997 thallium standard solution was used 197 for the instrumental mass-bias correction using the exponential law. Secondary standard 198 NIST RM-8160 (previously UM-Almadén standard) was used for validation of the 199 analytical session (Table S2).

200 Recoveries of extraction were verified for all samples by checking the signal intensity 201 obtained on the MC-ICPMS for diluted extracts relative to NIST 3133 standard (with an 202 approximate uncertainty of $\pm 15\%$). Total Hg concentrations in the extract solution were 203 compared to the concentrations found by AMA-254 analyses to assess method recovery. 204 Total Hg concentrations in the extract solution were compared to the concentrations found 205 by AMA-254 analyses to assess method recovery. Average recoveries obtained were 98 206 \pm 14% for feathers (n = 104) and 100 \pm 2% for blood samples (n = 102). Accuracy of Hg 207 isotopic analyses for keratin matrixes was evaluated with validated human hair material 208 NIES-13 isotopic composition ⁵⁷. Hg isotopic results for blood samples were validated 209 with reference values of Lake Michigan fish tissue NIST SRM 1947. Internal reference

samples of feathers (F-KP) and avian blood (RBC-KP) were also measured. Uncertainty
for delta values was calculated using 2SD typical errors for each internal reference
material (Table S2).

213 Carbon and nitrogen stable isotope analyses

214 Homogenized feather and blood subsamples (aliquots mass: ~ 0.3 mg) were weighed with a microbalance and packed in tin containers. Carbon (δ^{13} C) and nitrogen (δ^{15} N) stable 215 216 isotope ratios were determined with a continuous flow mass spectrometer (Thermo 217 Scientific Delta V Advantage) coupled to an elemental analyser (Thermo Scientific Flash 218 EA 1112). Results are in delta notation relative to Vienna PeeDee Belemnite and atmospheric N₂ for δ^{13} C and δ^{15} N, respectively. Replicate measurements of internal 219 220 laboratory standards (acetanilide) indicated measurement errors < 0.15% for both δ^{13} C 221 and δ^{15} N values. USGS-61 and USGS-62 reference materials were also analysed for 222 calibration.

223 2.3 <u>Statistical analyses</u>

224 Statistical analyses were performed using the software R 3.3.2 (R Core Team, 2018)⁵⁸. 225 Before statistical analyses, the data were checked for normality of distribution and 226 homogeneity of variances using Shapiro-Wilk and Breusch-Pagan tests, respectively. 227 Since data did not meet specificities of normality and homoscedasticity, non-parametrical 228 tests (Kruskal–Wallis with Conover-Iman *post-hoc* test) were performed. Statistically 229 significant results were set at $\alpha = 0.05$. Statistical significance of Hg concentration and 230 isotopic differences between head and body feathers were assessed using a randomization 231 procedure. A 99% confidence interval was calculated by means of bootstrap estimation 232 method (n=1000 iterations).

We examined the correlations between Hg concentrations, δ^{13} C, Hg MDF (δ^{202} Hg) and 233 MIF (Δ^{199} Hg and Δ^{200} Hg), latitude and longitude using linear regressions and Spearman 234 235 correlation rank tests. The influence of the latitude and longitude of their individual 236 breeding and non-breeding distribution on feather Hg isotopic signatures were tested 237 using linear mixed models (LMMs) with colonies as random effect on the whole data set, using the R package "lme4" ⁵⁹. Summer latitude, summer longitude and both summer 238 239 latitude + longitude together were used as predictors for Hg isotopic signatures of body 240 feathers. Similarly, median winter latitude, median winter longitude and both median 241 together were used as predictors of Hg isotopic signatures in head feathers. Variance inflation factors were always $< 3^{60}$, ensuring that there was not collinearity between 242 243 latitude and longitude in summer (breeding colonies) and median latitude and longitude in winter (wintering areas) ⁶¹. The different models were ranked based on Akaïke's 244 245 Information Criteria adjusted for small sample sizes (AICc) and compared using Δ AICc and Akaike weights (w) using the R package "wiqid" ⁶². To assess the explanative power 246 of these models, marginal R^2 was obtained using the R package "r2glmm" ⁶³. 247

- 248 **3 Results and discussion**
- 249 3.1 <u>Seasonal and geographical variations of feather MeHg concentrations related to</u>
 250 <u>changing foraging habits (δ¹⁵N)</u>

We observed that the dominant fraction of Hg was in the form of MeHg both in body feathers and blood (94 \pm 2%, n=20 and 90 \pm 3%, n=10; respectively) for all the studied populations of little auks (Table S3). This result is in good agreement with previous studies ^{55,56,64,65} and supports that both tissues of little auks principally present Hg as MeHg. Body and head feathers are known to be grown at different times and therefore the Hg excreted in head and body feathers reflects respectively the exposure during their wintering (October to April) and breeding (May to September) periods. Since birds are known to excrete between 70 and 90% of their Hg body burden by feather moult ⁴⁸, we cannot exclude that some residual Hg accumulated during the non-breeding period could also be excreted during body feather moult, and vice versa, but this fraction would be minor.

262 Overall, individuals presented higher Hg (MeHg) concentrations in head compared to 263 body feathers, exhibiting up to 2-fold higher concentrations in head feathers in the case 264 of East Greenland and Bear Island populations (Table S4). Higher Hg concentrations of head feathers are coherent with previous observations ²² and suggest a higher exposure to 265 266 MeHg during the non-breeding period outside the High Arctic. For instance, little auks 267 breeding in areas of Spitsbergen and East Greenland are known to mainly forage on copepods (*Calanus* spp.) during the breeding season ⁴³. However, the seasonal vertical 268 269 migration of their main prey Calanus spp. to inaccessible depths produces a seasonal shift 270 in their diet towards krill Meganyctiphanes norvegica, hyperiid amphipods Themisto spp., 271 and fish larvae ⁶⁶. The consumption of higher trophic level prey during winter could 272 explain the higher Hg levels excreted during the spring moult (head feathers), whereas 273 they are probably less exposed to Hg in summer.

We also observed high variations of Hg concentrations in head feathers among individuals of the same colony, especially in Bear Island (from 0.81 to 4.35 μ g g⁻¹) and Spitsbergen populations (from 1.67 to 3.79 μ g g⁻¹) (Fig. S1). This could be due to the wide-spread individual foraging specialisation during their non-breeding period and the consumption of a wider range of prey ⁶⁷. Conversely, little auks occupy more restricted foraging areas during the breeding season due to the need to frequently feed their chicks
and therefore feed on local prey captured near their respective colonies ⁶⁸, leading to less
intra-population variability of Hg concentrations in their body feathers.

We observed a consistent longitudinal trend of body feather Hg concentrations ($R^2=0.58$. 282 283 p<0.0001) with increasing Hg levels from eastern (Bear Island and FJL, 0.71 and 0.75 284 $\mu g \cdot g^{-1}$, respectively) to western colonies (NW Greenland, 2.07 $\mu g \cdot g^{-1}$) (Figure 1). When 285 applying mixed models, summer longitude was the most supported predictor of body 286 feather Hg concentrations (Table S6). Head feather Hg concentrations were positively 287 correlated both with winter latitude and longitude for the four spatially tracked 288 populations ($R^2=0.54$ and $R^2=0.60$, respectively; both p<0.0001) (Figure 1). Both 289 variables together were considered as predictors of Hg head feather concentrations by 290 linear mixed models (Table S7). Head feather concentrations were higher in populations 291 wintering in western zones (3.48 $\mu g \cdot g^{-1}$, East Greenland population) and decreased 292 gradually and significantly (H=20.13, p=0.001) in those wintering in northeast areas (1.74 293 $\mu g \cdot g^{-1}$, FJL population). The consistent longitudinal patterns both in summer and winter 294 reflect a higher accumulation of MeHg in little auks from western regions, whereas 295 colonies breeding in Arctic northern regions seem to be exposed to lower concentrations.



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Figure 1. Hg concentration ($\mu g \cdot g^{-1}$) of little auk body feathers (summer) as a function of latitude (A) and longitude of their breeding sites (B) and head feathers (winter) as a function of the median latitude (C) and longitude (D) of their winter grounds. Regression lines are A) Slope: 0.052±0.013, intercept: -3.420±0.997, R²=0.26, p=0.005; B) Slope: -0.012±0.001, intercept: 1.209±0.059, R²=0.58, p<0.0001; C) Slope: -0.059±0.011, intercept: 6.399±0.669, R²=0.54, p<0.0001; D) Slope: -0.022±0.003, intercept: 2.375±0.120, R²=0.60, p<0.0001. Regression lines presented only for significant relationship between the two variables.

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Seabird blood δ^{15} N values provide short- to medium-term information (about 1–5 weeks) while feathers δ^{15} N values reflect the diet at the time they were grown ^{50,69}. The distribution of little auk populations in winter was limited to the North Atlantic and the Arctic areas, where large-scale δ^{15} N values are known to be relatively homogeneous at the base of the food web ^{70,71}, then allowing the inter-population comparison. The lower

310 body feather Hg concentrations and blood δ^{15} N values observed in little auks (Table S5) 311 suggest that all birds from the different populations mostly feed at low trophic levels and 312 on *Calanus* copepods in summer. Contrarily, the interpopulation differences of $\delta^{15}N$ 313 values in winter (head feathers) were much more pronounced (Table S4). For instance, 314 little auk populations breeding in FJL and Bear Island exhibited a ~ 3 ‰ higher δ^{15} N 315 values in head feathers than in blood. This difference highlights the spatial variability of 316 δ^{15} N values in relation to the different winter distribution of little auks in winter. Previous 317 studies have reported significant seasonal variations in copepod δ^{15} N values (up to 6‰) 318 between late winter and spring (highly productive periods) relative to the summer and autumn periods ^{71,72}. The seasonal variability of zooplankton δ^{15} N values is common on 319 320 the eastern and western parts of the North Atlantic Ocean and needs to be considered here 321 due to the wide spatial distribution of little auks in winter. Therefore, the higher feather 322 Hg concentrations little auk colonies from western parts of the Arctic Ocean could be 323 influenced by their seasonal dietary shifts and different spatial distribution but also by the 324 complex Hg oceanic dynamics or distinct environmental sources that control the level of 325 exposure to MeHg at the different regions.

326 3.2 Spatio-temporal trends of Hg MDF (δ^{202} Hg) in feathers related to ecological aspects

327 Specific Hg integration times of seabird tissues may influence the seasonal incorporation 328 of MeHg from different spatial origin ⁷³. However, the geographical variations in δ^{202} Hg 329 values are generally difficult to distinguish since metabolic processes also induce Hg 330 MDF. Head and body feathers showed large ranges of δ^{202} Hg values, varying from -0.24 331 to 1.43 ‰ and from -0.11 to 1.28 ‰, respectively. Although we focused on the study of 332 multiple colonies of the same seabird species to minimize the metabolic or trophic-related 333 effects, we cannot exclude that the variability of δ^{202} Hg signatures among colonies is led 334 only by the specific isotopic baseline of their respective foraging habitats. For instance, FJL population exhibited significantly heavier δ^{202} Hg values relative to the other four 335 336 populations, both in head (H=29.42, p<0.0001) and body feathers (H=27.69, p<0.0001) 337 (Table S4). It is known that little auks from FJL are morphologically bigger than those of the populations from Svalbard due to more severe climate conditions in this area ⁷⁴. Thus, 338 339 potentially different morphological characteristics associated to their bigger size could 340 contribute to higher feather δ^{202} Hg values in this colony. Hg concentrations and δ^{202} Hg 341 values of head feathers were highly correlated for the overall populations ($R^2=0.52$, p<0.0001), while Δ^{199} Hg signatures were not related to Hg concentration in any type of 342 343 feather (Figure S2). This observation shows the completely decoupled behavior between δ^{202} Hg and Δ^{199} Hg signatures. The influence of biological and ecological factors on 344 345 δ^{202} Hg values shows the limitation of this type of signature to discern spatial MeHg 346 sources related to different migratory routes of seabird populations. The utilisation of feather δ^{202} Hg values as a proxy of geographical patterns or to changing environmental 347 348 conditions requires a complete knowledge of all the processes and factors driving Hg 349 MDF (i.e., trophic ecology and intrinsic metabolic/physiological processes).

350 3.3 <u>Hg odd-MIF (Δ¹⁹⁹Hg): seasonal and spatial differences of Hg marine</u> 351 <u>photochemistry</u>

Head and body feather odd-MIF values (Δ^{199} Hg) ranged from 0.72 to 1.26 ‰ and 0.90 to 1.91 ‰, respectively. Significantly higher Δ^{199} Hg values in body compared to head feathers (Table S8) suggest a seasonal variability in odd-MIF values. This could be primarily associated to the vertical migration of little auk main prey (copepods) and the consequent seasonal shift on their diet to krill/amphipods during the winter season. Their seasonal diet shift could enhance the accumulation of pelagic MeHg that is less connected

to the photic zone during winter, then leading to lower Δ^{199} Hg values of the MeHg 358 359 excreted into head feathers. A previous study on subantarctic penguins documented 360 significant differences of Δ^{199} Hg values as a function of their specific foraging depths ²⁴, 361 increasing around 0.4 ‰ from benthic to epipelagic populations. Although the little auk 362 populations studied here are known to mainly feed on the same prey items and forage at 363 similar depths, we should consider that changes on the availability of their prey among sites could also contribute to different feather Δ^{199} Hg values among populations. Further, 364 365 due to the diurnal migration of zooplankton from deep water to the surface, the mixed 366 pool of Hg accumulated in these organisms originate from different depths of the water column and therefore, their Δ^{199} Hg values represent a mixture from deep (low 367 photodemethylated) Hg and surface Hg uptake ⁷⁵. Together with the trophic and 368 369 ecological factors, we could expect that the seasonal variability of feather Δ^{199} Hg values 370 (body vs head feathers) of little auks could be also influenced by a higher extent of Hg 371 marine photochemistry occurring during summer. In summertime/ spring, little auks are 372 known to return to their breeding sites, located at northern latitudes, where they are 373 exposed to longer daily photoperiod at this moment of the year (polar day). Nevertheless, 374 the weak differences of Δ^{199} Hg values between body and head feathers of a same 375 population (from 0.26 to 0.50 ‰, Table S8) seem to indicate low variations of MeHg 376 photodemethylation extents between their summer and winter sites. Therefore, the 377 differences on daily photoperiod and/or light penetration between their summer and 378 wintering foraging grounds would have a minor role on Hg isotopic variations.

379 Concerning the spatial variability of Δ^{199} Hg values among colonies, we observed positive 380 linear relationships between body feather Δ^{199} Hg values and summer longitude (R²=0.20, 381 p<0.0001) and between head feather Δ^{199} Hg values and winter longitude (R²=0.22, 382 p<0.0001) (Figure 2). Summer and winter longitudes were respectively the most supported explanatory factors of body and head feather Δ^{199} Hg values (Tables S6 and 383 384 S7). No significant relationships were observed with latitude in summer ($R^2=0.01$, 385 p=0.20) nor winter (R²=0.07, p=0.06) (Figure 2). FJL population, the northern colony of this study, showed slightly higher body feather Δ^{199} Hg values (1.64 ± 0.15 ‰, n=10, 386 387 80°N) comparing to other studied colonies $(1.31 \pm 0.20 \text{ }\%, \text{ n}=37 \text{ }70\text{-}77^{\circ}\text{N})$ (H=11.96, p=0.018). FJL individuals also presented higher Δ^{199} Hg values of their head feathers (1.13) 388 389 ± 0.06 ‰) compared to the other colonies (1.00 ± 0.12 ‰) (H=18.55, p=0.001). Previous 390 studies on Alaskan seabirds reported around 2-fold higher mean Δ^{199} Hg signatures in low-391 ice-covered oceanic areas $(1.13 \pm 0.16 \text{ }\%; 56-58^{\circ}\text{N})$ than highly ice-covered regions (0.53 %)392 ± 0.15 %; 68°N) and revealed that the presence of sea ice inhibits light penetration and therefore, Hg marine photochemistry ²⁵. Compared to the latitudinal trend observed in 393 394 Alaska, the spatial variations of Δ^{199} Hg values between northern and southern populations 395 of little auks are relatively weak and, interestingly, presented an inversed tendency 396 between highly ice-covered (FJL) and non-ice-covered areas (North Atlantic regions). 397 Therefore, we cannot presume that the presence of sea ice is a driving factor controlling 398 MeHg photochemistry and the related odd-MIF signatures registered in feathers of little auks. The existence of an opposed north-to-south trend of Δ^{199} Hg values between the 399 400 Eastern and Western Arctic Ocean regions reveals different Hg dynamic systems, 401 especially for Hg accumulation pathways in food webs.

402 According to previous findings, the largest MeHg production in the Arctic water column 403 seems to occur in oxic surface waters just below the productive surface layer ^{6,13}. In the 404 Arctic, additional sources of Hg and carbon are provided by sea ice algae during spring 405 blooms ⁷⁶. The presence of terrestrial organic matter and sea ice layers that concentrates

406 phytoplankton near the MeHg production zone may favour the Hg microbial methylation at shallow depths of the Arctic water column^{12,77}. Shallower methylation occurring in 407 408 Arctic waters may result in higher photochemical impact on MeHg before its assimilation 409 in Arctic biota compared to North Atlantic marine food webs. This phenomenon could 410 contribute to the higher feather Δ^{199} Hg values of FJL little auks compared to populations 411 breeding at lower latitudes. The slight differences of Δ^{199} Hg values between northern and 412 southern colonies of little auks are similar to the ranges recently observed in seabirds 413 covering a wider latitudinal gradient (37 to 66°S) in the Southern Ocean, for which Δ^{199} Hg 414 values increased from Antarctic (1.31 to 1.73 ‰) to subtropical (1.69 to 2.04 ‰) populations ⁷⁸. The slight variations of Δ^{199} Hg values also found between these distant 415 416 sites of the Southern Ocean were translated into low differences of MeHg photochemical 417 demethylation extents among sites and a dominance of MeHg with a mesopelagic origin 418 in these remote environments. As previously discussed, the vertical daily migration of 419 copepods from deep water to the surface leads to the integration of Hg from relatively 420 deep zones of the water column therefore, contributing to the incorporation of low 421 photochemically impacted Hg in consumers ⁷⁵. Therefore, although the planktivourous 422 little auk feed at surface waters on the photic zone, their relatively low feather Δ^{199} Hg 423 values suggest that the Hg accumulated in their main prey could originate from Hg pools 424 from deeper zones of the water column.

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427 Figure 2. Hg odd-MIF (Δ^{199} Hg) of little auk body feathers (summer) as a function of latitude (A) and 428 longitude of their breeding sites (B) and head feathers (winter) as a function of the median latitude 429 (C) and longitude (D) of their wintering grounds. Regression lines are A) Slope: 0.014±0.011, 430 intercept: 0.319±0.822; R²=0.01, p=0.20; B) Slope: 0.002±0.001, intercept: 1.384±0.031, R²=0.20, 431 p<0.0001; C) Slope: 0.004±0.002, intercept: 0.784±0.123, R²=0.07, p=0.06; D) Slope: 0.001±0.001, 432 intercept: 1.069±0.022, R²=0.22, p=0.002. Regression lines presented only for significant relationship 433 between the two variables.

434

435 3.4 Spatial correlation of Hg MIF signatures and carbon stable isotopes (δ^{13} C)

The deposition of atmospheric Hg from mid-latitude anthropogenic emissions into the Arctic Ocean could contribute to the accumulation of MeHg from distinct origin in Arctic-North Atlantic food webs ⁷. Although body feathers of little auk presented a relative high range of Δ^{200} Hg signatures (from -0.23 to 0.17 ‰), the inter-population differences were not significant (H=3.685, p=0.45) (Figure S5). No substantial interpopulation variations 441 of Δ^{200} Hg values were neither observed for head feathers of little auks (-0.14 to 0.12 ‰), 442 therefore we cannot discriminate Hg sources from distinct atmospheric origin among 443 seabird wintering grounds. Nevertheless, the spatial trends of Δ^{200} Hg values observed in 444 little auks are more variable than those previously reported on Arctic marine mammals 445 and seabirds of Alaska (from -0.01 to 0.10 ‰; 71 to 54 °N ⁷⁹) and on Antarctic and 446 subtropical seabirds (from -0.02 to 0.04 ‰, 66 to 37 °S ⁷⁸).

447 Large-scale ocean circulation and vertical transport processes throughout the water 448 column could influence the distribution of distinct MeHg sources between the widely 449 distributed compartments used by little auks. The exploration of carbon stable isotopes 450 $(\delta^{13}C)$ of little auks could help discriminating the potential contributions of distinct MeHg 451 sources linked to the widely specific foraging habitats of little auks. Contrary to Hg 452 isotopes, body feather δ^{13} C values do not reflect the period of summer but the moult period in late summer/early autumn (September) when they are grown ⁴⁵. To ensure only 453 454 the integration of the summer, breeding period, we compared body feather Hg isotopes 455 with blood δ^{13} C values. Little auks from FJL exhibited the lowest blood δ^{13} C values (-456 23.13 ± 0.84 ‰) and NW Greenland individuals the highest (-20.07 ± 0.35 ‰) relative to the rest of the colonies (H=40.74, p<0.0001) (Table S5). Head feather δ^{13} C values 457 458 separated little auk populations in those overwintering in western areas of the North 459 Atlantic Ocean and those wintering in north-eastern areas (H=26.28, p<0.0001). The gradient of δ^{13} C values of head feathers increased from populations of FJL (-20.59 ± 0.40 460 461 ‰) and Bear Island (-20.35 \pm 0.38 ‰), to Northwest (-19.41 \pm 0.57 ‰) and East Greenland (-19.61 \pm 0.46 ‰) populations. Latitudinal gradients of δ^{13} C values of the 462 463 dissolved inorganic carbon are commonly observed in surface waters as an influence of the physical and biological processes ⁷¹. For instance, it is known that CO₂ solubility is 464

465 favoured in cold oceanic waters and consequently, surface waters at high latitudes have relatively low δ^{13} C values due to the introduction of isotopically light atmospheric CO₂. 466 467 By contrast, surface waters of outgassing upwelling equatorial areas become enriched on δ^{13} C values ^{80,81}. Parallelly, the δ^{13} C values of primary producers are strongly influenced 468 469 by the δ^{13} C values of dissolved inorganic carbon and therefore, by the temperature 470 gradients and CO₂ solubility ⁷¹. Spatial gradients of sea surface temperature and CO₂ concentrations could thus explain the more depleted $\delta^{13}C$ baseline in cold high Arctic 471 472 marine food webs and the enrichment in δ^{13} C values when going southward to North 473 Atlantic oceanic areas. Furthermore, the dominance of distinct marine currents between the different wintering seabird sites could strongly determine the $\delta^{13}C$ at the base of the 474 475 food webs. The FJL archipelago and surrounding high Arctic areas are strongly impacted 476 by the Makarov and Arctic cold currents flowing southward from the north, and 477 potentially contributing to transport isotopically depleted carbon from high latitude areas. 478 In contrast, little auk wintering regions near the Newfoundland Island (East and West 479 Greenland populations) are affected by the Gulf Stream and North Atlantic Current ⁴⁵ 480 which could supply carbon organic matter from warmer water masses ⁸².

Significant negative linear relationships were obtained between Δ^{199} Hg and δ^{13} C values 481 both in summer (R²=0.17, p=0.003) and winter (R²=0.31, p<0.0001) (Figure 3). 482 Interestingly, the negative relationship between Δ^{199} Hg and δ^{13} C values of little auks 483 484 contrasts with those previously reported on eggs from guillemot species (or murres, Uria aalge and U. lomvia) breeding in the Alaskan Arctic²⁶. These authors reported a co-485 enrichment of egg δ^{13} C and Δ^{199} Hg values linked to the transition from terrestrial to 486 487 marine Hg sources and the subsequent reduction of Hg photochemistry in coastal reservoirs due to higher turbidity ²⁶. However, the wintering areas of little auks mainly 488

489 correspond to more opened oceanic areas as the study in the Bering Sea and probably do 490 not present such a remarkable coastal-oceanic gradient. The significant correlation obtained here between Δ^{199} Hg and δ^{13} C signatures both in body and head feathers of little 491 492 auks reflect common spatial trends summer and winter foraging grounds. This interesting 493 relationship seems be associated to both the spatial gradient of physical parameters 494 controlling C isotopic baselines (temperature and CO₂ exchange in surface waters) and to 495 the extent of Hg photochemical processes. Probably, a higher stratification and impact of 496 sea ice cover in high Arctic oceanic zones favours the methylation of Hg in surface waters ¹³, and the extent of photochemical reactions leading to slightly positive Δ^{199} Hg values 497 498 and more negative δ^{13} C values of biota. The dominance of northern marine currents in this area would also contribute to depleted δ^{13} C values. Although we could consider the 499 500 existence of distinct carbon inputs transported by the marine currents on these ecosystems 501 (i.e. external carbon supply, planktonic production), the complex interaction of 502 oceanographic and physical parameters governing these areas does not allow to provide 503 conclusive evidence from our data.

504

505 Figure 3. Carbon (δ^{13} C) vs MIF Hg signatures for A) summer (body feathers) and B) winter (head 506 feathers) periods. Regression lines are A) Slope: -0.122±0.026, intercept: -1.045±0.558, R²=0.31, 507 p<0.0001; B) Slope: -0.088±0.028, intercept: -0.746±0.566, R²=0.17, p=0.003.

508 3.5 <u>Geographically distinct Hg source mixing across the Arctic and North Atlantic</u> 509 <u>Oceans</u>

510 Our results suggest that the variations in relation to longitude of the Hg concentrations,

- 511 Δ^{199} Hg, and δ^{13} C values of little auks is linked to the assimilation of isotopically distinct
- 512 MeHg depending on their wintering grounds. Figure 4 shows a compilation of Hg odd-

513 MIF values observed in little auks compared to previous studies in Arctic biota over a 514 wide spatial scale. The observed isotopic spatial variability across the different regions of 515 the Arctic Ocean suggests the existence of two different Hg systems between East (Atlantic) and West (Pacific) Arctic Ocean regions. Our opposed trend of Δ^{199} Hg from 516 north to south populations values relative to Western Arctic compartments ^{25,35} indicates 517 518 that the presence of sea ice cover is not the only driving factor controlling Hg 519 photochemistry in the Eastern Arctic Ocean. Possibly, an additional supply of Hg and 520 carbon sources by sea-ice algae may enhance the microbial/photochemical methylation 521 and demethylation processes at shallower depths ^{6,13} in East Arctic regions, therefore 522 contributing to the higher odd-MIF values of Hg accumulated in biota. The inversed relationship of Δ^{199} Hg and Δ^{200} Hg values of little auks with latitude is opposed to the 523 latitudinal covariation of Δ^{199} Hg and Δ^{200} Hg in biota from Western Arctic regions ⁷⁹ and 524 525 from Antarctic regions ⁷⁸ and evidences a completely different functioning of Hg cycling 526 compared to other polar marine environments. Complex Hg dynamics and ocean control 527 factors seem to drive the increasing pattern of Hg isotopes from west to east regions of 528 the Arctic Ocean. Future research assessing large scale and long-term Hg contamination 529 are necessary to have a complete understanding of the Hg exposure pathways and of the associated risks for the whole marine Arctic environments. 530

531

533 Figure 4. Compilation of Hg odd-MIF values (Δ^{199} Hg) of marine biota from spatially distant Arctic 534 Regions. The map comprises both little auk breeding sites (orange), individual little auk median 535 winter positions (yellow) and previous published data (green) including seabirds ^{25,26}, beluga whale, 536 seals and polar bears ⁷⁹ from Alaskan Regions; pilot whales from Faroe Islands ²⁸, and fish from the 537 Labrador Sea ⁸³.

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843