1	Diet influences on growth and mercury concentrations of two salmonid species from lakes
2	in the eastern Canadian Arctic
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22 ABSTRACT

Diet, age, and growth rate influences on fish mercury concentrations were investigated for Arctic 23 24 char (Salvelinus alpinus) and brook trout (Salvelinus fontinalis) in lakes of the eastern Canadian Arctic. We hypothesized that faster-growing fish have lower mercury concentrations because of 25 growth dilution, a process whereby more efficient growth dilutes a fish's mercury burden. Using 26 27 datasets of 55 brook trout and 133 Arctic char, linear regression modelling showed fish age and diet indices were the dominant explanatory variables of muscle mercury concentrations for both 28 29 species. Faster-growing fish (based on length-at-age) fed at a higher trophic position, and as a result, their mercury concentrations were not lower than slower-growing fish. Muscle 30 RNA/DNA ratios were used as a physiological indicator of short-term growth rates (days to 31 weeks). Slower growth of Arctic char, inferred from RNA/DNA ratios, was found in winter 32 versus summer and in polar desert versus tundra lakes, but RNA/DNA ratio was (at best) a weak 33 predictor of fish mercury concentration. Net effects of diet and age on mercury concentration 34 35 were greater than any potential offset by biomass dilution in faster-growing fish. In these resource-poor Arctic lakes, faster growth was associated with feeding at a higher trophic 36 37 position, likely due to greater caloric (and mercury) intake, rather than growth efficiency. 38

Keywords: somatic growth dilution; RNA:DNA; stable isotopes; climate change

40 **Capsule:** Age and diet, but not growth rate, are primary drivers of mercury concentrations in two

41 salmonid fish species in Canadian Arctic lakes.

42 **1. Introduction**

Mercury contamination in fishes is a widespread environmental issue due to global emissions 43 44 and long-range transport that may result in health risks to fishes (Fuchsman et al., 2016; Kidd and Batchelar, 2011) and human populations that consume them (Dellinger et al., 2012). 45 Research since the 1990s has identified key environmental, ecological and biological processes 46 47 controlling mercury concentrations in freshwater fishes including the supply of methylmercury and its uptake into aquatic food webs (Chételat et al., 2018; Pickhardt et al., 2002; Watras et al., 48 1998; Wu et al., 2019), biomagnification (Cabana and Rasmussen, 1994; Lavoie et al., 2013), 49 habitat-specific trophic transfer (Eagles-Smith et al., 2008; Karimi et al., 2016; Power et al., 50 2002), and bioaccumulation with age due to slow elimination rates (Van Walleghem et al., 2007; 51 Van Walleghem et al., 2013). One aspect of mercury bioaccumulation in fishes that remains 52 uncertain (with conflicting findings in the literature and a lack of consensus) is the importance of 53 growth rates in controlling mercury concentrations. 54 55 Bioenergetic modelling indicates that faster growth rates can reduce fish mercury concentrations through somatic growth dilution (SGD), a process where greater biomass 56 production per unit of food (and mercury) consumed results in the dilution of a fish's mercury 57 58 burden (Carroll and Warwick, 2017; Karimi et al., 2016; Li et al., 2018; Trudel and Rasmussen, 2006). Empirical evidence for the importance of growth dilution in controlling fish mercury 59 60 concentrations is more ambiguous. Some studies have concluded that the mercury concentration 61 of a fish's diet is more important than its growth rate (Chumchal et al., 2010; Essington and Houser, 2003; Harris and Bodaly, 1998; Marusczak et al., 2011; Stafford and Haines, 2001; 62 63 Wiener et al., 1990), while other studies have found lower mercury concentrations in fish with 64 faster growth rates (Doyon et al., 1998; Sharma et al., 2008; Simoneau et al., 2005; Verta, 1990;

Wang and Wang, 2012; Ward et al., 2010). Further refinement of theory on growth rate effects in
relation to other drivers of fish mercury concentration is needed to evaluate multiple stressors
currently impacting aquatic ecosystems such as climate change and eutrophication, which may
alter fish growth (Eagles-Smith et al., 2018).

Climate warming is anticipated to improve growth conditions for fishes in Arctic fresh waters 69 70 through warmer temperatures and increased primary production (Reist et al., 2006b). Species with Arctic distributions such as lake trout (Salvelinus namaycush) and Arctic char (Salvelinus 71 *alpinus*) are typically slow-growing and long-lived, which has been reported as a contributing 72 73 factor for higher contaminant concentrations (Evans et al., 2005; Schindler et al., 1995). More efficient growth could reduce mercury concentrations of Arctic fishes through somatic growth 74 dilution (Stern et al., 2012). However, with limited information on growth-mercury dynamics of 75 Arctic fishes and different reported findings (Braaten et al., 2014; Keva et al., 2017; Swanson 76 77 and Kidd, 2010; van der Velden et al., 2013; van der Velden et al., 2012), further research is 78 needed for Arctic ecosystems.

Divergent findings in the literature on the importance of growth rates and SGD may be 79 influenced by different study designs. van der Velden et al. (2012) noted that SGD of mercury 80 81 has been more typically found in studies of juvenile fish compared to those examining older adults. Recent growth rates, such as those measured in studies of juvenile (Ward et al., 2010) or 82 83 small-bodied fish (Wang and Wang, 2018) could provide different information on growth 84 dilution effects than growth based on length-at-age of adult fishes. There may also be seasonal 85 effects of mercury dilution in summer relative to winter (Braaten et al., 2014; Keva et al., 2017). 86 Growth dilution is the result of efficient growth, when more energy is allocated to biomass 87 production for a given intake of energy (Trudel and Rasmussen, 2006; Ward et al., 2010). A

faster growth rate may not lead to lower mercury concentrations if the growth is due to higher
consumption rates or a diet containing more mercury (Lepak et al., 2009; Trudel and Rasmussen,
2006). Field studies on fish often lack detailed information on the mercury content of their prey,
which may hamper the interpretation of growth effects.

This study examined the influence of age, diet and growth rate on mercury concentrations in 92 93 two salmonid fish species, Arctic char and brook trout (Salvelinus fontinalis), in lakes of the eastern Canadian Arctic. Several aspects of the study design provided new insights on the 94 95 relative importance of of growth, age, and diet in those Arctic fishes: (1) mercury concentrations 96 were contrasted between short-lived, fast-growing brook trout and long-lived, slow-growing Arctic char; (2) short-term growth rates (days to weeks) were examined using a physiological 97 indicator, the RNA/DNA ratio of fish muscle (Chícharo and Chícharo, 2008; Ciotti et al., 2010) 98 in addition to long-term growth rates (years) estimated by length-at-age; and (3) lake-specific 99 100 mercury exposure was quantified by measuring methylmercury (MMHg) concentrations near the 101 base of the food web. We used statistical modelling to test the hypothesis that faster-growing fish have lower mercury concentrations after quantifying the influence of age and diet. 102

103 **2. Materials and methods**

104 *2.1. Study Area*

Fishes were investigated in 12 lakes along a latitudinal gradient in the eastern Canadian Arctic
between 2012 and 2014 (Supplemental Table S1, Fig. S1). The lakes are identified by numbers
(lake 1-12) because only a portion have official names. Five subarctic taiga lakes were sampled
in 2012 for brook trout near the community of Kuujjuaraapik-Whapmagoustui in northern
Quebec (55°N), located approximately 200 km south of the treeline boundary. In three of the five
subarctic lakes, one other fish species–the small-bodied ninespine stickleback (*Pungitius*)

pungitius)-was present and consumed by brook trout. Lake-dwelling Arctic char were sampled
in three Arctic tundra lakes near the community of Iqaluit, Nunavut (64°N) in 2013-14 and four
High Arctic polar desert lakes near the community of Resolute Bay, Nunavut (75°N) in 2014.
Arctic char was the only fish species present in the lakes, with the exception of one tundra lake
(near Iqaluit) which also contained nine-spine stickleback.

Cold temperatures in the Canadian Arctic result in short open-water seasons of ~1–4 months 116 117 for lakes, with the majority of the year under ice cover. None of the study lakes were thermally stratified at the time of sampling, and July water temperatures were 10–17°C in the taiga lakes 118 119 and $\leq 10^{\circ}$ C in the tundra and polar desert lakes. The five taiga lakes with brook trout were small in surface area $(0.01-0.39 \text{ km}^2)$ and had shallow mean depths (0.3-1.9 m). The seven lakes 120 containing Arctic char were larger $(0.15-4.5 \text{ km}^2)$ and deeper (3-19 m). All of the lakes were 121 unproductive (oligotrophic) ecosystems with low water concentrations of total phosphorus (1-10)122 μ g/L), although latitudinal gradients in water-column organic matter, measured as dissolved 123 organic carbon (DOC) and chlorophyll, were observed among the ecoregions (Chételat et al., 124 125 2018; Richardson et al., 2020). In particular, the taiga lakes had higher water concentrations of DOC (4.4 \pm 0.7 mg/L) and chlorophyll (1.2 \pm 0.7 μ g/L) than the tundra lakes (DOC: 1.9 \pm 1.0 126 mg/L; chlorophyll: $0.5 \pm 0.04 \,\mu$ g/L) and polar desert lakes (DOC: $1.6 \pm 0.7 \,$ mg/L; chlorophyll: 127 $0.3 \pm 0.1 \,\mu$ g/L). Geographic coordinates, physical characteristics and water chemistry (measured 128 129 at the time of fish sampling) for each of the study lakes are provided in Supplemental Table S1. 130 2.2. Field sampling

Fish were collected from each lake during one sampling period in July or August, with the exception of three tundra lakes where additional fish were collected during winter (February) under ice. Multi-panel gill nets of varying mesh size (1-5 inches) were used to capture fish of

different size classes in taiga and tundra lakes. For polar desert lakes, only large-sized Arctic 134 char were targeted by gill net as part of an annual monitoring program in those lakes (Hudelson 135 et al., 2019). Total sample sizes per study lake ranged from 10–15 for brook trout and 7–33 for 136 Arctic char (Table 1). Captured fish were euthanized on site according to authorized animal care 137 protocols and measured for mass (g) and total length (cm). Three 2 mm or two 4 mm biopsy 138 139 punches were immediately collected from dorsal muscle, stored in cryovials with RNAlater stabilization solution for preservation of nucleic acid content, and frozen on the same day. 140 141 Muscle biopsies were not collected from some fish (e.g., if they were dead in the net upon collection), and no biopsies were obtained from Arctic char in Lake 9. In the laboratory, fish 142 were examined qualitatively for stomach contents (brook trout only) and dissected for otoliths, 143 scales, and a muscle sample. 144

Collections of sediment-dwelling benthic invertebrates (chironomid larvae: Insecta, Diptera) 145 from the study lakes were reported previously (Chételat et al., 2014; Chételat et al., 2018). 146 147 Invertebrate sampling was conducted around the same time as the fish collection for all lakes except polar desert Lake 9 (fish in 2014, invertebrates in 2011). For two other polar desert lakes, 148 invertebrate data from 2011 were added to increase sample size (Supplemental Information, 149 150 Table S2). Invertebrates were collected offshore with an Ekman grab and 500 µm sediment sieve, and occasionally from shoreline areas using a D-framed kicknet. Chironomid larvae were 151 152 used as a proxy for MMHg bioaccumulation near the base of the food web, providing a consistent taxonomic comparison among lakes of food web exposure to MMHg. Previous 153 research on the same study lakes demonstrated that chironomid MeHg concentrations were 154 related to water MeHg concentrations and were a robust indicator of baseline MeHg 155 bioavailability (Chételat et al., 2018). The tundra and polar desert study lakes had a low diversity 156

of macroinvertebrate taxa, the food webs were supported by benthic carbon, and chironomids
were overwhelmingly the dominant invertebrate prey item of Arctic char (Chételat et al., 2010;
Gallagher and Dick, 2010; Gantner et al., 2010; Lescord et al., 2015). The food webs of the
subarctic taiga lakes with brook trout had not been previously characterized, and
macroinvertebrate taxa (amphipods, dragonfly nymphs, caddisflies, mayflies, and march flies)
were collected at the same time as chironomids to measure MMHg concentrations in potential
prey.

164 *2.3. Laboratory analysis*

Fish age was estimated with otoliths using the crack and burn method. Otoliths were not 165 available for a small subset of brook trout and Arctic char, in which case age was estimated from 166 scale annuli (for < 10% of fish). We found that scales slightly underestimated age (by 0–2 years) 167 relative to otoliths for young fish, as reported elsewhere (Stolarski and Hartman, 2008). We used 168 species-specific poisson regressions derived using scale and otolith data from a subset of study 169 170 fish to estimate otolith age from scales of 6 brook trout and 9 Arctic char aged 2–6 years (Supplemental Information). Scales were not used to estimate age for fish older than 6 years. 171 Total mercury concentration (THg, μ g/g) was measured in freeze-dried and homogenized 172 173 muscle by thermal decomposition, gold trapping, and atomic absorption spectrometry at either the National Wildlife Research Centre (Ottawa, Canada; n = 151) or the Canada Centre for 174 175 Inland Waters (Burlington, Canada; n = 45). For the latter samples, analytical method and quality 176 assurance details are provided in Hudelson et al. (2019), and raw data are available from an open data portal (Environment and Climate Change Canada, 2019). For THg measurements conducted 177 178 at the National Wildlife Research Centre, analytical duplicates had a relative standard deviation 179 of $2 \pm 1\%$ (mean \pm SD, n = 17). Recoveries of THg from certified reference materials were $109 \pm$ 180 9 % (n = 37) for TORT-2 (lobster hepatopancreas) and 100 ± 6 % (n = 21) for DOLT-4 (dogfish 181 liver) from the Natural Research Council of Canada. Muscle THg concentrations were presented 182 on a wet weight basis assuming 80% moisture and were assumed to be representative of muscle 183 MMHg concentrations (Lescord et al., 2015).

Methylmercury concentrations (ng/g dry weight) of benthic invertebrates were measured in 184 185 freeze-dried and homogenized material by pretreatment with an alkaline then acidic digestion, extraction in dichloromethane, isolation with sodium thiosulfate and detection by capillary gas 186 chromatography coupled with atomic fluorescence spectrometry. The results of chironomid 187 MMHg measurements were originally reported in Chételat et al. (2014; 2018) and method and 188 QA/QC details are found therein. MMHg concentrations for other macroinvertebrate taxa in the 189 subarctic lakes have not been previously published although the samples were analyzed at the 190 same time as chironomid samples reported in Chételat et al. (2018), and therefore the same 191 method and QA/QC details apply. Lake-specific sample sizes and concentrations of invertebrate 192 193 MMHg are provided in Supplemental Table S2 and S3.

Dried and homogenized fish muscle and chironomid samples were analyzed for carbon and nitrogen stable isotopes on a a DeltaPlus XP Isotope Ratio Mass Spectrometer interfaced by a Conflo II to a Vario EL III elemental analyzer. Stable isotope ratios were expressed in delta notation (δ) as the parts per thousand (∞) deviation from standards of atmospheric N₂ gas for δ^{15} N and Vienna PeeDee Belemnite for δ^{13} C. Muscle C:N ratios (on a molar basis) were calculated using the %C and %N values generated by the elemental analyzer during the isotopic measurements.

201 Muscle plugs preserved in RNAlater (and stored frozen) were measured for RNA and DNA 202 content. The RNA:DNA ratio of muscle is a quantitative physiological proxy for short-term

203	growth rate of fish because RNA content increases during protein synthesis which drives growth
204	(Chícharo and Chícharo, 2008; Ciotti et al., 2010). DNA content is used to correct for variation
205	in muscle cell density. Muscle RNA and DNA of 157 fish were extracted from 5–10 mg of bead-
206	milled muscle using spin column kits (miRNeasy Mini Kit or DNeasy Blood and Tissue Kit,
207	Qiagen, Hilden, Germany), and the yields were measured by spectrophotometry on a NanoDrop
208	(Thermoscientific, Massachusetts, USA). RNA and DNA absorbance peaks were inspected and
209	the absorbance ratios of 260:280 nm and 260:230 nm were measured to assess the quality of
210	extractions. RNA and DNA yields were divided by the sample mass to calculate muscle
211	concentration (μ g/mg fresh weight). Duplicate measurements of muscle samples showed good
212	reproducibility for both RNA (RSD = 9 ± 7 %, $n = 18$) and DNA (RSD = 13 ± 11 %, $n = 20$).
213	2.4. Data Analysis
214	Fish trophic position was calculated using individual $\delta^{15}N$ values of fish and lake-specific
215	δ^{15} N values of benthic invertebrates according to the following equation:
216	Trophic position = 2 + $(\delta^{15}N_{fish} - \delta^{15}N_{chironomid})/TFF$
217	where $\delta^{15}N_{fish}$ and $\delta^{15}N_{chironomid}$ are the $\delta^{15}N$ values for an individual fish and the lake-specific
218	mean for sediment-dwelling chironomid larvae, and TFF is an assumed $\delta^{15}N$ trophic
219	fractionation factor of 3.4 between a predator and its prey (Post, 2002). Lake-specific $\delta^{15}N$ values
220	for chironomids are provided in supplemental Table S2.
221	Long-term growth rates of Arctic char and brook trout were examined in two different ways,
222	using a continous and a categorical variable to characterize the length-at-age patterns. First, fish
223	were pooled across lakes to generate a species-specific length-at-age distribution, which was fit
224	with an exponential curve (Length=a*(1-b ^{Age})) using the dynamic curve function in SigmaPlot
225	11.0 (Systat Software Inc., San Jose, California). A continuous variable for growth rate, termed

226	relative length-at-age (RLAA), was calculated as the difference between total length of
227	individual fish (L_{fish}) and the modelled total length for fish with the same age (L_{age}) determined
228	from the exponential curve (RLAA= L_{fish} - L_{age}). Negative RLAA values indicated that a fish was
229	smaller than the modelled length for a given age, while positive values indicated a fish was
230	larger. For each species, fish were categorized as "slower", "unclassified" or "faster" growing
231	based on the 0–25 th percentiles, 26–74 th percentiles, and 75–100 th percentiles of RLAA values,
232	respectively, to compare growth categories in a similar manner to van der Velden et al. (2012)
233	Note that all Arctic char in the study lakes grew relatively slowly (e.g., median length at age 10
234	was 28 cm), and the term "faster" grower in this context only refers to a comparison with other
235	Arctic char. Data for a one-year old brook trout was excluded from the analysis because it was
236	the only fish in that age class. In addition, the influence of short-term growth rates was examined
237	using the RNA/DNA ratio of fish muscle. Measurements of age or muscle RNA/DNA ratios
238	were not available for all fish (4% and 20% of values missing, respectively).
239	Statistical models were tested with R (http://cran.r-project.org). Analysis of Covariance
240	(ANCOVA, lm function in stats package) tested differences in log-transformed muscle THg
241	concentration between slower- and faster-growing fish categories, after quantifying the effect of
242	age. Separate tests were conducted for Arctic char and brook trout with fish pooled across lakes.
243	Multiple linear regression models were tested for each species to identify the relative importance
244	of explanatory variables (age, diet, growth) in relation to log-transformed muscle THg
245	concentration. Regression models were tested with the lm function, and the significance of
246	individual variables was determined using type III sums of squares (car package). The relative
247	importance of explanatory variables in the regression models was evaluated by an "lmg" index
248	using the relaimpo package, which calculates the average partial r^2 for all possible orderings of

the explanatory variables because of the influence of variable sequence on the partial r^2

250 (Grömping, 2006). The lmg index (refered to here as the independent partial r^2) is equivalent to

the independent contribution of each predictor variable to the variance of the response variable in

a multiple regression model (Chevan and Sutherland, 1991). Comparisons between two groups

253 were made with *t*-tests or Mann-Whitney U tests (if normality or homogeneity of variance were

not satisfied), and Pearson correlations were used to test for associations among explanatoryvariables.

256 **3. Results**

257 *3.1. Characteristics of the fish populations*

Brook trout across the study lakes had lengths of 12.2–36.7 cm and ages of 1–6 years (Table 258 1). In contrast, Arctic char reached larger sizes and were longer-lived than the brook trout, with 259 260 lengths of 13.5–64.5 cm and ages of 3–26 years (Table 1). Both species showed a range of trophic position estimates consistent with some individuals feeding on aquatic invertebrates 261 while others also likely consumed fish. The lake-mean δ^{13} C values of brook trout and Arctic char 262 ranged from -27.2 to -23.3‰ and -23.9 to -19.7‰, respectively, and there was generally little 263 264 variability within a lake. While the benthic food webs supporting Arctic char have been previously characterized in the study lakes, new stomach content data of brook trout indicated 265 they fed on a variety of macroinvertebrates (e.g., amphipods, caddisflies, chironomids, dragonfly 266 nymphs, and snails) (Supplemental Table S4), and the δ^{13} C values of macroinvertebrates were 267 similar to the fish (Table 1, Supplemental Table S3). The muscle C:N ratios (a proxy for lipid 268 content) showed little variation among fish for either brook trout (3.9 ± 0.1) or Arctic char (2.8 ± 0.1) 269 0.1). Muscle THg concentrations of brook trout varied over an order of magnitude from 270 0.036–0.528 µg/g (wet weight). Arctic char THg concentrations varied even more from 271

272	0.030–3.49 μ g/g and 16% of the 133 char had concentrations > 0.5 μ g/g. The MMHg
273	concentrations of benthic chironomids from the brook trout lakes and Arctic char lakes ranged
274	from 32–160 and 4–173 ng/g (dry weight), respectively, (Supplemental Table S2), indicating
275	that MMHg exposure to the base of food webs differed among the study lakes. For the subarctic
276	taiga lakes, MMHg concentrations of other macroinvertebrates were similar to and correlated
277	with MMHg concentrations of chironomids in the same lakes (Supplemental Table S3, Fig. S2).
278	Those measurements justify the use of chironomid MMHg concentrations as a proxy for among-
279	lakes differences in food web MMHg exposure.

Table 1. Mean (range) of total length, age, trophic position (TP), muscle total mercury

concentration (THg, wet weight)), and δ^{13} C values of brook trout from taiga lakes and Arctic

Lake	Fish	Length	Age	ge TP		$\delta^{13}C$
	n	(cm)	(years)		$(\mu g/g w.w.)$	
			Taiga b	brook trout		
Lake 1	11	23.9	3	3.0	0.137	-25.9
		(17.7 - 29.1)	(2-5)	(2.9 - 3.1)	(0.072-0.176)	(-26.425.2)
Lake 2	10	18.6	4	3.2	0.294	-27.2
		(17.4–19.6)	(2-6)	(3.1 - 3.3)	(0.209 - 0.487)	(-27.626.8)
Lake 3	15	27.0	4	4.0	0.115	-23.5
		(14.8–36.6)	(2-6)	(3.7 - 4.4)	(0.042 - 0.248)	(-26.122.2)
Lake 4	15	26.4	3	3.6	0.079	-24.8
		(12.2–36.2)	(2-4)	(3.3 - 3.9)	(0.036 - 0.128)	(-25.724.1)
Lake 5	10	29.4	4	3.6	0.167	-23.3
		(20.2-36.7)	(1-6)	(3.1 - 4.1)	(0.074 - 0.528)	(-24.722.5)
			Tundra .	Arctic char		
Lake 6	28	30.1	8	3.9	0.197	-20.2
		(14.5 - 58.0)	(3–20)	(3.2–4.6)	(0.043 - 1.17)	(-22.017.6)
Lake 7	29	31.9	11	3.2	0.481	-21.3
		(13.9–59.0)	(4–19)	(2.5 - 3.9)	(0.053 - 3.49)	(-25.718.9)
Lake 8	33	25.8	10	3.6	0.124	-19.7
		(13.5–64.5)	(5-26)	(3.1 - 4.5)	(0.030 - 1.21)	(-23.917.3)
			Polar dese	rt Arctic char		
Lake 9	11	50.5	16	4.0	1.50	-23.2
		(45.9–59.0)	(12–25)	(3.8–4.3)	(0.801 - 2.65)	(-23.822.7)
Lake 10	7	42.1	16	2.5	0.230	-23.9
		(39.9–53.2)	(15–17)	(2.1 - 2.9)	(0.139–0.386)	(-27.3 – -22.3)
Lake 11	14	42.0	17	3.2	0.130	-22.2
		(37.1–45.8)	(15–20)	(2.8 - 3.6)	(0.058 - 0.244)	(-22.921.6)
Lake 12	ie 12 13 36.4 14 3.3 0.102 -2		-23.5			
		(33.5–41.5)	(12–17)	(3.0–3.7)	(0.062-0.212)	(-24.223.0)

char from tundra and polar desert lakes.

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284 *3.2. Diet influences on long-term growth rates and mercury concentration*

Growth rates based on length-at-age measurements showed large variation among Arctic char 285 or brook trout pooled across lakes (Fig. 1). Exponential curves fit to the length-at-age data were 286 highly significant for both species (p < 0.001). The length of brook trout ranged ~10–15 cm for 287 any given age while those of Arctic char ranged even more, from ~20-30 cm for older fish. On 288 289 average, brook trout categorized as faster-growing were 14 cm longer than slower-growing fish 290 of the same age (mean \pm SD: 31.9 \pm 3.8 cm versus 18.3 \pm 3.1 cm, *t*-test, *p* < 0.001, *n* = 30). 291 Similarly, Arctic char categorized as faster-growing were on average 20 cm longer than slower-292 growing fish of the same age (46.1 \pm 10.3 cm versus 25.7 \pm 10.5 cm, *t*-test, *p* < 0.001, *n* = 72). Note that both fast-growing and slow-growing Arctic char were found in each of the tundra and 293

294 polar desert ecoregions (Supplemental Fig. S2).



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Figure 1. Length-at-age of brook trout from 5 lakes (n = 57) and Arctic char from 7 lakes (n = 133) in the eastern Canada Arctic. Individual fish were categorized as faster- or slower-growing or unclassified based their length relative to the modelled exponential growth curve (see methods).

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Diet influenced the growth of brook trout and Arctic char. For both species, faster-growing fish fed at a higher trophic position than slower-growing fish (Mann-Whitney rank sum tests, p < 0.001, n = 30 or 72; Supplemental Fig. S3). The median trophic position of brook trout and Arctic char was 0.5 and 0.8 units higher, respectively, in faster-growing fish. Faster-growing brook trout had higher δ^{13} C values than slower growers (*t*-test, p = 0.006, n = 30) but that was

not the case for Arctic char (*t*-test, p = 0.26, n = 72). Thus, the growth of both fish species was related to their diet, with trophic position and carbon source (for brook trout) contributing to enhanced growth.



Figure 2. Comparison of log muscle THg concentration between growth categories after controlling for the effect of age for Arctic char (n = 72) or brook trout (n = 30) (see text for ANCOVA results).

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Total mercury concentrations were not lower in fish categorized as faster-growing (Fig. 2). For Arctic char, log THg concentrations of fish muscle increased with age and were higher in faster than slower growing fish (ANCOVA, age p < 0.001, growth category p < 0.001, n = 72). In contrast, log THg concentrations of brook trout increased with age but there was no growth effect (ANCOVA, age p < 0.001, growth category p = 0.83, n = 30).

Long-term growth rates estimated by a continuous variable, RLAA, showed similar patterns 319 with diet and fish THg concentration compared to the analysis with growth categories. RLAA 320 was positively correlated with trophic position for both Arctic char and brook trout (Fig. 3), 321 322 indicating that faster-growing fish tended to have a higher trophic position. For brook trout, RLAA was also positively correlated with muscle δ^{13} C (Pearson r = 0.50, p < 0.001, n = 57), but 323 324 that was not the case for Arctic char (Pearson r = 0.05, p = 0.56, n = 133). RLAA was positively 325 correlated with the log THg concentration of Arctic char (Pearson r = 0.52, p < 0.001, n = 133) and there was no correlation for brook trout (Pearson r = -0.10, p = 0.46, n = 57). Thus, this 326 327 continuous variable for growth (RLAA) similarly showed that growth rates were influenced by diet and faster-growing fish did not have lower THg concentrations. 328



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Figure 3. Correlations between trophic position and long-term growth rates (relative length-at-age) for brook trout and Arctic char.

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Explanatory variables of log-transformed fish THg concentrations were tested with multiple linear regression models that included age, long-term growth rate (RLAA), trophic position, δ^{13} C, and lake-specific MMHg accumulation near the base of the food web (using chironomid MMHg concentrations). Multiple regression models were highly significant for both salmonid species, explaining almost 75% of the variation in log THg concentration of fish within and among lakes (Table 2). Age, RLAA, and diet (chironomid) MMHg concentrations were highly significant explanatory variables and had positive effects for both brook trout and Arctic char.

340 Trophic position was also a significant positive explanatory variable in the Arctic char model.

Independed partial r^2 values indicated that age and diet (trophic position and chironomid MMHg)

342 were stronger explanatory variables, accounting more than 50% of the variance in fish THg

343 concentration, while growth rate (RLAA) was less important (Table 2). Positive correlations

between RLAA and fish THg concentrations in the models were contrary to a hypothesized

345 negative biodilution effect.

346

Table 2. Results of multiple linear regression models testing the relative importance of
explanatory variables of log-transformed muscle THg concentration for brook trout and Arctic
char.

Explanatory	Brook trout				Arctic char			
variable	Model $r^2_{adj} = 0.74$, $p < 0.001$, $n = 57$		Model $r_{adj}^2 = 0.71, p < 0.001, n = 133$					
	F	Р	Independent	Std. Co.	F	Р	Independent	Std. Co.
			partial r^2				partial r^2	
Age (years)	23.9	< 0.001	0.21	0.36	71.3	< 0.001	0.26	0.46
RLAA (cm)	31.6	< 0.001	0.10	0.58	30.4	< 0.001	0.16	0.33
Diet MMHg	34.9	< 0.001	0.45	0.97	23.2	< 0.001	0.19	0.30
(ng/g)								
TP	0.1	0.72			10.0	0.002	0.11	0.21
δ ¹³ C (‰)	0.6	0.44			0.0	0.984		

RLAA = relative length-at-age, Diet MMHg = lake-mean MMHg concentration of benthic

chironomids, TP = trophic position, Std. Co. = standardized coefficient, Independent partial r^2 =

352 contribution averaged over all possible orderings of explanatory variables (see methods)

353

354 *3.3. Influence of short-term growth rates on fish mercury concentrations*

355 Muscle RNA/DNA ratios were lower in brook trout (mean \pm SD=1.8 \pm 0.8, n = 57) than in

Arctic char (mean \pm SD = 2.9 \pm 1.1, *n* = 100), suggesting species-specific differences for this

between Arctic char ($1.2 \pm 0.5 \,\mu\text{g/mg}$) and brook trout ($0.9 \pm 0.3 \,\mu\text{g/mg}$), and therefore, species-358 specific differences in RNA/DNA ratios were due to the extent of RNA expression in muscle 359 360 cells. There was large (five-fold) variation in muscle RNA/DNA ratios among individual fish 361 within a species (Supplemental Fig. S4). The muscle RNA/DNA ratio of brook trout (n = 55-58) was not correlated with age, trophic 362 position, or RLAA (p > 0.05,) but decreased with length (Pearson r = -0.35, p = 0.007) and mass 363 (Pearson r = -0.28, p = 0.035), and increased with δ^{13} C values (Pearson r = 0.27, p = 0.037). For 364 Arctic char (n = 98-100), muscle RNA/DNA ratio increased with δ^{13} C (Pearson r = 0.23, p =365 0.022) and trophic position (Pearson r = 0.23, p = 0.022) but was not correlated with length, 366 mass, age or RLAA (p > 0.05). These results showed a positive, albeit weak, correlation between 367 368 diet and muscle RNA/DNA ratio for both species and a negative correlation with size for brook 369 trout. The log-transformed THg concentration of brook trout muscle was negatively correlated with its RNA/DNA ratio (Pearson r = -0.31, p = 0.021, Supplemental Fig. S5). This pattern 370 371 would be expected if biodilution from faster growth rates was resulting in lower THg concentrations. However, muscle RNA/DNA ratio was not a significant explanatory variable of 372 THg concentration for either species in multiple regression models that included age and diet 373 effects (Supplemental Table S3). 374

physiological proxy of growth rate (Supplemental Fig. S4). Muscle DNA content was similar

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The muscle RNA/DNA ratio of Arctic char varied seasonally in two tundra lakes where fish were collected in both winter and summer. Winter-sampled fish had a lower RNA/DNA ratios than those sampled in summer (*t*-test, p = 0.013, n = 51; Fig. 4), suggesting slower short-term muscle growth of char in winter. Muscle RNA/DNA ratio was not correlated (p = 0.23) with log THg concentration for the Arctic char in those two lakes, after including age and diet variables in

a multiple regression model (Supplemental Table S4).



Figure 4. Comparison of RNA/DNA ratios (a physiological proxy of short-term growth rates) in
Arctic char muscle between seasons (A) and ecoregions (B).

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381

Arctic char also showed regional variation in muscle RNA/DNA ratios. The subset of fish 385 collected in summer from more productive tundra lakes had higher ratios than those from polar 386 desert lakes (Mann-Whitney U test, p = 0.003, n = 68; Fig. 4). Those Arctic char from tundra 387 lakes also had higher length-at-age (RLAA) than char from polar desert lakes (Mann-Whitney U 388 test, p = 0.003, n = 68), consistent with the interpretation of RNA/DNA results that the char from 389 tundra lakes had faster summer growth rates. However, muscle RNA/DNA ratios of Arctic char 390 did not correlate (p = 0.19) with log THg concentrations for this subset of fish in a multiple 391 regression model with age and diet co-variates (Supplemental Table S5). Thus, inferred 392

differences in short-term growth rates between tundra and polar desert lakes did not explainvariation in muscle THg concentrations of Arctic char.

395 4. Discussion

396 *4.1. Explanatory variables of fish mercury concentrations*

Diet and age best explained muscle THg concentrations of two salmonid species in the Arctic 397 398 study lakes. Highly significant linear regression models accounted for close to 75% of the within- and among-lake variation of THg concentration in Arctic char or brook trout, of which 399 400 more than 50% was related to age and diet variables (trophic position, diet MMHg 401 concentration). Food web exposure to MeHg differed among sites while growth rates and diet varied among fish both within and between populations. The fishes were supported by benthic-402 derived carbon as indicated by their carbon stable isotopes, stomach contents (of brook trout), 403 and diet characterization in previous studies (of landlocked Arctic char; Gallagher and Dick, 404 2010; Gantner et al., 2010; Lescord et al., 2015). In contrast to our hypothesis, faster-growing 405 406 fishes had higher THg concentrations because their long-term growth rates (RLAA) were related to trophic position. Short-term growth rates, inferred from muscle RNA/DNA ratios, did not 407 explain or very weakly explained fish THg concentration. These findings were generated in 408 409 resource-poor Arctic lakes with low fish diversity (i.e. 1 or 2 fish species) and simple, benthicdominated food webs. Drivers of salmonid THg concentrations may be different in other Arctic 410 411 lakes with more complex food webs and greater pelagic production such as in northern Europe 412 (Eloranta et al., 2015; Kahilainen et al., 2016) or western Canada (Evans et al., 2005; Rohonczy 413 et al., 2020).

414 *4.2. Diet influences on growth*

A major finding of this study is that diet explained why faster growth rates were not 415 416 associated with lower THg concentrations in two Arctic salmonid species. Fish with faster growth for a given age fed at a higher trophic position, which was equivalent to approximately 417 one trophic level higher in the fastest-growing fish relative to the slowest growers (Fig. 4, 418 419 Supplemental Fig 3). This increase in trophic position can be explained by the consumption of fish. In resource-poor Arctic lakes, fish consumption confers an advantage for growth through 420 421 greater caloric intake (Power et al., 2009). Stomach contents of the brook trout showed some fish 422 were feeding on ninespine stickleback, which indicates fish consumption was occurring to some extent. For Arctic char, canibalism is well documented for populations in lakes of the eastern 423 Canadian Arctic based on stomach contents, nitrogen stable isotopes, and fish parasite tracers, 424 and the largest lacustrine morphs of char are piscivorous (Gallagher and Dick, 2010; Guiguer et 425 al., 2002; Hobson and Welch, 1995; Power et al., 2009). In one tundra lake with Arctic char 426 427 (Lake 6) where ninespine stickleback were also present, the higher trophic position of fastergrowing fish may have also been supported by consumption of stickleback (Gallagher and Dick, 428 2010). While fish consumption allowed for greater growth by a given age, it also promoted 429 430 mercury bioaccumulation because of greater mercury intake from eating higher trophic level prey. For brook trout, there was no significant difference in THg concentration between fish 431 432 categorized as slower- or faster-growing (Fig. 2), nor was trophic position a significant variable 433 in the multiple regression model (Table 2). Further, muscle RNA/DNA ratios of brook trout were 434 negatively correlated with THg concentration (Fig. S5). Those results possibly reflect a dilution 435 effect where greater mercury intake from feeding at a higher trophic position was somewhat 436 offset by biomass dilution from faster growth for brook trout. Nevertheless, the multiple

regression modelling indicated that the net effect of dietary exposure and age on muscle mercury 437 concentration was greater than any potential offset by biomass dilution. In contrast with this 438 439 study, faster growth rates in fish are not necessarily associated with consuming more contaminated prey, and biodilution of mercury can occur in temperate lakes from consumption of 440 higher caloric invertebrate prey with lower mercury concentrations (Karimi et al., 2016). 441 442 Earlier research by van der Velden et al. (2012) on an Arctic char population in Heintzelman Lake (Ellesmere Island, Canada) found that muscle THg concentration did not differ between 443 444 slow- and fast-growing fish. In that study, the lake-dwelling char were insectivores, and slower growth was likely due to greater energy allocation to reproduction rather than a difference in 445 trophic position. Their research is in agreement with findings of the current study regarding the 446 lack of growth rate influence on mercury fish concentration although the underlying drivers of 447 growth in Arctic char were different. Lavigne et al. (2010) similarly did not find an effect of 448 449 biodilution on THg concentrations of another salmonid species, lake trout, in northern Quebec 450 (Canada) but biodilution was reported for two other fish species.

451 4.3. Application of RNA/DNA ratio in muscle to test short-term growth rates

To our knowledge, this study is the first to use RNA/DNA ratios in fish muscle to examine 452 453 short-term growth effects on mercury concentration. On average, differences in inferred shortterm growth were found for Arctic char between summer and winter, and between tundra and 454 455 polar desert lakes using this physiological proxy. However, muscle RNA/DNA values varied 456 considerably among individual fish within the same season or ecoregion, and high variability has been reported elsewhere for this measurement in fish, including Arctic char (Ferguson et al., 457 458 1990). Fish muscle RNA/DNA ratios can respond rapidly (i.e., days to weeks) to changes in food 459 availability or environmental conditions (Chícharo and Chícharo, 2008; Stierhoff et al., 2009),

460 and that time-scale may be shorter than rates of change in muscle mercury concentration.

461 Additional factors such as fish age and diet, and water temperature can also influence fish muscle

462 RNA/DNA ratios (Chícharo and Chícharo, 2008; Roberts et al., 2011). Research is

463 recommended to further constrain the appropriate time scale and conditions for using muscle

464 RNA/DNA ratios to track potential influences of short-term growth on mercury concentration.

465 *4.4. Implications for somatic growth dilution theory*

This theory predicts that more efficient growth will result in a lower body mercury 466 concentration (Trudel and Rasmussen, 2006). More specifically, the prediction has been phrased 467 468 as "fish that grow to a larger size for a given age and prey intake will have a lower mercury concentration" to acknowledge the influence of prey type and age (van der Velden et al., 2012; 469 Ward et al., 2010). This study did not specifically test SGD because the theory is predicated on 470 growth efficiency, and key variables for energy allocation (consumption rate, activity rate, 471 reproductive effort) were not measured (Trudel and Rasmussen, 2006). Nevertheless, our 472 473 findings on diet-growth influences suggest that bioenergetic assumptions of SGD are not realistic in the case of lake-dwelling salmonids in the eastern Canadian Arctic. Size variation and 474 morphological diversity are common within populations of Arctic char, and differences in 475 476 growth are probably not determined by efficiency of energy use but rather by factors such as diet, resource availability, and habitat (Guénard et al., 2010; Power et al., 2009), which also 477 478 influence mercury exposure. Muscle biodilution of mercury has been reported for other Arctic 479 fish species and regions such as lake whitefish in a Finnish subarctic lake (Keva et al., 2017), and perch, roach and northern pike in a Norwegian subarctic lake (Sharma et al., 2008). A common 480 481 element of studies that found biodilution effects is the rapid onset of growth which occurred for: 482 (1) juvenile fish (Ward et al., 2010), (2) between winter starvation and summer growth (Keva et

483 al., 2017), (3) after the removal of predation pressure (Sharma et al., 2008), or (4) from stocking

484 of prey fish (Lepak et al., 2012). Future research should focus on identifying the key

485 environmental conditions where fish mercury concentrations are influenced by SGD.

486 *4.5. Fish mercury concentrations in a changing Arctic environment*

The length-at-age of lake-dwelling Arctic char declines with latitude in the Canadian Arctic 487 488 (Chavarie et al., 2010), reflecting slower growth in colder, lower productivity ecosystems. Our study lakes spanned a 20° latitudinal gradient, with warmer water temperatures and greater 489 490 organic matter in subarctic taiga lakes compared with tundra and polar desert lakes. Although the 491 effects of temperature and productivity were not explicitly tested, similar modelling results were found (i.e., the importance of age and diet for fish bioaccumulation of mercury) for both a fast-492 growing fish species in the taiga lakes and a slower growing species in the tundra and polar 493 desert lakes. Previous reviews of Arctic climate change have suggested altered growth conditions 494 495 for fish may have a long-term impact on their contaminant concentrations (Reist et al., 2006a; 496 Stern et al., 2012). However, this study for two salmonid species and earlier work on Arctic char (van der Velden et al., 2013; 2012) have demonstrated age (period of exposure) is a more 497 important determinant of mercury concentration than length (growth rate). Higher mercury in 498 499 fish that take longer to reach a specific length (length-adjusted concentration) is due to the older age of the fish rather than SGD. This study found MMHg concentrations near the base of the 500 501 food web were a key driver of among-lake differences in fish mercury concentrations. Together, 502 these findings suggest environmental change in the Arctic that affects food web exposure to MMHg (e.g., atmospheric mercury deposition, MMHg production, uptake at the base of the food 503 504 web, food web structure) may have more of a long-term impact on mercury bioaccumulation in 505 fish than improved growth from warmer temperatures or increased lake productivity (Reist et al.,

2006b; Stern et al., 2012). Bioenergetic modelling of mercury bioaccumulation in lake-dwelling
Arctic char similarly showed their mercury concentrations were highly responsive to changes in
dietary mercury and not changes in water temperature (Drevnick, 2014). Since warming
enhances the production of methylmercury in Arctic lake sediment (Hudelson et al., 2020),
tracking future changes in MMHg exposure to food webs in Arctic lakes is warranted.

511 **5.** Conclusions

Using data on two salmonid fish species from lakes in three ecoregions of the eastern 512 513 Canadian Arctic, we tested the hypothesis that faster-growing fish have lower mercury 514 concentrations after quantifying the influence of age and diet. We found age and diet indices were the dominant explanatory variables of muscle mercury concentrations. Faster-growing fish 515 had higher mercury concentrations because those fish were feeding at a higher trophic 516 517 position. In these resource-poor Arctic lakes, differences in growth rate were influenced by diet rather than growth efficiency. These findings contribute to the evolving science on growth rate 518 519 effects in relation to other drivers of fish mercury concentration, which is needed to evaluate multiple stressors currently impacting aquatic ecosystems such as climate change in the Arctic. 520

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535

536 Supplemental Information

537 The supplemental information to this article can be found online at:

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Highlights

- Age and diet best explained mercury concentrations of fishes in Arctic lakes.
- Long-term growth rates (length-at-age) were related to fish trophic position.
- Faster-growing fish had a higher trophic position and mercury concentration.
- Muscle RNA/DNA ratios indicated differences in short-term growth rates of fishes.
- Muscle RNA/DNA ratios were a poor predictor of fish mercury concentrations.



Diet influences on growth and mercury concentrations of two salmonid fish species from lakes in the eastern Canadian Arctic

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Ecoregion	region Nearest community		Local lake name	Latitude	Longitude	Lake area	Mean	Water DOC	Water Chl
				(°N)	(° W)	(km2)	depth (m)	(mg/L)	(µg/L)
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 1	Christmas Lake	55°22'10"	77°37'04"	0.11	1.0	4.2	0.9
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 2		55°18'16"	77°42'56"	0.01	0.3	4.1	0.8
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 3	Kachishayoot Lake	55°20'03"	77°37'31"	0.29	1.9	4.0	1.7
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 4		55°20'01"	77°35'48"	0.39	0.9	5.8	2.5
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 5		55°20'14"	77°36'42"	0.10	1.2	3.9	2.1
Arctic tundra	Iqaluit	Lake 6	Iqalugaajuruluit Lake	63°41'06"	68°22'34"	0.54	8	3.0	0.4
Arctic tundra	Iqaluit	Lake 7		63°49'30"	68°36'14"	0.56	12	1.6	0.4
Arctic tundra	Iqaluit	Lake 8	Tasirluk (Crazy) Lake	63°52'29"	68°28'40"	4.5	7.5	1.0	0.5
High Arctic polar desert	Resolute Bay	Lake 9	Amituk Lake	75°02'44"	93°47'15"	0.38	19	1.7	0.3
High Arctic polar desert	Resolute Bay	Lake 10	North Lake	74°46'37"	95°05'47"	0.63	7	0.9	0.3
High Arctic polar desert	Resolute Bay	Lake 11	Resolute Lake	74°41'15"	94°56'33"	1.21	9	1.4	0.2
High Arctic polar desert	Resolute Bay	Lake 12	Small Lake	74°45'33"	95°03'37"	0.15	3	2.5	0.4

Table S1. Ecoregion, location, physical characteristics and water chemistry of the study lakes in the eastern Canadian Arctic.

DOC = dissolved organic carbon, Chl = chlorophyll; water chemistry data are from Chételat et al. (2018), except for Lake 9 (Amituk) for which water chemistry data were collected earlier by Gantner et al. 2010)



Figure S1. Map of lake locations at the subarctic taiga (Kuujjuaraapik-Whapmagoustui), Arctic tundra (Iqaluit) and High Arctic polar desert (Resolute Bay) study sites.

Age estimations using scales

Fish age was estimated with otoliths using the crack and burn method. Otoliths were not available for a small subset of brook trout and Arctic char (<10%), in which case fish age was estimated from scale annuli. We found that scales gave similar age estimates to otoliths for young fish (<6 years) although slightly underestimated (by 0-2 years) relative to otoliths, as reported elsewhere (Stolarski et al. 2008). We therefore used species-specific poisson regressions derived using scale and otolith data from our study fish to estimate otolith age from scales for young fish (6 brook trout and 9 Arctic char).

Comparison of age estimates from scales and otoliths for brook trout (n=13) and Arctic char (n=32) from study fish.

Fish species	Scale Age	Otolith Age
Brook trout	2	2
Brook trout	2	2
Brook trout	3	3
Brook trout	3	5
Brook trout	3	5
Brook trout	3	3
Brook trout	3	4
Brook trout	3	3
Brook trout	4	6
Brook trout	4	4
Arctic char	3	3
Arctic char	3	5
Arctic char	3	4
Arctic char	3	5
Arctic char	4	7
Arctic char	4	7
Arctic char	4	6
Arctic char	4	6
Arctic char	4	8
Arctic char	4	7
Arctic char	4	9
Arctic char	4	5
Arctic char	4	6
Arctic char	5	8
Arctic char	5	5
Arctic char	5	7

Arctic char	5	5
Arctic char	5	5
Arctic char	5	6
Arctic char	5	6
Arctic char	5	5
Arctic char	5	9
Arctic char	5	6
Arctic char	6	7
Arctic char	6	8
Arctic char	6	7
Arctic char	6	6
Arctic char	6	6
Arctic char	6	7
Arctic char	6	8
Arctic char	6	7
Arctic char	6	8

Poisson regressions were fitted to the data (using the glm function in R) to estimate fish age based on scales, when otolith age estimates were not available.

Brook trout poisson regression model predictions (ages for 6 fish were predicted based on scale ages from 2-4 years):

- Scale age = 2, predicted otolith age = 2
- Scale age = 3, predicted otolith age = 3
- Scale age = 4, predicted otolith age = 5

Arctic char poisson regression model predictions (ages for 9 fish were predicted based on scale ages from 3-6 years):

- Scale age = 3, predicted otolith age = 5
- Scale age = 4, predicted otolith age = 6
- Scale age = 5, predicted otolith age = 7
- Scale age = 6, predicted otolith age = 7

Table S2. Lake-specific sample sizes, MMHg concentrations, and δ^{15} N values of chironomids used to estimate fish trophic position and dietary MMHg exposure near the base of the food web. Chironomids were collected around the same time as fish in all lakes except Lake 9 (fish in 2014, chiromids in 2011). Additional chironomid data from collections in 2011 were included for Lake 11&12 to supplement the 2014 measurements.

Ecoregion	Nearest community	Lake ID	Mean ± SD [MMHg]	$\frac{\text{Mean} \pm \text{SD}}{\delta^{15} \text{N} (\%)}$	Data Source
			ng/g dry weight	• • • • • • • • • • • • • • • • • • • •	
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 1	75 ± 13 (3)	4.5 [4.7, 4.3] (2)	1
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 2	160(1)	4.0 (1)	1
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 3	32 (1)	2.8 (1)	1
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 4	37 ± 8 (3)	2.8 [2.9, 2.7] (2)	1
Subarctic taiga	Kuujjuaraapik-Whapmagoustui	Lake 5	$32 \pm 9 (3)$	2.1 [2.4, 1.8] (2)	1
Arctic tundra	Iqaluit	Lake 6	4 ± 2 (4)	6.0 ± 0.7 (4)	1
Arctic tundra	Iqaluit	Lake 7	19 [9, 29] (2)	5.7 [5.9, 5.5] (2)	1
Arctic tundra	Iqaluit	Lake 8	16 ± 7 (3)	3.7 ± 0.2 (3)	1
High Arctic polar desert	Resolute Bay	Lake 9	173 ± 83 (9)	5.6 ± 0.3 (7)	2
High Arctic polar desert	Resolute Bay	Lake 10	78 ± 19 (4)	10.8 ± 0.9 (4)	1
High Arctic polar desert	Resolute Bay	Lake 11	38 ± 13 (15)	6.8 ± 1.5 (16)	1,2
High Arctic polar desert	Resolute Bay	Lake 12	40 ± 28 (10)	4.6 ± 2.3 (8)	1,2

Sample sizes for mean \pm SD are provided in parentheses. Individual values are provided in square brackets when n=2.

Data Sources:

1) Chételat et al. (2018)

2) Chételat et al. (2014)

Lake ID	Macroinvertebrate	Mean ± SD [MMHg]	Mean ± SD	Mean \pm SD
	taxon	ng/g dry weight	δ ¹³ N (‰)	δ ¹³ C (‰)
Lake 1	Amphipods (<10 mm)	79 ± 21 (3)	2.9 ± 0.1 (3)	-27.3 ± 0.4 (3)
	Amphipods (>10 mm)	144 (1)		
	March flies*	54 [70, 37] (2)	5.4 [5.5, 5.2] (2)	-26.4 [-26.2,-26.6] (2)
	All taxa	$81 \pm 37(6)$	3.9 ± 1.3 (5)	$-26.9 \pm 0.6 (5)$
Lake 2	Dragon fly nymphs	482 (1)	5.0(1)	-29.0 (1)
Lake 3	Amphipods (<10 mm)	43 (1)	1.0(1)	-20.3 (1)
	Amphipods (>10 mm)	53 [49, 58] (2)	2.7 [2.8, 2.5] (2)	-23.0 [-24.8, -21.2] (2)
	Mayflies	20.2 (1)	0.5 (1)	-20.2 (1)
	All taxa	$42 \pm 16(4)$	1.7 ± 1.1 (4)	-21.6 ± 2.2 (4)
Lake 4	Amphipods (<10 mm)	43 [40, 45] (2)	1.6 [1.6, 1.5] (2)	-23.2 [-23.0, -23.3] (2)
	Amphipods (>10 mm)	71 (1)	3.0(1)	-24.5
	Caddisflies	17 (1)	1.2 (1)	-23.3
	All taxa	43 ± 22 (4)	1.9 ± 0.8 (4)	$-23.5 \pm 0.7 (4)$
Lake 5	Caddisflies	49(1)	2.1 (1)	-25.6 (1)

Table S3. MMHg concentrations and values of δ^{15} N and δ^{13} C for macroinvertebrate taxa sampled in subarctic taiga lakes containing brook trout. See Table S2 for chironomid data also collected in those lakes.

Sample sizes for the mean \pm SD are provided in parentheses. Individual values are provided in square brackets when n=2.

*Obtained from gut contents



Figure S2. Comparison of MMHg concentrations in chironomids with other macroinvertebrate taxa from the same lake. Values are lake means ± 1 standard deviation (see Tables S2&S3 for data).

	Lake 1	Lake 2	Lake 3	Lake 4	Lake 5
Stomach content item	n=11	n=10	n=13	n=14	n=10
Unidentified flies	9	10	54	57	10
March flies (Bibionidae)	100	70	54	7	10
Chironomids (Chironomidae)	9	10	0	50	40
Beetles (Coleoptera)	18	40	23	7	0
Backswimmers (Notonectidae)	9	0	0	0	0
Amphipods (Amphipoda)	81	0	15	36	0
Dragonfly nymphs (Anisoptera)	0	30	15	0	50
Snails (Gastropoda)	0	50	7	0	0
Caddisflies (Trichoptera)	0	0	54	36	0
Ninespine stickleback	0	0	7	14	20
(Pungitius pungitius)					

Table S4. Frequency of occurrence (%) of prey items in the stomachs of brook trout from subarctic taiga lakes. Sample sizes (n) are the number of fish examined in each lake.



Figure S3. Length-at-age of Arctic char from 7 Arctic lakes (n = 133) separated out by ecoregion (polar desert, tundra) in the eastern Canadian Arctic. Individual fish were categorized as faster- or slower-growing or unclassified based on their length compared to the modelled exponential growth curve.



Figure S4. Comparison of trophic position for brook trout and Arctic char categorized as either faster-growing or slower-growing fish.



Figure S5. Comparison of muscle RNA/DNA ratios of brook trout and Arctic char.



Figure S6. Negative correlation between log muscle THg concentration of brook trout and their muscle RNA/DNA ratio. This pattern would be expected if biodilution from faster growth rates was resulting in lower THg concentrations. However, muscle RNA/DNA ratio was not a significant explanatory variable of THg concentration for brook trout after accounting for the influence of age, trophic position, diet MMHg, and δ^{13} C in multiple regression models (full model results in Table S5).

Table S5. Results of multiple linear regression models testing the relative importance of muscle

 RNA/DNA ratios in explaining log-transformed muscle THg concentration for brook trout and

 Arctic char, after quantifying the influence of diet and age.

Explanatory	tory Brook tro				Arctic char			
variable	Mod	el $r^{2}_{adj} = 0$.55, p < 0.001, n	$p, p < 0.001, n = 54$ Model $r^{2}_{adj} = 0$.47, p < 0.001, n = 98	
	F P		Independent	Std.	F	Р	Independent	Std.
			partial <i>r</i> ²	Co.			partial <i>r</i> ²	Co.
Age (years)	13.2	< 0.001	0.23	0.43	35.3	< 0.001	0.35	0.60
TP	1.8	0.19			8.4	0.005	0.13	0.37
Diet MMHg	6.2	0.016	0.34	0.54	0	0.96		
(ng/g)								
RNA/DNA ratio	0.6	0.43			1.4	0.23		
δ ¹³ C (‰)	2.8	0.10			1.2	0.27		

TP = trophic position, Diet MMHg = lake-mean MMHg concentration of benthic chironomids,

Std. Co. = standardized coefficient, Independent partial r^2 = partial r^2 contribution averaged over

all possible orderings of explanatory variables

Table S6. Results of a multiple linear regression model testing the relative importance of muscle RNA/DNA ratios in explaining log-transformed muscle THg concentration for summer and winter captured Arctic char in two lakes, after quantifying the influence of diet and age.

Explanatory variable	Arctic char Model $r^2 = 0.87$, $n < 0.001$, $n = 51$						
variable	F	Std. Co.					
Age (years)	29.1	< 0.001	0.49	0.46			
ТР	40.9	< 0.001	0.20	0.62			
Diet MMHg (ng/g)	28.8	< 0.001	0.19	0.58			
RNA/DNA ratio	1.5	0.23					
δ ¹³ C (‰)	0	0.88					

TP = trophic position, Diet MMHg = lake-mean MMHg concentration of benthic chironomids, Std. Co. = standardized coefficient, Independent partial r^2 = partial r^2 contribution averaged over all possible orderings of explanatory variables **Table S7.** Results of a multiple linear regression model testing the relative importance of muscle RNA/DNA ratios in explaining log-transformed muscle THg concentration Arctic char in tundra and polar desert lakes (captured during summer), after quantifying the influence of age and diet.

Explanatory variable	Arctic char Model $r^{2}_{adj} = 0.65, p < 0.001, n = 68$						
	F	P	Independent partial ^{r2}	Std. Co.			
Age (years)	92.5	< 0.001	0.44	0.94			
Diet MMHg (ng/g)	41.2	< 0.001	0.12	-0.60			
ТР	13.9	< 0.001	0.10	0.32			
δ^{13} C (‰)	0.2	0.69					
RNA/DNA ratio	0	0.91					

TP = trophic position, Diet MMHg = lake-mean MMHg concentration of benthic chironomids, Std. Co. = standardized coefficient, Independent partial r^2 = partial r^2 contribution averaged over all possible orderings of

explanatory variables

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