

# Effect of water warming on the structure of biofilm-dwelling communities

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## 1 Effect of water warming on the structure of biofilm-dwelling

## 2 communities

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#### 26 Abstract

27 Water warming, either resulting from global or local drivers, is of growing concern for 28 aquatic environments. Changes in thermal regimes may alter the functioning and the trophic 29 dynamics of river ecosystems since most aquatic biota like micro-algae, invertebrates, or fish are poikilotherms. Since biofilms integrate basal and intermediate components of aquatic 30 31 food webs, examination of their responses to warming should contribute to a better 32 understanding of the effects of ongoing climate change on river ecosystems. In this study, we considered sites that were or were not exposed directly to thermal effluents from a nuclear 33 34 power plant as "in situ laboratories" of the effect of climate change and water warming. To 35 compare the structure of algal and invertebrate communities dwelling in epilithic biofilms subjected to different thermal regimes in the Garonne River, sets of concrete tiles were placed 36 37 in the river and the biofilm growing on these standard substrates was sampled after two 38 months of colonisation. The site located within the thermal plume directly downstream of the 39 thermal effluents showed warmer average water temperature (mean  $\Delta T$  of about 1.5°C over 40 the study period) and greater daily temperature fluctuations in comparison to the control site. 41 Algal and invertebrate biomass did not differ significantly, but the structure of communities 42 differed markedly between sites. At the heated site, the proportion of cyanobacteria was 43 higher, while there were fewer micro-crustaceans and larvae of insects and bivalves. However, the population of chironomids showed a greater proportion of relatively large 44 45 individuals. Interestingly, nematodes were more abundant, and their communities comprised more species at the heated site. Their feeding-types were also more diverse showing a greater 46 47 proportion of algivores, large-sized omnivorous and predatory species. Changes in the 48 abundance and age structure further showed that some bacterivore species, such as 49 Eumonhystera vulgaris, thrived especially well at the heated site, whereas another 50 bacterivore, Monhystrella paramacrura, thrived well at the control site. In the context of

- 51 climate change, our results highlight that warmer and more fluctuating water temperatures
- 52 have the potential to alter the body-size spectrum, species composition, distribution of
- 53 feeding specialisations and age structure within biofilm-dwelling communities.
- 54

### 55 Keywords

56 Meiofauna, Warming, Freshwater, Community ecology, Species distribution

#### 57 **1. Introduction**

The question of how organisms and ecosystems respond to a rise in temperatures has 58 attracted growing attention from the scientific community because of our need to understand 59 60 and foresee the effects of the ongoing climate change on biological diversity and ecosystem 61 functionality. In aquatic environments, spatial variability in water temperature can be natural 62 (e.g. geothermal activity, presence of a water source) or result from direct local (e.g. 63 deforestation, industrial activities) or indirect global (i.e. climate change) anthropogenic disturbances (Caissie, 2006; Van Vliet et al., 2011). Alteration of thermal regimes can be a 64 65 major determinant of changes in the diversity and resilience of aquatic biota (Steel et al., 66 2017; Reid et al., 2019). For example, Mouthon and Daufresne (2006) recorded a dramatic reduction in density and diversity of mollusc communities in the Saône River (France) during 67 68 the exceptional European heat wave in the summer of 2003. The recovery of the mollusc 69 communities studied was not fully observed even eight years after the heat wave, and 70 profound changes in community structure have persisted over the long term (Mouthon and 71 Daufresne, 2015). Streams and rivers that are naturally or artificially heated can be 72 considered relevant and realistic "in situ laboratories" to study the effects of climate change 73 and water warming on aquatic organisms from the individual to the ecosystem level 74 (Woodward et al., 2010). Several examples of such experiments have contributed to a better 75 understanding of the effects of global warming on aquatic biodiversity and ecosystem 76 functioning. In their review, O'Gorman et al. (2012) summarised a decade of research conducted in an active geothermal region (Hengill, Iceland). The underground geothermal 77 78 heating of streams in this region allowed for comparative studies between naturally heated 79 and non-heated systems. These works have revealed strong influences of water temperature 80 on the structure of macroinvertebrate and algal assemblages, and on ecosystem functioning. 81 Sandblom et al. (2016), Svensson et al. (2017), and Huss et al. (2019) have documented some 82 of the most remarkable results obtained from the "Biotest Enclosure" experiment carried out 83 in the Baltic Sea (Sweden). For research purposes, heated cooling water from a nuclear power 84 plant was diverted into an isolated atoll in the late 1970s. Since then, a temperature difference 85 of 5–10°C has been maintained between the inside and outside of the enclosure. This made 86 possible the study of the physiological responses of the European perch (Perca fluviatilis) to 87 water warming, highlighting the fact that global warming could outpace the adaptive capacity 88 of ectotherms (Sandblom et al., 2016). Lastly, Boulêtreau et al. (2014) compared biofilms collected inside or outside the thermal plume of a nuclear power plant located on the Garonne 89 90 River (France) and demonstrated that water warming affected the denitrification activity of 91 the biofilms through changes in the composition of their bacterial community. Despite the 92 key insights brought by these studies into the responses of aquatic ecosystems to climate 93 change, we still lack a comprehensive understanding of the warming-induced ecological 94 effects that can be expected in the future on more complex and realistic communities made 95 up of different interacting biological compartments.

96 Biofilms are colonies of microscopic organisms that become large enough to create 97 visible, macroscopic structures. In aquatic ecosystems, biofilms are hotspots of biodiversity 98 supporting pivotal ecosystem functions such as primary production and nutrient cycling. 99 Thereby, biofilms fix and transform drifting particles and dissolved inorganic nutrients into 100 attached biomass that can then be used by a variety of consumers (Battin et al., 2016; Lyautey 101 et al., 2013; Risse-Buhl et al., 2012; Romaní et al., 2004; Teissier et al., 2007; Weitere et al., 102 2018). Ecological successions in river biofilms are characterised by an accretion phase 103 followed by a detachment phase as the thickness and porosity of mature biofilms increase 104 their sensitivity to shear stress and grazers (Biggs, 1996). Nevertheless, biofilms can also 105 self-detach in summer under warmer temperatures and low flow because of bacterial 106 decomposition of senescent basal algal layers (Boulêtreau et al., 2006). The density and

107 biomass of biofilm-dwelling grazers correlate well with the total biomass and chlorophyll a 108 content of the biofilm because biofilms represent both a source of food and a habitat for 109 micro-grazers like nematodes (Majdi et al., 2011, 2012a; Peters and Traunspurger, 2005; 110 Schroeder et al., 2012; Vidakovic et al., 2011). Overall, most ecosystem services provided by 111 aquatic biofilms could be strongly affected by global warming (Romaní et al., 2016). 112 Experimental studies of the response of biofilm communities to water warming have shown 113 faster accretion phases (especially higher bacterial growth rates), greater heterotrophic use of 114 organic compounds (Norf et al., 2007; Villanueva et al., 2011), and enhanced filtration rates 115 of peritrich ciliates and rotifers under warmer conditions (Kathol et al., 2009). In a 21-days in 116 *situ* warming experiment (on average +2.5°C), warmed-up biofilms showed faster 117 colonisation and maturation, and greater respiration and denitrification rates due to changes 118 in the structure of bacterial communities at the expense of phototrophic biomass (Boulêtreau et al., 2014). 119

120 The responses of species to water warming essentially depends on their physiological 121 ability to cope with warmer temperatures, or fluctuations thereof, and their capacity to cope 122 with changes in the abundance of the species with which they interact (i.e. their predators, 123 competitors, and prey) (O'Gorman et al., 2012; Woodward et al., 2010). Our aim with this 124 study was to compare heated and non-heated natural river biofilms, with their complex 125 assemblage of prey, micro-grazers, and predators to examine whether their community 126 structure showed coherent responses to different thermal conditions. We exposed biofilms to 127 heated effluents from the nuclear power plant of Golfech (Garonne River, France). At the 128 chosen experimental site, the warming averages +2.5°C (Boulêtreau et al., 2014), which lies in the range of most Earth surface warming scenarios expected for the end of the 21<sup>st</sup> century 129 130 (IPCC, 2014). Moreover, the amplitude of daily fluctuating temperatures is increased because 131 of the passage of discharged water in natural draft wet-cooling towers. We monitored the

132 algal and invertebrate communities in biofilms, with particular emphasis on the taxonomic 133 and functional structure of nematodes (numerically dominant metazoans). The biofilms grew 134 during a low-flow period during the summer-fall, right after a substantial heat wave had hit 135 Europe (29 July - 7 August 2018). Our hypotheses regarding the responses of biofilm-136 dwelling communities to warmer water temperature conditions were: (1) lower algal biomass 137 and changes in the structure of algal communities; (2) changes in the structure of invertebrate communities, especially through the loss of thermosensitive taxa; (3) shift in the size-138 139 spectrum of insect larvae due to changes in cohort maturation rates; (4) greater taxonomic 140 and trophic diversity of biofilm-dwelling nematodes, as previously observed in the summer in 141 the Garonne River (Majdi et al., 2011).

142

#### 143 **2. Material and methods**

144 *2.1. Study site* 

145 The experiment was conducted from August to October 2018 downstream of the nuclear 146 power plant of Golfech on the Garonne River (SW France, 44°07'04.3"N, 0°50'02.7"E). This 147 nuclear power plant is operated in an open loop with recirculation and is equipped with 148 natural draft wet-cooling towers to reduce the temperature of the blowdown water. The water 149 is withdrawn from the river for its cooling system and the blowdown water is then discharged 150 heated at a series of outlet pipes located directly in the middle of the riverbed over a width of 151 about 100 m (Fig. 1). The resulting thermal plume covers a cone of a few hundred meters long by a hundred meters wide at its base. At this site, the Garonne River is 200 m wide with 152 153 a relatively shallow (from 0.5 to 1 m depth) riverbed consisting of clayey bedrocks with large 154 stone slabs, cobble bars, or sand/pebbles banks depending on local hydrodynamics, with 155 substantial coverage of several hydrophyte species (e.g., Potamogeton spp., Myriophyllum 156 spicatum L.).

157

#### 158 2.2. Experimental design

159 Forty concrete tiles (60 x 40 x 4 cm, 21.5 Kg each) were used as standard substrates for the 160 growth of biofilm and were immerged on 7 August 2018. They were arranged in two blocks of 20 tiles each (5 rows of 4 tiles) facing the flow direction at a depth of 50 cm. One block 161 162 (heated site) was positioned within the thermal plume, 20 m downstream from the outlet 163 pipes of the nuclear power plant and about 40 m from the right bank of the river (Fig. 1). The 164 other block (control site) was positioned outside the thermal plume on the same transect 165 about 120 m from the right bank of the river (Fig. 1). We observed that each tile in both 166 blocks showed homogeneous and similar depths and bottom flow velocities (depth =  $50 \pm 2$ cm; flow velocity =  $30 \pm 5$  cm s<sup>-1</sup>). A temperature data-logger (measurement frequency = 5 167 168 min) linked to a buoy was positioned 4 cm above the riverbed and fixed to a steel rod 169 hammered in the riverbed at the centre of each block. Oxygen concentration, conductivity, 170 and pH were recorded monthly using a portable probe (U50, HORIBA) (Table S1 in 171 Appendix). When sampling biofilms, water samples were also collected at each site and analysed after filtration on glass fibre filters (GF/C, Whatman) to measure the concentration 172 of dissolved organic carbon using the combustion catalytic oxidation method (TOC-L, 173 174 Shimadzu), and of anions and cations using high-performance ionic chromatography (ISC-175 5000+ and DX-120, Dionex).

176

177 2.3. Sampling of biofilm

All tiles were sampled on 2 October 2018 (i.e. for a total of 40 samples: one sample per tile
with 20 tiles for each of the two sites) after 56 days of immersion; a period deemed long
enough to allow for the settlement of mature biofilm communities (Majdi et al., 2012b; Norf
et al., 2009). We collected 3.14 cm<sup>2</sup> patches of biofilm directly underwater using a brush-

182 sampler (Peters et al., 2005). For invertebrates, four patches were collected at random 183 locations on each tile, poured on 20 µm meshes, and pooled in a 50 ml centrifuge tube. 184 Invertebrate samples were preserved in a buffered formaldehyde solution (final concentration 185 4%) and stained with a few drops Rose Bengal. For algal pigments, two patches were collected and poured in a 50 ml centrifuge tube. Samples were returned to the laboratory in 186 187 an icebox. The river discharge was very stable (low-flow conditions) during the biofilm 188 colonisation period (i.e. from 7 August to 2 October 2018), but a major flood occurred two 189 weeks after collecting the samples (16 October 2018), which prevented us from continuing 190 the experiment.

191

#### 192 2.4. Invertebrate identification and data pretreatment

193 Preserved invertebrates were counted, categorised into size classes (based on maximal length 194 and width), and identified under a dissecting microscope at 40x magnification. While 195 counting, 50 nematode individuals were picked out randomly from each sample, 196 progressively transferred to glycerol, and mounted on slides following the method of 197 Seinhorst (1959). Nematodes were identified at the species level and classified into age and 198 sex categories. Nematode individuals were also classified according to their feeding type 199 based on the morphology of their buccal cavity, after the classification of Traunspurger 200 (1997): deposit-feeders (mainly bacterivorous given the small size of their unarmed buccal 201 cavity), epistrate-feeders (potent algivores given the presence of small teeth in their buccal 202 cavity allowing them to crack diatom frustules), chewers (omnivores or strict predators given 203 their wide mouth armed with large teeth), and suction-feeders (omnivorous or fungal/plant-204 feeders given that their hollow stylet allows them to perform asymmetrical predation on other 205 animals and hyphae). Thirty individuals of chironomids were also randomly sorted out from 206 each sample during the counting, photographed under the microscope, and measured to the

207	nearest $\mu$ m using the ImageJ image analysis software. Invertebrate biomass was estimated
208	from allometric relationships found in the literature (see Table S4 in Supplementary
209	Information) and expressed as dry mass per area of biofilm (µgDM 10cm <sup>-2</sup> ).
210	
211	2.5. Pigment analysis
212	In the laboratory, biofilm samples were pelletized by centrifugation (3000 g, 20 min), freeze-
213	dried, and stored at -80°C until analysis. The pellets were then homogenised and 200 mg
214	subsamples were extracted in 5 mL 98% cold-buffered methanol (with 2% of 1 M ammonium
215	acetate) in a 4°C sonication tub followed by a 15 min extraction in the dark at –20°C. The
216	solution was centrifugated (3000 g, 20 min), the supernatant was kept aside at $-20^{\circ}$ C, and the
217	pellet was extracted again following the same steps as described above. Thereby, for each
218	sample, a total of 10 mL pigment solution was obtained from which 2 mL was subsampled
219	and filtered through a 0.2 $\mu$ m polytetrafluoroethylene syringe filter. The pigment solution
220	was analysed using a high-performance liquid chromatograph consisting of a 100 mL loop
221	autosampler and a quaternary solvent delivery system coupled to a diode array
222	spectrophotometer (LC1200 series; Agilent). The mobile phase was set after the protocol
223	described by Barlow et al. (1997). Algal pigments were identified and quantified by
224	comparing their retention time and absorption spectra with those of pure standards (DHI LAB
225	products; see Majdi et al., 2011 for further details). We compared the concentration of

biomarker pigments typical of specific algal taxonomic groups to estimate potential changes 226

in the composition of the algal community (Jeffrey et al., 1997). 227

228

2.6. Statistical analyses 229

All statistical analyses were performed using the 'vegan' (Oksanen et al., 2013) and 230

'indicspecies' (Cáceres and Legendre, 2009) packages of the R software (R Development 231

232 Core Team, 2018). Changes in the pigment composition of the biofilm and the structure of 233 biofilm-dwelling invertebrate and nematode communities between heated and control sites 234 were assessed using permutational analyses of variance based on the Bray-Curtis distance 235 matrices (PERMANOVA, 9999 permutations) calculated from Hellinger-transformed abundance data. Multivariate homogeneity of group dispersion was tested using Anderson's 236 237 PERMDISP2 procedure which is a multivariate analog of Levene's test and which was met in 238 all cases. We used non-metric multidimensional scaling (nMDS) based on Hellinger-239 transformed abundance data and Bray-Curtis similarity to ordinate the species in a bi-240 dimensional space. Sample scores were then plotted as an illustrative variable in this 2D-241 space to visually assess how different the communities were between the heated and control 242 sites. We used multilevel pattern analysis (9999 permutations) to test the significance of 243 species-site group associations and derive candidate indicator species for each site (Cáceres 244 and Legendre, 2009).

245 We calculated a set of diversity indices based on the untransformed abundances of 246 each taxon in the nematode community. First, the Index of Trophic Diversity (ITD) was computed as ITD =  $\Sigma \theta^2$  with  $\theta$  being the relative contribution of each of the four functional 247 feeding guilds identified (Heip et al., 1985). This index ranges from 0.25 to 1, which 248 249 corresponds to the highest (each trophic group contributes 25%) and lowest (only one trophic 250 group present) feeding type diversity, respectively. Species richness (S), Shannon's entropy 251 (H, ln-based), Simpson's dominance (D), and Pielou's evenness (J) were calculated. 252 We used either one-way ANOVA when normality assumptions were met (tested using Shapiro-Wilk and Bartlett test), or non-parametric Kruskal-Wallis rank sum test on 253 254 untransformed data to test for differences in total abundance and biomass, pigment 255 concentration, diversity indices, the proportion of nematode feeding types, and age ratios between the heated and control sites. Regarding the temperature data and the body mass of 256

chironomid individuals, we compared cumulative density functions between sites using theKolmogorov-Smirnov test.

259

#### 260 **3. Results**

#### 261 *3.1. Abiotic background*

262 Oxygen, pH, streambed flow velocity, and water depth showed very little variations between 263 sites and along the biofilm colonisation period (Table S1). The most noticeable difference in 264 water chemistry between sites, but minor, was the slightly higher concentrations of major ions like  $Mg^{2+}$ ,  $Na^+$ ,  $Ca^{2+}$ ,  $Cl^-$ , and  $SO_4^{2-}$ , which increased water conductivity by 16.3% on 265 266 average at the heated site (Tables S1 and S2). This difference was probably due to the partial 267 evaporation of the water as it passed through the natural draft wet-cooling towers, thus 268 concentrating the ions in the water discharged upstream of the heated site. Nevertheless, the 269 relative concentration of ions were very similar across sites (Table S2).

270 At the beginning of the colonisation period (7 August 2018), the water temperature 271 was very high (> 28°C) at both sites (Fig 2A). Water temperature then progressively 272 diminished and the temperature contrast between the control and heated sites became more 273 pronounced in September (Fig. 2C). Overall, the temperature regime differed across the two 274 sites over the 56 days exposure period (total: 1344 h). The control site showed an average temperature of 23.81°C (min: 19.85°C; max: 28.85°C). The heated site showed an average 275 276 temperature of 25.32°C (min: 19.85°C; max: 30.86°C). Thus, the heated site was, on average, 1.5°C warmer than the control site during the study period (min measured  $\Delta T = -2.69$ °C; 277 278 max  $\Delta T = +4.38$ °C; Fig. 2C). When comparing the cumulative distribution of temperature 279 functions, there was a significant difference in the local thermal regime between the control and heated sites (Kolmogorov-Smirnov test,  $D^+ = 0.47$ , P < 0.001, Fig. S1). The average daily 280 281 fluctuation of water temperature (i.e., the range between the minimum and maximum water

temperature recorded each day) was also significantly higher at the heated site (on average 283 2.69°C) than at the control site (0.72°C) (Kruskal-Wallis test,  $\chi^2 = 90$ , P < 0.001) (Fig. 2B).

284

#### 285 *3.2. Biofilm pigment composition*

Biofilm pigment composition was significantly different between the heated and control sites (PERMANOVA,  $F_{1,38} = 15.8$ ,  $R^2 = 0.29$ , P < 0.001) (Fig. 3). More specifically, the biofilms showed significantly higher concentrations of chlorophyllide *a*, zeaxanthin,  $\beta$ -carotene, myxoxanthophyll, and echinenone at the heated site in comparison to the control site, while concentrations of chlorophyll *b*, violaxanthin, and lutein were significantly lower at the heated site (Table 1). The concentrations of the most abundant pigments, chlorophyll *a* and fucoxanthin, were not significantly different across sites.

293

#### 294 *3.3. Invertebrate community*

The total abundance of invertebrates was not significantly different across sites (Table 1) 295 with  $335.2 \pm 38.3$  (mean  $\pm$  SD) and  $295.9 \pm 17.3$  ind 10 cm<sup>-2</sup> at the control and heated sites, 296 297 respectively. Nematodes were numerically dominant in the invertebrate community, 298 representing 27.2% and 47.3% of all individuals found at the control and heated sites, respectively (Table 1). The structure of the invertebrate community differed significantly 299 across sites (PERMANOVA,  $F_{1,38} = 13.9$ ,  $R^2 = 0.26$ , P < 0.001) (Fig. 4). Nematodes and 300 oligochaetes were significantly more abundant at the heated site, while larvae of chironomids, 301 Corbicula, and Ephemeroptera, as well as harpacticoid copepods and their nauplii larvae, 302 303 were significantly more abundant at the control site (Table 1). No significant differences in 304 the abundances of rotifers, ostracods, and cladocers were found across sites (Table 1). The estimated total biomass of invertebrates was not significantly different across 305 sites, with  $385.8 \pm 51.4$  (mean  $\pm$  SD) and  $284.5 \pm 58.1 \mu$ gDM 10cm<sup>-2</sup> at the control and 306

heated sites, respectively (Table 1). Chironomids were strongly dominant in terms of biomass
and represented 59.1% and 86.9% of the total invertebrate biomass at the control and heated
sites, respectively. The biomasses of Ephemeroptera larvae, *Corbicula* larvae, harpacticoid
copepods, ostracods, and nauplii were significantly dampened at the heated site, whereas
nematodes showed significantly higher biomass (Table 1). Full test-statistics are provided in
Table S3.

The individual body-mass of chironomid larvae was on average 5.37 and 9.11 µgDM ind<sup>-1</sup> at the control and heated sites, respectively (Fig. 5A). Most individuals showed a bodymass < 1 µgDM (Fig. 5A). However, the cumulative distribution function of the chironomid population at the control site was significantly above that of the heated site (Kolmogorov-Smirnov test, D+ = 0.09, P = 0.045) (Fig. 5B), showing a greater proportion of relatively large chironomid individuals with a body-mass > 15 µgDM at the heated site (Fig. 5B).

320 *3.4. Nematode community* 

321 Of the 2,000 nematode individuals that have been identified, a total of 21 species have been

found, with 14 and 17 species at the control and heated sites, respectively (Table 2).

323 Monhystrella paramacrura (Meyl, 1954), Eumonhystera vulgaris (de Man, 1880), and

324 *Chromadorina bioculata* (Schultze *in* Carus, 1857) were the three dominant species at both

sites, accounting cumulatively for approximately 92 and 80% of the total community at the

326 control and heated sites, respectively (Table 2). However, the structure of the nematode

327 community differed significantly across sites (PERMANOVA,  $F_{1,38} = 24.8$ ,  $R^2 = 0.39$ , P <

- 328 0.001) (Fig. 6). The multilevel pattern analysis highlighted three species that were
- 329 significantly associated with the heated site (test statistics > 0.56, P <0.008): the two large
- 330 predatory-omnivorous *Mylonchulus lacustris* (Cobb, 1915) and *Mesodorylaimus bastiani*
- 331 (Bütschli, 1873), as well as the bacterivorous *Eumonhystera dispar* (Bastian, 1865).

332 *Mylonchulus lacustris* was absent at the control site while representing approximatively 6.3%333 of the total number of individuals in the community at the heated site. *Mesodorylaimus* 334 *bastiani* was absent at the control site while having a relative abundance of about 5.1% at the 335 heated site. Finally, *E. dispar* was > 10 times more abundant at the heated site (1.71 ind 336  $10 \text{cm}^{-2}$ ) in comparison to the control site (0.13 ind  $10 \text{cm}^{-2}$ ) (Table 2).

337 Nematode species richness was significantly higher at the heated site (on average 6.95 338 species found in each sample) than at the control site (on average 4.40 species) (Table 1). 339 Shannon's entropy (H) was also significantly higher at the heated site, while Simpson's 340 dominance (D) was lower (Table 1). Pielou's evenness (J) did not differ significantly across 341 sites (Table 1). The index of trophic diversity (ITD) was significantly lower at the heated site, 342 indicating a more homogeneous distribution between the different feeding types (Table 1). 343 Nematode communities were strongly dominated by deposit-feeders at the control site 344 (average relative abundance of 84.2%), while they represented on average only 54.2% of the 345 total abundance at the heated site. In contrast, the relative abundance of epistrate-feeders, 346 chewers, and suction-feeders was significantly higher at the heated site than at the control site 347 (Table 1, Fig. 7).

With a substantial number of individuals sampled, we could compare age ratios in the two dominant bacterivorous species, *Monhystrella paramacrura* and *Eumonhystera vulgaris*, across sites. *M. paramacrura* showed a ratio of juveniles:adults significantly lower at the heated site (0.75 and 1.20 at the heated and control sites, respectively; ANOVA,  $F_{1,38} = 5.42$ , P = 0.024). *E. vulgaris* showed a ratio gravid:non-gravid females almost 10x higher at the heated site (0.65 and 0.07 at the heated and control sites, respectively; Kruskal-Wallis,  $\chi^2 =$ 5.3, P = 0.021). Other ratios did not differ significantly across sites.

#### 355 **4. Discussion**

#### 356 *4.1. A realistic in situ warming experiment?*

357 Several studies have used naturally or artificially heated streams or river as "in situ 358 laboratories" to study the effects of water warming and climate change on aquatic organisms 359 and ecosystem functioning (e.g. Woodward et al., 2010, O'Gorman et al., 2012, Boulêtreau et 360 al., 2014, Worthington et al., 2015, Sandblom et al., 2016). In this study, we carried out an 361 experiment within the thermal plume downstream of heated effluents from a nuclear power 362 plant because we expected that all environmental factors that could influence the structure of 363 the biofilm community (e.g., light, nutrients, hydrodynamics), other than water temperature, 364 would be similar at the two experimental sites (inside or outside the thermal plume). The 365 results on the abiotic environment showed that, given our caution regarding local 366 hydrodynamics, this hypothesis was verified, and thus water temperature appeared to be the 367 major environmental determinant that differed between the heated and control sites. During 368 the experiment, the average difference in water temperature between the control and heated 369 sites was about 1.5°C, which is within the range of optimistic projections of temperature 370 increases by the end of the century (IPCC, 2014). The passage through natural draft wet-371 cooling towers implies a partial evaporation of the cooling water, thus concentrating 372 dissolved substances in the released effluent. We observed this effect also for dissolved 373 nutrients like nitrate and phosphate. However, when comparing their relative concentrations, 374 we found minor differences between sites, suggesting that the nutrient balance was similar for 375 biofilm-dwelling organisms at both sites. For example, the nitrate:phosphate ratio was 42.5 376 and 41.9 at the control and heated sites, respectively. The passage of the water through natural draft wet-cooling towers, whose efficiency primarily varies with air humidity, has 377 378 also resulted in greater daily fluctuations in water temperature. Those secondary effects 379 probably enhanced the environmental contrast found across sites. Nevertheless, they also are compatible with future climate change scenarios as the frequency of heat waves are expected 380

381 to increase in line with the rise in temperature and evaporation (Michel et al., 2019; Morrill et 382 al., 2005; Webb, 1996). It is worth noting that the experimental heated site was located within 383 the thermal plume only 20 m downstream from the outlet pipes of the nuclear power plant. 384 This area does not correspond to the regulatory downstream area (which is located a few 385 kilometres further downstream, after complete mixing between the river and the cooling 386 water released) where ecological differences potentially induced by the operation of the 387 nuclear power plant are under regulatory control, in accordance with local legislation and the 388 requirements of the Water Framework Directive.

389

#### 390 4.2. Effects of water warming on the microphytobenthic community of biofilms

391 At both sites, we found relatively high concentrations of fucoxanthin and diadinoxanthin, 392 which are two biomarker pigments for diatoms. This result is consistent with previous studies that have highlighted that epilithic biofilms of the Garonne River were strongly dominated by 393 diatoms in terms of biomass (Boulêtreau et al., 2014; Majdi et al., 2011, 2012a; Tekwani et 394 395 al., 2013). However, the absence of significant differences in fucoxanthin and diadinoxanthin 396 concentrations across sites showed that diatom biomass was not affected by the temperature 397 contrast studied. Similarly, chlorophyll *a* concentration (as a proxy of phototrophic biomass) 398 did not vary significantly across sites. However we found that the concentrations of 399 chlorophyllide a, myxoxanthophyll, and echinenone were more than twice as high at the 400 heated site in comparison to the control site. These three latter pigments are reported 401 biomarkers for cyanobacteria (Hertzberg et al., 1971; Hodgson et al., 2004; Takaichi and 402 Mochimaru, 2007). Several studies showed that warming can selectively promote 403 cyanobacterial growth in rivers (Butterwick et al., 2005; Watkinson et al., 2005). Majdi et al. 404 (2011, 2012a) and Tekwani et al. (2013) also reported an increase of the relative biomass of cyanobacteria in the plankton and in biofilms of the Garonne River during summer low-flow 405

406 periods. Further studies should focus on identifying which species of cyanobacteria are
407 favoured under warmer conditions and how such changes in the structure of the
408 microphytobenthic community could affect biofilm grazers and the ecosystem-level functions
409 supported by biofilm communities.

410

#### 411 *4.3. Effects of water warming on invertebrate communities*

We found that the total abundance of invertebrates did not differ across sites. However, the 412 413 structure of the community was different, with a relatively diverse community at the control 414 site whereas, at the heated site, it was crowded with nematodes and oligochaetes but lacked micro-crustaceans, larvae of *Corbicula*, and Ephemeroptera. Given the standard procedures 415 416 used to sample biofilm and the relatively straightforward changes observed at coarse 417 taxonomic-level, our results point out a potential to indicate water warming effects based on 418 changes in the taxonomic structure of biofilm-dwelling invertebrates. It is possible that the 419 warmer temperature and/or the greater daily fluctuation of temperature could have negatively 420 affected thermosensitive taxa such as Ephemeroptera, which need time to acclimatise to 421 critically high temperatures (Houghton and Shoup, 2014). For Corbicula larvae, Weitere et 422 al. (2009) showed that a 3°C increase in water temperature could reduce body-mass, 423 reproduction rate, and increase mortality. Such a rise in temperature occurred almost daily at 424 the heated site during the experimental period. In contrast, biofilm-dwelling nematodes and, 425 in a lesser extent, oligochaetes thrived well at the heated site. In addition to warmer average 426 temperatures, greater thermal fluctuations may also represent a stress for aquatic biota (Coulter et al., 2016; Thome et al., 2016), which could lead to the selection of species able to 427 428 cope with temperature extremes and large temperature fluctuations. 429

In our study, chironomids were the major contributors to invertebrate biomass.However, the reduction of their abundance observed at the heated site did not result in a

431 reduction in total invertebrate biomass. We further showed that this was due to the presence of chironomid individuals that were, on average, twice as large at the heated site in 432 433 comparison to the control site. Ectotherms can develop adaptive responses to warming such 434 as increased investment in reproduction, decreased body-size at sexual maturity, or higher 435 heat tolerance limits (Daufresne et al., 2009; Loisel et al., 2019; Majdi et al., 2019; Nelson et 436 al., 2017; Ohlberger, 2013; Sandblom et al., 2016; Yvon-Durocher et al., 2011). We did not 437 observe such a reduction of the size spectrum for chironomids. The potential rationales to this result are probably complex and must take into account the fact that (1) we have no 438 439 information at the species-level for chironomids, therefore we cannot rule out the hypothesis that different species assemblages at the heated site may account for the observed body-size 440 441 differences. (2) Chironomids complete only part of their life cycle in the aquatic 442 environment. For larvae destined to metamorphose and leave the aquatic environment, 443 warmer temperatures have the potential to boost larval growth and reduce the duration of the 444 larval stage, thereby hastening cohort succession (Cheney et al., 2019). The increase in the 445 size of chironomids observed in our study could therefore also result from the hastening of early summer cohorts. Temperature-boosted larvae could also emerge earlier, with potential 446 far-ranging consequences for species' ecology and biogeography. Nevertheless, this is likely 447 context- and time-dependent. Further large-scale studies examining chironomid communities 448 449 at a finer taxonomic-level are needed before envisaging the development of a bio-indicator of 450 river warming based on the shape of the size spectrum of chironomids (but see Walker et al., 451 1991 for bio-indication of past climate based on fossil chironomid species assemblages preserved in lake sediments). 452

453

454 *4.4. Changes in the structure of nematode communities* 

455 We observed that the structure of nematode communities differed across sites, both in terms 456 of species composition and in terms of feeding-type composition. There is evidence that the 457 taxonomic and feeding-type structure of freshwater nematode assemblages can shift in 458 response to broad-scale mechanisms like ecosystem productivity and disturbance regime 459 (Barbuto and Zullini, 2005; Kazemi-Dinan et al., 2014; Majdi et al., 2011; Michiels and 460 Traunspurger, 2005; Ristau et al., 2013; Höss et al., 2011). Here we observed that nematode 461 diversity was greater at the heated site, and we identified a few species that were clearly 462 associated with heated biofilms: Mylonchulus lacustris, Mesodorylaimus bastiani and 463 Eumonhystera dispar. The latter is a rather small opportunistic species, thus it made sense to find it at the heated site. However, the former two species are relatively large and it was 464 465 surprising that they thrived well at the heated site. Whether those species can be considered 466 indicators of thermal effects in river biofilms needs further studies. However, it has to be 467 noted that several freshwater nematode species have been already classified as sensitive or 468 tolerant to toxic pollution in freshwater sediments leading to the development of a "nematode 469 species at risk" index (NEMAspear; Höss et al., 2011). Among the most common species 470 dwelling river sediments, Höss et al. (2011) showed that Eumonhystera dispar was also one 471 of the most resistant to metal and organic pollutions.

Warming can affect biomass production and food web structure with mixed effects, 472 473 from body size-reductions and shortening of food chains in laboratory microbial food webs 474 (Petchey et al., 1999), to body-size increases and elongation of food chains in natural 475 geothermal streams (Woodward et al., 2016). For nematodes, we observed that the distribution between the different feeding-types was more even at the heated site than at the 476 477 control site. The community was also made up of a greater number of large-sized omnivorous 478 or predatory species at the heated site. In addition, *Chromadorina bioculata*, an algivorous 479 species typical of epilithic biofilms of the Garonne River (Majdi et al. 2011), was about five

480 times more abundant at the heated site than at the control site. These results suggest that 481 warming fostered the establishment of an assemblage of nematodes with more diverse 482 feeding-types, which would presumably play a more important role in trophic transfers in 483 comparison with the control site. Previous studies examining the biofilm ecological 484 succession in the Garonne River have shown that warmer temperatures during summer low-485 flow periods can trigger profound change in the structure of the algal and bacterial 486 communities (Leflaive et al., 2008; Lyautey et al., 2005; Tekwani et al., 2013) with 487 consequences on nutrient cycling and standing stocks available for grazers (Boulêtreau et al., 488 2014; Graba et al., 2014; Teissier et al., 2007). Free-living nematodes show complex trophic 489 behaviours including feeding-selectivity and attraction to food patches (Derycke et al., 2016; 490 Estifanos et al., 2013; Höckelmann et al., 2004; Moens et al., 2013; Schroeder et al., 2010). 491 Although more studies are needed to establish robust causalities, we may presume that the 492 presence of a more functionally diverse assemblage of nematode species could be explained 493 by at least two mechanisms: (1) A diversification of food resources that nematodes could 494 exploit in the biofilm, hence agreeing with studies showing the importance of bottom-up 495 effects at structuring nematode communities (Michiels and Traunspurger, 2005; Ristau et al., 496 2013). (2) Predation and/ or competition for resources as several studies have shown that many organisms can prey on or compete with freshwater nematodes (Majdi and 497 498 Traunspurger, 2015). A decrease in competition for biofilm basal resources (due e.g. to the 499 reduction of Ephemeroptera and Corbicula larvae) may have increased the number of trophic 500 niches available for algivorous nematodes. Furthermore, the decrease in intraguild 501 competition between omnivores and predators (as the numbers of micro-crustaceans and 502 insect larvae were dampened at the heated site) may have favoured the establishment of 503 relatively large omnivorous and predatory nematode species instead. Nematodes are extremely abundant in biofilms and their feeding type does not require specific taxonomic 504

expertise to be recognized under the microscope. Previous studies have shown that the
distribution of feeding types within freshwater nematode communities can be influenced by
key environmental drivers such as ecosystem productivity (see Traunspurger et al., 2020, and
references therein). Whether nematode feeding types may be envisaged as a straightforward
indicator of disturbance needs further confirmation, but this is promising as we showed a
more even distribution of feeding types at the heated site.

511 Not all nematode species thrived well in warmed-up biofilms. The case of 512 Monhystrella paramacrura was especially striking. Although this species was commonly 513 found at both sites, the abundance of this species was half as low at the heated site as at the 514 control site. This difference was accompanied by a much lower number of juveniles in 515 comparison to the control site. This result suggests that populations of *M. paramacrura* were 516 seemingly less successful under a warmer and more fluctuating temperature regime. This 517 response may be physiological as free-living nematodes show species-specific thermal niches 518 with associated patterns of population growth rate, age structure, egg production, or body-519 size at maturity (Majdi et al., 2019). An average temperature difference of 1.5°C across 520 experimental sites did not seem huge at first. However, Majdi et al. (2019) have evidenced 521 the existence of temperature thresholds between 25 and 30°C above which survival and 522 population growth rate of several freshwater nematode species could be severely reduced. 523 Such upper temperature tolerance thresholds were more likely to be exceeded at the heated 524 site due to an overall higher temperature and a greater amplitude of daily temperature 525 fluctuations. Over the whole 1344 h of study, water temperatures were >25°C for 229 h (i.e. 17% of time) and 814 h (i.e. 61%) at the control and heated sites, respectively. This likely 526 527 explains the reduced success of *M. paramacrura*, at the heated site. Furthermore, nematodes 528 can associate a certain temperature with conditions of food availability and thereby exhibit active attraction or repulsion to a given temperature where appropriate (Mohri et al., 2005). 529

530 *M. paramacrura* is a small species with a very slender body and a minute mouth cavity 531 without teeth. This may constrain the diet of *M. paramacrura*. On the other hand, 532 Eumonhystera vulgaris belongs to the same feeding guild as M. paramacrura, but unlike M. 533 paramacrura, E. vulgaris was very successful at the heated site showing higher abundances 534 and a much greater number of gravid females. In comparison, to M. paramacrura, E. vulgaris 535 is a bit larger showing a larger mouth cavity. Thereby, its diet could have been less limited by 536 prey size, which might be advantageous in warmed-up biofilms. Indeed, at this stage, it is 537 premature to suggest what precisely causes such a species turnover, laboratory studies 538 examining the effects of different conditions of food and temperature on single species and 539 species mixtures could help identify underlying mechanisms more accurately.

540

#### 541 **5. Conclusion**

The studied differences in thermal regime did not affect neither the biomass of biofilm-542 543 dwelling phototrophs nor the total abundance and biomass of invertebrates. However, the 544 community structure was different with a higher proportion of cyanobacteria and nematodes 545 and lower abundances of crustaceans and insects under warmer conditions. The response of 546 the dominant invertebrate groups in terms of biomass (chironomids) and abundance 547 (nematodes) was further examined, and the results suggested that the body-size structure of chironomids and the taxonomic and functional structure of nematode assemblages could be 548 549 good tools to indicate the impact of warming on river biofilm communities. The highlighted 550 differences in body-size and community structure may be explained by direct modification of 551 the metabolism and different specific tolerances to temperatures among biofilm-dwelling 552 organisms. The shifted biofilm landscape may, in return, create conditions that are beneficial 553 (or not) for the further settlement of colonists. In summary, our results suggest that water 554 warming coupled with larger daily temperature fluctuations have the potential to profoundly

affect biofilm communities. Further studies are recommended to correlate those community
shifts to potential changes in important ecosystem functions supported by river biofilms.

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#### 809 Tables

810 Table 1. Mean, standard deviation (SD), and minimum (Min) and maximum (Max) values of

811 the variables measured or indices computed at each site (N = 20 samples for each site).

812 Differences were tested using one-way ANOVA or Kruskal-Wallis rank sum test depending

813 on whether the data met the linear assumptions or not, respectively. The level of significance

814 (p-values: \* = 0.01-0.05; \*\* = 0.001-0.01; \*\*\* < 0.001) is given. Full test statistics are

815 provided in Table S3. Mean values highlighted in bold show significantly higher values at a

816 given site.

	Control site				Heated site				Test
Variables –	Mean	SD	Min	Max	Mean	SD	Min	Max	signif.
Pigment concentrations (ng.mgBiofilmDM <sup>-1</sup> )									
Chlorophyll a	254.49	42.41	21.51	713.53	240.71	23.21	66.59	418.75	
Fucoxanthin	83.72	13.10	4.49	221.34	71.59	8.95	21.78	154.10	
Chlorophyll b	25.15	5.51	1.77	119.52	7.36	1.11	1.11	18.40	***
Diadinoxanthin	13.05	2.31	1.31	39.67	10.19	1.46	2.35	24.22	
Chlorophyllide a	10.48	7.58	0.00	156.57	46.15	14.45	0.00	190.65	**
Hexanoyloxy fucoxanthin	8.44	1.45	0.00	18.65	10.99	1.28	0.29	21.91	
Zeaxanthin	7.66	1.48	0.46	25.06	13.48	1.48	3.06	24.42	*
Violaxanthin	5.78	1.24	0.45	26.40	2.27	0.20	1.01	4.61	*
Lutein	5.67	1.03	0.00	19.23	1.98	0.26	0.00	4.35	**
Pheophytin a	4.79	0.79	0.00	13.81	6.26	0.64	0.00	13.09	
Alloxanthin	1.82	0.38	0.00	5.38	2.05	0.36	0.00	5.36	
β Carotene	0.99	0.14	0.00	2.57	1.64	0.18	0.57	3.14	**
Myxoxanthophyll	0.81	0.58	0.00	10.99	1.80	0.32	0.00	4.49	***
Echinenone	0.00	0.00	0.00	0.00	0.89	0.25	0.00	3.80	***
Abundance (ind.10cm <sup>-2</sup> )									
Total invertebrates	335.22	38.30	157.17	839.54	295.87	17.34	145.23	467.52	
Nematodes	91.12	8.66	37.80	194.96	139.98	11.80	27.85	228.79	**
Rotifers	85.05	6.09	27.85	137.27	94.30	7.86	41.78	185.02	
Chironomids	60.38	6.31	1.99	123.35	39.81	4.92	6.37	97.48	*
Corbicula	55.80	16.82	0.00	326.27	3.76	1.16	0.00	17.90	***
Harpacticoids	19.70	6.09	0.00	109.42	3.68	1.00	0.00	17.90	**
Nauplii	8.06	2.14	0.00	33.82	2.19	0.54	0.00	9.95	*
Oligochaetes	7.56	2.36	0.00	33.82	9.67	1.34	0.00	21.88	*

Ephemeroptera	5.67	1.04	0.00	17.90	1.39	0.45	0.00	7.96	***
Ostracods	1.59	0.41	0.00	5.97	0.90	0.36	0.00	5.97	
Cladocers	0.30	0.16	0.00	1.99	0.20	0.13	0.00	1.99	
Biomass (µgDM.10cm <sup>-2</sup> )									
Total invertebrates	385.75	51.43	43.17	996.37	284.54	58.11	12.01	1080.08	
Chironomids	227.82	31.61	0.28	472.95	247.14	56.17	3.14	1041.08	
Ephemeroptera	79.59	16.98	0.00	294.06	16.97	5.67	0.00	92.53	***
Harpacticoids	48.19	17.02	0.00	345.32	5.69	1.94	0.00	33.99	***
Corbicula	13.57	4.09	0.00	79.36	0.91	0.28	0.00	4.35	***
Oligochaets	6.83	2.65	0.00	40.73	6.27	1.07	0.00	21.46	
Ostracods	5.23	1.36	0.00	19.62	2.94	1.18	0.00	19.62	*
Nauplii	3.06	0.82	0.00	12.67	0.82	0.20	0.00	3.73	*
Nematodes	0.94	0.11	0.21	2.20	3.28	0.38	0.36	7.16	***
Rotifers	0.41	0.03	0.13	0.63	0.45	0.04	0.19	0.85	
Cladocers	0.10	0.05	0.00	0.64	0.06	0.04	0.00	0.64	
Nematode feeding types (%)									
Deposit-feeders	84.23	2.35	54.55	100.00	54.19	2.49	28.21	75.00	***
Epistrate-feeders	12.81	2.50	0.00	45.45	33.12	2.78	10.00	61.54	***
Chewers	1.91	0.83	0.00	15.38	6.79	1.14	0.00	18.75	***
Suction-feeders	0.92	0.43	0.00	6.25	5.90	0.97	0.00	12.50	***
Nematode diversity indices									
Species richness (S)	4.40	0.24	3.00	7.00	6.95	0.32	5.00	10.00	***
Shannon's entropy (H)	1.09	0.04	0.73	1.45	1.48	0.05	0.94	1.90	***
Simpson's dominance (D)	0.40	0.02	0.27	0.61	0.30	0.02	0.19	0.54	***
Pielou's eveness (J)	0.76	0.03	0.53	0.96	0.77	0.02	0.59	0.92	
Trophic diversity (ITD)	0.75	0.03	0.50	1.00	0.44	0.02	0.30	0.59	***

Table 2. Mean (ind 10cm<sup>-2</sup>) and relative (%) abundance of biofilm-dwelling nematode
species. Abbreviations for nematode species names are provided in Table S5. Suffixes refer
to species feeding types (DF = deposit-feeders; EF = epistrate-feeders; C = chewers; SF =
suction-feeders).

Co	ontrol site		Heated site					
Species	Mean (ind.10cm <sup>-2</sup> )	%	Species	Mean (ind.10cm <sup>-2</sup> )	%			
MonParaDF	38.06	41.77	EuVulgDF	46.89	33.50			
EuVulgDF	37.90	41.59	ChroBiocEF	45.86	32.76			
ChroBiocEF	7.84	8.61	MonParaDF	18.95	13.54			
MonPaluDF	1.92	2.10	MyloLacC	8.80	6.29			
EuSimpDF	1.66	1.82	DoryBasSF	7.20	5.14			
TobStefC	1.10	1.21	EuSimpDF	2.49	1.78			
DoryStagSF	1.03	1.13	RhabAquaDF	2.00	1.43			
IroTenuC	0.42	0.46	EuDispDF	1.71	1.22			
EuPseuDF	0.38	0.42	EuPseuDF	1.52	1.08			
EthmoPratEF	0.27	0.30	RhabTerrDF	1.25	0.90			
TripGloC	0.15	0.16	TobStefC	1.16	0.83			
EuDispDF	0.13	0.15	DoryStagSF	0.68	0.49			
Mermi-Par	0.13	0.15	MonPaluDF	0.54	0.39			
PlecAquaDF	0.13	0.14	PlecAquaDF	0.36	0.26			
			EuLongDF	0.24	0.17			
			ChroVirEF	0.20	0.15			
			DoryAgiSF	0.14	0.10			

#### 841 **Figure caption**

Fig. 1. Aerial view of the study site (source: IGN 2020) and photographs of the experimentalset-up at the heated and control sites.

844

**Fig. 2. A.** Hourly water temperature recorded at the control (green line) and heated (orange line) sites over the entire experimental period (1344 h, starting on 7 August 2018 at 1 PM and ending when the samples were retrieved on 2 October 2018 at 1 PM). **B.** Boxplots of the daily fluctuation of water temperature (i.e., the range between the minimum and maximum water temperature for each day) at each site over the entire period. **C.** Hourly  $\Delta T$  (difference in water temperature between heated and control sites).

851

Fig. 3. Non-metric multidimensional scaling (nMDS) of the Bray–Curtis similarities of
pigment composition across samples (N = 40 concrete tiles). Pigment concentrations are
ordinated on the biplot. 'spider web' extremities indicate the position of the samples in the
2D-space. The labels in black (control or heated) indicate the centroid of the site they belong
to, and the labels in red indicate the centroid of each pigment concentration (pigment
abbreviations can be found in Table S5).

858

Fig. 4. Non-metric multidimensional scaling (nMDS) of the Bray–Curtis similarities of
invertebrate group abundances across samples. Group abundances are ordinated on the biplot.
'spider web' extremities indicate the position of the samples in the 2D-space. The label in
black (control or heated) indicate the centroid of the site they belong and the labels in red
indicate the centroid of each invertebrate group (group abbreviations can be found in Table
S5).

866	<b>Fig. 5. A.</b>	Chironomid body	y-mass distribution	at each site (	heated site:	dark orange.	, control
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site: green). Vertical dotted lines indicate the mean body-mass value for each site. **B.** 

868 Cumulative density functions for chironomid individual body-mass.

869

	870	<b>Fig. 6.</b> Non	-metric multidi	mensional	scaling (	(nMDS)	of the ]	Bray–Cu	ırtis sin	nilarities	of
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871 nematode species abundances across samples. Species abundances are ordinated on the

872 biplot. 'spider web' extremities indicate the position of the samples in the 2D-space. The

873 label in black (control or heated) indicate the centroid of the site they belong and the labels in

red indicate the centroid of each species (species abbreviations can be found in Table S5;

suffixes refer to species feeding type, see Table 2).

876

Fig. 7. Relative abundance of nematode feeding types across sites. Values are medians, boxesand whiskers show interquartile range.







![](_page_43_Figure_0.jpeg)

![](_page_44_Figure_0.jpeg)

## Cumulative body-mass distribution of chironomids

![](_page_44_Figure_2.jpeg)

Fig. 5

![](_page_45_Figure_0.jpeg)

![](_page_46_Figure_0.jpeg)

## In situ warming experiment in the Garonne River (France) Heated site Control site

![](_page_47_Picture_1.jpeg)

![](_page_47_Picture_2.jpeg)

## **Biofilm-dwelling communities**

![](_page_47_Picture_4.jpeg)

![](_page_47_Picture_5.jpeg)

Comparable standing stocks Different assemblages