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

### **communities**

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#### **Abstract**

Water warming, either resulting from global or local drivers, is of growing concern for aquatic environments. Changes in thermal regimes may alter the functioning and the trophic 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 food webs, examination of their responses to warming should contribute to a better 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 power plant as "in situ laboratories" of the effect of climate change and water warming. To 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 in the river and the biofilm growing on these standard substrates was sampled after two months of colonisation. The site located within the thermal plume directly downstream of the thermal effluents showed warmer average water temperature (mean ΔT of about 1.5°C over the study period) and greater daily temperature fluctuations in comparison to the control site. Algal and invertebrate biomass did not differ significantly, but the structure of communities differed markedly between sites. At the heated site, the proportion of cyanobacteria was 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 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 proportion of algivores, large-sized omnivorous and predatory species. Changes in the abundance and age structure further showed that some bacterivore species, such as *Eumonhystera vulgaris*, thrived especially well at the heated site, whereas another bacterivore, *Monhystrella paramacrura*, thrived well at the control site. In the context of

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

Meiofauna, Warming, Freshwater, Community ecology, Species distribution

#### **1. Introduction**

The question of how organisms and ecosystems respond to a rise in temperatures has attracted growing attention from the scientific community because of our need to understand and foresee the effects of the ongoing climate change on biological diversity and ecosystem functionality. In aquatic environments, spatial variability in water temperature can be natural (e.g. geothermal activity, presence of a water source) or result from direct local (e.g. 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 major determinant of changes in the diversity and resilience of aquatic biota (Steel et al., 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 the exceptional European heat wave in the summer of 2003. The recovery of the mollusc communities studied was not fully observed even eight years after the heat wave, and profound changes in community structure have persisted over the long term (Mouthon and Daufresne, 2015). Streams and rivers that are naturally or artificially heated can be considered relevant and realistic "*in situ* laboratories" to study the effects of climate change and water warming on aquatic organisms from the individual to the ecosystem level (Woodward et al., 2010). Several examples of such experiments have contributed to a better understanding of the effects of global warming on aquatic biodiversity and ecosystem 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 heating of streams in this region allowed for comparative studies between naturally heated and non-heated systems. These works have revealed strong influences of water temperature on the structure of macroinvertebrate and algal assemblages, and on ecosystem functioning. 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 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 of 5–10°C has been maintained between the inside and outside of the enclosure. This made possible the study of the physiological responses of the European perch (*Perca fluviatilis*) to water warming, highlighting the fact that global warming could outpace the adaptive capacity 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 River (France) and demonstrated that water warming affected the denitrification activity of the biofilms through changes in the composition of their bacterial community. Despite the key insights brought by these studies into the responses of aquatic ecosystems to climate change, we still lack a comprehensive understanding of the warming-induced ecological effects that can be expected in the future on more complex and realistic communities made up of different interacting biological compartments.

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

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

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

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

**2. Material and methods** 

*2.1. Study site* 

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 nuclear power plant is operated in an open loop with recirculation and is equipped with natural draft wet-cooling towers to reduce the temperature of the blowdown water. The water is withdrawn from the river for its cooling system and the blowdown water is then discharged heated at a series of outlet pipes located directly in the middle of the riverbed over a width of 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 a relatively shallow (from 0.5 to 1 m depth) riverbed consisting of clayey bedrocks with large stone slabs, cobble bars, or sand/pebbles banks depending on local hydrodynamics, with substantial coverage of several hydrophyte species (e.g., *Potamogeton* spp., *Myriophyllum spicatum* L.).

#### *2.2. Experimental design*

Forty concrete tiles (60 x 40 x 4 cm, 21.5 Kg each) were used as standard substrates for the 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 (heated site) was positioned within the thermal plume, 20 m downstream from the outlet pipes of the nuclear power plant and about 40 m from the right bank of the river (Fig. 1). The other block (control site) was positioned outside the thermal plume on the same transect 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 =  $50 \pm 2$ ) 167 cm; flow velocity =  $30 \pm 5$  cm s<sup>-1</sup>). A temperature data-logger (measurement frequency =  $5$ min) linked to a buoy was positioned 4 cm above the riverbed and fixed to a steel rod hammered in the riverbed at the centre of each block. Oxygen concentration, conductivity, and pH were recorded monthly using a portable probe (U50, HORIBA) (Table S1 in 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 of dissolved organic carbon using the combustion catalytic oxidation method (TOC-L, Shimadzu), and of anions and cations using high-performance ionic chromatography (ISC-5000+ and DX-120, Dionex).

*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 181 et al., 2009). We collected 3.14 cm<sup>2</sup> patches of biofilm directly underwater using a brushsampler (Peters et al., 2005). For invertebrates, four patches were collected at random locations on each tile, poured on 20 µm meshes, and pooled in a 50 ml centrifuge tube. Invertebrate samples were preserved in a buffered formaldehyde solution (final concentration 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 an icebox. The river discharge was very stable (low-flow conditions) during the biofilm colonisation period (i.e. from 7 August to 2 October 2018), but a major flood occurred two weeks after collecting the samples (16 October 2018), which prevented us from continuing the experiment.

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

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

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

nearest µm using the ImageJ image analysis software. Invertebrate biomass was estimated

*2.6. Statistical analyses* 

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

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

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

We calculated a set of diversity indices based on the untransformed abundances of each taxon in the nematode community. First, the Index of Trophic Diversity (ITD) was 247 computed as ITD =  $\Sigma \theta^2$  with θ being the relative contribution of each of the four functional feeding guilds identified (Heip et al., 1985). This index ranges from 0.25 to 1, which corresponds to the highest (each trophic group contributes 25%) and lowest (only one trophic group present) feeding type diversity, respectively. Species richness (S), Shannon's entropy (H, ln-based), Simpson's dominance (D), and Pielou's evenness (J) were calculated. 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 untransformed data to test for differences in total abundance and biomass, pigment 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

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

#### **3. Results**

#### *3.1. Abiotic background*

Oxygen, pH, streambed flow velocity, and water depth showed very little variations between sites and along the biofilm colonisation period (Table S1). The most noticeable difference in water chemistry between sites, but minor, was the slightly higher concentrations of major 265 ions like Mg<sup>2+</sup>, Na<sup>+</sup>, Ca<sup>2+</sup>, Cl<sup>-</sup>, and SO<sub>4</sub><sup>2-</sup>, which increased water conductivity by 16.3% on average at the heated site (Tables S1 and S2). This difference was probably due to the partial evaporation of the water as it passed through the natural draft wet-cooling towers, thus concentrating the ions in the water discharged upstream of the heated site. Nevertheless, the relative concentration of ions were very similar across sites (Table S2).

At the beginning of the colonisation period (7 August 2018), the water temperature 271 was very high ( $> 28^{\circ}$ C) at both sites (Fig 2A). Water temperature then progressively diminished and the temperature contrast between the control and heated sites became more pronounced in September (Fig. 2C). Overall, the temperature regime differed across the two 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 temperature of 25.32°C (min: 19.85°C; max: 30.86°C). Thus, the heated site was, on average, 277 1.5°C warmer than the control site during the study period (min measured  $\Delta T = -2.69$ °C; 278 max  $\Delta T = +4.38$ °C; Fig. 2C). When comparing the cumulative distribution of temperature functions, there was a significant difference in the local thermal regime between the control 280 and heated sites (Kolmogorov-Smirnov test,  $D^+ = 0.47$ , P < 0.001, Fig. S1). The average daily 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  $2.69^{\circ}$ C) than at the control site (0.72°C) (Kruskal-Wallis test,  $\chi^2 = 90$ , P < 0.001) (Fig. 2B).

#### *3.2. Biofilm pigment composition*

Biofilm pigment composition was significantly different between the heated and control sites 287 (PERMANOVA,  $F_{1,38} = 15.8$ ,  $R^2 = 0.29$ ,  $P \le 0.001$ ) (Fig. 3). More specifically, the biofilms showed significantly higher concentrations of chlorophyllide *a*, zeaxanthin, β-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.

#### *3.3. Invertebrate community*

The total abundance of invertebrates was not significantly different across sites (Table 1) 296 with  $335.2 \pm 38.3$  (mean  $\pm$  SD) and  $295.9 \pm 17.3$  ind  $10 \text{cm}^{-2}$  at the control and heated sites, respectively. Nematodes were numerically dominant in the invertebrate community, 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 300 across sites (PERMANOVA,  $F_{1,38} = 13.9$ ,  $R^2 = 0.26$ , P < 0.001) (Fig. 4). Nematodes and oligochaetes were significantly more abundant at the heated site, while larvae of chironomids, *Corbicula*, and Ephemeroptera, as well as harpacticoid copepods and their nauplii larvae, were significantly more abundant at the control site (Table 1). No significant differences in the abundances of rotifers, ostracods, and cladocers were found across sites (Table 1). The estimated total biomass of invertebrates was not significantly different across 306 sites, with  $385.8 \pm 51.4$  (mean  $\pm$  SD) and  $284.5 \pm 58.1$  µgDM  $10cm^{-2}$  at the control and

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 body-mass < 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-317 Smirnov test,  $D<sub>+</sub> = 0.09$ ,  $P = 0.045$ ) (Fig. 5B), showing a greater proportion of relatively large chironomid individuals with a body-mass > 15 μgDM at the heated site (Fig. 5B). 

*3.4. Nematode community* 

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).

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

*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

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 <

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

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

Nematode species richness was significantly higher at the heated site (on average 6.95 species found in each sample) than at the control site (on average 4.40 species) (Table 1). Shannon's entropy (H) was also significantly higher at the heated site, while Simpson's dominance (D) was lower (Table 1). Pielou's evenness (J) did not differ significantly across sites (Table 1). The index of trophic diversity (ITD) was significantly lower at the heated site, indicating a more homogeneous distribution between the different feeding types (Table 1). Nematode communities were strongly dominated by deposit-feeders at the control site (average relative abundance of 84.2%), while they represented on average only 54.2% of the total abundance at the heated site. In contrast, the relative abundance of epistrate-feeders, chewers, and suction-feeders was significantly higher at the heated site than at the control site (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 351 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 353 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.

#### **4. Discussion**

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

Several studies have used naturally or artificially heated streams or river as "*in situ* laboratories" to study the effects of water warming and climate change on aquatic organisms and ecosystem functioning (e.g. Woodward et al., 2010, O'Gorman et al., 2012, Boulêtreau et al., 2014, Worthington et al., 2015, Sandblom et al., 2016). In this study, we carried out an experiment within the thermal plume downstream of heated effluents from a nuclear power plant because we expected that all environmental factors that could influence the structure of the biofilm community (e.g., light, nutrients, hydrodynamics), other than water temperature, would be similar at the two experimental sites (inside or outside the thermal plume). The results on the abiotic environment showed that, given our caution regarding local hydrodynamics, this hypothesis was verified, and thus water temperature appeared to be the major environmental determinant that differed between the heated and control sites. During the experiment, the average difference in water temperature between the control and heated sites was about 1.5°C, which is within the range of optimistic projections of temperature increases by the end of the century (IPCC, 2014). The passage through natural draft wet-cooling towers implies a partial evaporation of the cooling water, thus concentrating dissolved substances in the released effluent. We observed this effect also for dissolved nutrients like nitrate and phosphate. However, when comparing their relative concentrations, we found minor differences between sites, suggesting that the nutrient balance was similar for biofilm-dwelling organisms at both sites. For example, the nitrate:phosphate ratio was 42.5 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 also resulted in greater daily fluctuations in water temperature. Those secondary effects 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

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

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

At both sites, we found relatively high concentrations of fucoxanthin and diadinoxanthin, 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 diatoms in terms of biomass (Boulêtreau et al., 2014; Majdi et al., 2011, 2012a; Tekwani et al., 2013). However, the absence of significant differences in fucoxanthin and diadinoxanthin concentrations across sites showed that diatom biomass was not affected by the temperature contrast studied. Similarly, chlorophyll *a* concentration (as a proxy of phototrophic biomass) did not vary significantly across sites. However we found that the concentrations of chlorophyllide *a*, myxoxanthophyll, and echinenone were more than twice as high at the heated site in comparison to the control site. These three latter pigments are reported biomarkers for cyanobacteria (Hertzberg et al., 1971; Hodgson et al., 2004; Takaichi and Mochimaru, 2007). Several studies showed that warming can selectively promote cyanobacterial growth in rivers (Butterwick et al., 2005; Watkinson et al., 2005). Majdi et al. (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

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

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

We found that the total abundance of invertebrates did not differ across sites. However, the structure of the community was different, with a relatively diverse community at the control 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 used to sample biofilm and the relatively straightforward changes observed at coarse taxonomic-level, our results point out a potential to indicate water warming effects based on changes in the taxonomic structure of biofilm-dwelling invertebrates. It is possible that the warmer temperature and/or the greater daily fluctuation of temperature could have negatively affected thermosensitive taxa such as Ephemeroptera, which need time to acclimatise to critically high temperatures (Houghton and Shoup, 2014). For *Corbicula* larvae, Weitere et 422 al. (2009) showed that a  $3^{\circ}$ C increase in water temperature could reduce body-mass, reproduction rate, and increase mortality. Such a rise in temperature occurred almost daily at the heated site during the experimental period. In contrast, biofilm-dwelling nematodes and, in a lesser extent, oligochaetes thrived well at the heated site. In addition to warmer average 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 cope with temperature extremes and large temperature fluctuations. 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

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 comparison to the control site. Ectotherms can develop adaptive responses to warming such as increased investment in reproduction, decreased body-size at sexual maturity, or higher heat tolerance limits (Daufresne et al., 2009; Loisel et al., 2019; Majdi et al., 2019; Nelson et al., 2017; Ohlberger, 2013; Sandblom et al., 2016; Yvon-Durocher et al., 2011). We did not 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 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 differences. (2) Chironomids complete only part of their life cycle in the aquatic environment. For larvae destined to metamorphose and leave the aquatic environment, warmer temperatures have the potential to boost larval growth and reduce the duration of the larval stage, thereby hastening cohort succession (Cheney et al., 2019). The increase in the 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 far-ranging consequences for species' ecology and biogeography. Nevertheless, this is likely context- and time-dependent. Further large-scale studies examining chironomid communities at a finer taxonomic-level are needed before envisaging the development of a bio-indicator of river warming based on the shape of the size spectrum of chironomids (but see Walker et al., 1991 for bio-indication of past climate based on fossil chironomid species assemblages preserved in lake sediments).

*4.4. Changes in the structure of nematode communities* 

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

Warming can affect biomass production and food web structure with mixed effects, from body size-reductions and shortening of food chains in laboratory microbial food webs (Petchey et al., 1999), to body-size increases and elongation of food chains in natural 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 control site. The community was also made up of a greater number of large-sized omnivorous or predatory species at the heated site. In addition, *Chromadorina bioculata,* an algivorous species typical of epilithic biofilms of the Garonne River (Majdi et al. 2011), was about five

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

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.

Not all nematode species thrived well in warmed-up biofilms. The case of *Monhystrella paramacrura* was especially striking. Although this species was commonly found at both sites, the abundance of this species was half as low at the heated site as at the control site. This difference was accompanied by a much lower number of juveniles in comparison to the control site. This result suggests that populations of *M. paramacrura* were seemingly less successful under a warmer and more fluctuating temperature regime. This response may be physiological as free-living nematodes show species-specific thermal niches with associated patterns of population growth rate, age structure, egg production, or body-size at maturity (Majdi et al., 2019). An average temperature difference of 1.5°C across 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 population growth rate of several freshwater nematode species could be severely reduced. Such upper temperature tolerance thresholds were more likely to be exceeded at the heated 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  $>25^{\circ}$ C for 229 h (i.e. 526 17% of time) and 814 h (i.e. 61%) at the control and heated sites, respectively. This likely explains the reduced success of *M. paramacrura*, at the heated site. Furthermore, nematodes 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).

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

#### **5. Conclusion**

The studied differences in thermal regime did not affect neither the biomass of biofilm-dwelling phototrophs nor the total abundance and biomass of invertebrates. However, the community structure was different with a higher proportion of cyanobacteria and nematodes and lower abundances of crustaceans and insects under warmer conditions. The response of the dominant invertebrate groups in terms of biomass (chironomids) and abundance (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 good tools to indicate the impact of warming on river biofilm communities. The highlighted differences in body-size and community structure may be explained by direct modification of the metabolism and different specific tolerances to temperatures among biofilm-dwelling organisms. The shifted biofilm landscape may, in return, create conditions that are beneficial (or not) for the further settlement of colonists. In summary, our results suggest that water 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.

#### **6. Aknowledgements**

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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.
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**Table 2.** Mean (ind  $10 \text{cm}^{-2}$ ) and relative (%) abundance of biofilm-dwelling nematode 830 species. Abbreviations for nematode species names are provided in Table S5. Suffixes refer 831 to species feeding types (DF = deposit-feeders;  $EF$  = epistrate-feeders;  $C$  = chewers;  $SF$  = 832 suction-feeders).

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#### **Figure caption**

**Fig. 1.** Aerial view of the study site (source: IGN 2020) and photographs of the experimental 843 set-up at the heated and control sites.

**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 ΔT (difference in water temperature between heated and control sites).

**Fig. 3.** Non-metric multidimensional scaling (nMDS) of the Bray–Curtis similarities of 853 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).

**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).

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

Cumulative density functions for chironomid individual body-mass.

- **Fig. 6.** Non-metric multidimensional scaling (nMDS) of the Bray–Curtis similarities of
- nematode species abundances across samples. Species 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
- 874 red indicate the centroid of each species (species abbreviations can be found in Table S5;
- suffixes refer to species feeding type, see Table 2).
- 
- **Fig. 7.** Relative abundance of nematode feeding types across sites. Values are medians, boxes 878 and whiskers show interquartile range.











## **Cumulative body-mass distribution of chironomids**



Fig. 5





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





## Biofilm-dwelling communities





Comparable standing stocks Different assemblages