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#### 1 Meroplanktic phytoplankton play a crucial role in responding to peak discharge events in the

- 2 middle lowland section of the Loire River (France)
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#### 16 Abstract

- 17 Meteorological and hydrological forcings influence phytoplankton at very short time scales. The effect
- 18 of turbulence, dilution, light and nutrients are highly dynamic. Yet, our knowledge of short-term

19 phytoplankton dynamics associated with discharge disturbances and nutrient inputs remains elusive,

- 20 especially in large rivers.
- 21 Based on every three-day monitoring, we studied phytoplankton in the middle Loire River (France)
- and related to the daily variations in water discharge and the physical and chemical parameters. We
- 23 focused on summer phytoplankton (2013 and 2014), where dissolved inorganic phosphorus
- 24 concentration was potentially limiting growth. We identified eight discharge events, which increased
- suspended matter concentration and decreased chlorophyll-a concentration. The most significant

26 environmental drivers of phytoplankton composition were discharge and water temperature, a

27 sensitive proxy for meteorological forcing at short-time scale.

28 The phytoplankton composition responded to changes in hydrology along with three distinct

assemblage types, where even small water discharge increase induced a community response.

30 Meroplanktic algae being able to withstand sedimentation and resuspension could take advantage of

31 hydrological peaks, following the benthic retention hypothesis. Our results suggest that short-term

32 dynamics are crucial to understanding community organisation and functioning in large river plankton,

33 with meroplankton playing a decisive role in maintaining phytoplankton diversity and ecosystem

34 functioning.

**Keywords:** large rivers; short-term events; benthic retention hypothesis; Reynolds' phytoplankton

36 functional groups; functional traits

37

#### 38 Introduction

Large rivers are critical players in the biogeochemical dynamics, exerting control over the transport of 39 organic matter and associated chemical elements along the land-sea continuum. Large spatial and 40 41 temporal variations occur in hydrology and biogeochemical processes in rivers, which also influence the diversity of aquatic communities (Johnson et al., 1995). Large rivers exceeds the sixth river order 42 (Wehr & Descy, 1998) with extended braided arms towards the lowland river section. Their hydrology 43 44 is subjected to significant variations in discharge in response to meteorological variations and water 45 flow management. Water discharge also controls connectivity among adjacent pools composed by water 46 bodies with different water quality. The longitudinal component from the headwaters to the lowland 47 reach can be described by a continuous change in environmental conditions, well summarised in the 48 'River Continuum Concept - RCC' (Vannote et al., 1980). The RCC also provides a natural gradient to 49 scale the diversity of aquatic communities. The RCC describes changes in river metabolism, in the type of primary producers and the macro-invertebrates and fishes' community structure, associated with 50 changes in the slope (current velocity), allochthonous inputs, light exposure and water residence time. 51 The large lowland river sections are influenced by upstream processes and aquatic communities, with 52

53 phytoplankton dominance mainly from the middle river section (Reynolds & Descy, 1996). Temporal fluctuations associated with river discharge should affect the taxonomic and functional diversity of 54 55 aquatic communities, by connecting benthic and planktic communities, in accordance with the benthic retention hypothesis (Istvànovics & Honti, 2011). This latter suggests that fast-sinking meroplanktic 56 diatoms can take advantage of sedimentation and resuspension if the retention time is sufficiently short 57 and can compensate against light deficient conditions (Istvànovics & Honti, 2011). How this hypothesis 58 59 applies to other taxonomic groups, such as coccal green algae that may be favoured during summer low 60 flow conditions, remains unclear.

Phytoplankton play a crucial role in the ecology and functioning of large rivers, reproducing within river 61 62 (Wu et al., 2011) and exhibiting a rapid response time to environmental conditions ranging from a few hours to a few days. Phytoplankton biomass and composition are controlled by a set of biotic and abiotic 63 factors, such as flushing rate (Reynolds et al., 1994), turbulence (Bormans et al., 2004), light availability 64 into the water column (linked to atmospheric conditions, turbidity and river depth) (Várbíró et al., 2018), 65 water temperature (Peterson & Stevenson, 1989), nutrient availability (Minaudo et al., 2021), 66 67 competition for nutrients with macrophytes (Sabater et al., 2008), and grazing by zooplankton (Lair & Reves-Marchant, 1997). Studies often identify and rank controlling factors at a seasonal time scale, 68 69 focusing on hydrological forcing with high versus low-flow periods, on bottom-up aspects (seasonal 70 changes in resource availability) or, more exceptionally on top-down control (Picard & Lair, 2005). 71 While the seasonal and longitudinal dynamics of phytoplankton have been well studied in large rivers (Reynolds & Descy, 1996; Istvánovics et al., 2010), short-term dynamics associated with discharge 72 forcing, have mainly been neglected. Biological responses, in terms of growth and biomass, are known 73 (Kiss et al., 1996), and are increasingly studied based on high-frequency data (Bowes et al., 2016; Pathak 74 75 et al., 2021). However, much less has been derived in terms of community responses (Dubelaar et al., 76 2004).

Changes in phytoplankton assemblages result from two key interacting processes (Cloern, 1996; Loreau & Mouquet, 1999; Hiltunen et al., 2006): (1) the allogenic processes (also called larger-scale or regional processes), with physical transport, mixing and water advection, which mix populations of different origins and waters with different chemical compositions; (2) the autogenic processes (also called local

processes), with biological processes such as species-dependent growth and losses due to competition, predation or response to abiotic changes. Temporal response to allogenic processes is considered immediate whereas autogenic processes need time to produce a visible response in the ecosystem (Reynolds, 1984; Cloern, 1996; Pannard et al., 2008b).

85 In lakes, it is known that short-term dynamic of phytoplankton is mainly controlled by wind forcing (Carrick et al., 1993; Reynolds, 2006; Winder et al., 2008). Wind-induced mixing controls the physical 86 87 and chemical environment (resource availability, stratification intensity, deepening on the mixed layer) and the development of species adapted to well-mixed conditions (Pannard et al., 2008a). In temperate 88 89 lakes, the short-term dynamic nests into the seasonal dynamic, leading to a new community structure 90 without any return to the previous one, because of constantly changing environmental conditions 91 (thermocline depth or temperature, for instance). Such 'resetting events' are more common in tropical 92 lakes (Lewis, 1978).

93 In coastal areas, the short-term dynamic of phytoplankton is also controlled by mixing events with 94 estuarine waters due to windy circumstances, river flow and tidal dynamics (Lohrenz et al., 2003). 95 Mixing events lead to an abrupt change in the total phytoplankton biomass and concentration of 96 dominant species with irregularly high growth rates (Pannard et al., 2008b). Consequently, varying 97 intensity and frequency in physical forcing help maintain high phytoplankton diversity in lakes and 98 coastal waters by mixing populations, suppressing dominance and favouring rare species.

99 Water discharge can swiftly regulate autogenic and allogenic processes in river phytoplankton. An elevation in river discharge can alter environmental conditions such as turbulence and turbidity, while 100 101 facilitating the connection of neighbouring water bodies with distinct nutrient loads and communities, 102 including the benthic-pelagic interface (Istvànovics & Honti, 2011). During flood events, phytoplankton 103 is enriched by meroplankton, which refers to benthic and epiphytic species that have been resuspended, 104 and species from adjacent aquatic ecosystems reconnected by hydrological forcing. Rojo et al. (1994) 105 have shown that meroplankton can represent a significant fraction of the total phytoplankton biomass 106 (up to 50% of suspended taxa). Benthic and epiphytic taxa observed in phytoplankton should be a good proxy of river discharge peak events. Mixing-induced meroplankton occurrence should increase 107 108 diversity and at the same time may help maintain rare species.

109 River discharge is a crucial driver of phytoplankton dynamics and should promote diversity maintenance 110 by i) transporting new species from adjacent river pools and benthic areas, ii) diluting the resident 111 community, dampening the competitive exclusion by dominant species and therefore sustaining rare 112 species, iii) changing light by increasing turbidity, thus limiting growth except for shade-adapted species, iv) increasing nutrient availability, either directly through nutrient inputs from the catchment 113 area, or indirectly by reducing the boundary layer around large cells through turbulence (Barton et al., 114 115 2014) and by reducing the competition for nutrients when dominant species are suppressed. While nutrient pulses favour fast-growing species, large cells might take advantage of these short-term events, 116 thanks to nutrient storage. Thus, species-specific characteristics (i.e. traits) should primarily reflect the 117 species- and community-level responses to discharge-mediated disturbances. 118

119 Functional traits are morphological, physiological, or phenological features - measurable at the individual level - affecting fitness (Violle et al., 2007). Measuring mechanistic traits at the individual 120 (and even species) level in natural communities is rather impossible. Morphological traits that reflect 121 122 mechanistic traits are traditionally used in phytoplankton ecology (Litchman & Klausmeier, 2008), and 123 often result in the use of some key traits, such as cell size and organization. Morphological traits also 124 enable classifying phytoplankton taxa into ecological groups (Salmaso et al., 2015), which can help 125 better relate ecosystem functioning to community diversity (Abonyi et al., 2018a). Grouping species 126 into ecological, often called 'functional groups', is increasingly adopted by ecologists (Nock et al., 127 2016). A frequently used functional approach is the classification sensu Reynolds (FG) based on habitat templates (Reynolds et al., 2002), which has been advocated as a trait-based approach (Kruk et al., 128 2021); although we still need to get a mechanistic understanding to link ecosystem-level processes to 129 130 community composition based on FGs (Abonyi et al., 2021).

At the scale of hydrodynamic events, short and long responses can be expected with a modification in phytoplankton species composition and dominance, and a corresponding alteration in functional trait composition. Short-term dynamics in water discharge should promote larger biodiversity in river phytoplankton, partly explained by meroplankton. A bi-monthly sampling in the Moselle river already demonstrated that diversity and the composition of phytoplankton assemblages are clearly linked to river discharge (Descy, 1993).

137 The Loire River is one of the four largest Atlantic rivers with 200-300 m of width in its lower sections and about 1,012 km of length from its source to the estuarine zone (Moatar et al., 2022). Phytoplankton 138 139 from the Loire have been widely studied since 1920, but its short-term dynamics have never been shown. 140 Many efforts have been conducted since 1991 to limit anthropogenic phosphorus point sources in the hydrosystem to counteract eutrophication which was reaching some extreme levels in the 1980s with 141 chlorophyll-a concentration often over 250 µg L<sup>-1</sup> (Moatar & Meybeck, 2005; Minaudo et al., 2015). 142 143 Over time, the phosphorus concentration dropped by a factor of three and the Loire phytoplankton now seem to be highly P-limited in late summer (Oudin et al., 2009; Floury et al., 2012; Minaudo et al., 2015, 144 2018, 2021). Longitudinal changes in functional composition, coinciding with the main hydro-145 ecoregions, have been highlighted, and associated with regional differences in the N:P and Si:P ratios 146 147 (Abonyi et al., 2014). The Loire River phytoplankton have three distinct assemblage types, upstream detached benthic pennate diatoms, planktic centric diatoms in the middle section (in accordance with an 148 increase in water residence time) and coccal green algae together with small centric diatoms downstream 149 150 (in accordance with shallowness and increased light availability) (Abonyi et al., 2012; Descy et al., 151 2012). The Middle Loire River (465 to 772 river km from the headwaters) combines most of the conditions that may favour both pelagic and benthic growth: high light availability with low water level 152 153 in summer ( $\approx$ 1m), high water temperature (Seyedhashemi et al., 2022) and a river morphology with 154 multiple channels, adjacent pools, and numerous islands, slowing down the flow velocity (Latapie et al., 155 2014). The Middle Loire receives 40% of the total basin contribution, with very little amount of lateral water inputs, making this section the most representative of the large lowland Loire River. Based on the 156 presence of true planktic species (centric diatoms), one can say that autogenic process dominates from 157 158 the Middle Loire, which makes it an excellent site to study short-term dynamics. A single station was 159 chosen in the Middle Loire and its short-term temporal dynamics were analysed over two hydrological 160 summers in 2013 and 2014. Based on an every 3-day sampling frequency, the short-term variations in 161 river discharge were related to the variations in both local physical and chemical parameters and 162 phytoplankton assemblages.

163 The objective of this study was (1) to highlight the changes in physical, chemical and biological164 parameters associated with water discharge increase (even in low amplitude), (2) to identify short-term

response of phytoplankton assemblage and characterize key taxa and their traits, (3) to highlight a potential effect of repeated disturbances on phytoplankton assemblage, and (4) to evaluate the role of meroplanktic algae on phytoplankton diversity in connection with the benthic retention hypothesis. Studies combining the monitoring of water quality variables (nutrients, physical and chemical parameters) with community data on phytoplankton at a near-daily scale remain scarce.

170

#### 171 Method

#### 172 Study area and sampling location

The geographical and physical characteristics of the Loire River basin have formerly been described extensively in the literature (Descy et al., 2009; Moatar et al., 2022). The sampling station, located at Cinq-Mars-La-Pile near Tours in France, is considered representative of the Middle-Loire (Minaudo et al., 2015). It covers 40 % (43.6 x10<sup>3</sup> km<sup>2</sup>) of the total basin contribution with an annual mean river discharge ~355 m<sup>3</sup> s<sup>-1</sup>. The Loire River does not receive any significant inflow over 272 km upstream before arriving to Cinq-Mars-La-Pile, except some groundwater inputs from the Beauce aquifers under high agricultural pressure (Lalot et al., 2015).

#### 180 Daily river discharge and physical and chemical data

Daily river discharge (Q) at gauging station was downloaded from the online open national water monitoring database (http://www.hydro.eaufrance.fr/). Water temperature, pH, dissolved oxygen and conductivity were monitored hourly by Electricité de France (EDF), upstream of the Avoine nuclear power plant, 20 km downstream from our sampling site. Global solar radiation originated from SAFRAN dataset re-analysis (Quintana-Segui et al., 2008), and were averaged over the previous week to account for prior conditions that might had impacted phytoplankton at a given date (Várbíró et al., 2018).

#### 188 Sampling design

Water was sampled at the thalweg from a bridge in the main river channel. Total suspended solid concentrations (SM) were measured every day. The following parameters were analyzed on a 3-day frequency basis from November 2011 to October 2014: dissolved and particulate organic carbon (DOC and POC), total and dissolved inorganic phosphorus (TP and DIP), total nitrogen (TN), total dissolved

193 nitrogen (TDN) and nitrates (N-NO<sub>3</sub>), dissolved silicon (DSi), chlorophyll-a and pheopigment 194 concentrations. Filtrations were immediately made on-site with 0.45 µm cellulose acetate membrane 195 filters for chemical parameters and with 0.70 µm glass filter (Whatman GFF) previously burned at 196 500°C during 6 hours for chlorophyll-a and pheopigment. SM was determined by filtration of a precise volume of each water sample through pre-weighed filters and by drying them at 105°C. After filtration, 197 198 water samples and filters were stored at -80°C in polypropylene tubes after acidification of aliquots for 199 nitrates, DIP and DOC analysis. Tubes and filters were unfrozen the day of the analysis. Dissolved 200 organic carbon, TN and TDN concentrations were measured with a carbon and nitrogen analyzer (Shimadzu TOC-V CSH/CSN with TN unit) according to standard protocols from manufacturer 201 (Shimadzu TOC-VCSH/CSN User Manual). Nitrates concentration was determined by ionic 202 203 chromatography. Phosphorus was measured by colorimetry after solid digestion (potassium-persulfate digestion) when measuring TP (Grasshoff et al., 1983). Dissolved silicon was measured by colorimetry 204 205 (Aminot & Kérouel, 2004). For POC analyses, the filters were first treated with HCl 2N to remove carbonates, dried at 60°C for 24 hours and then measured with a C/S analyzer (LECO C-S 200). 206 207 Chlorophyll-a (Chl-a) and pheopigment were measured by fluorimetry at a wavelength > 665 nm after an excitation step between 340 and 550 nm (Strickland & Parsons, 1972). To estimate living 208 209 phytoplankton and meroplankton in terms photosynthetic activity, we calculated the particulate organic 210 carbon to chlorophyll-a ratio (POC:chl-a) for each sample. It is used as an indicator for ecological and physiological state of phytoplankton, with low ratio when particulate organic matter is dominated by 211 living phytoplankton and high values when there is a lot of organic dead materials (Bibi et al., 2020). 212

Phytoplankton were collected from April to October, with 12 samples in 2012, 44 in 2013 and 44 in 213 2014. Phytoplankton community composition was only related to hydrology in summers 2013 and 2014. 214 215 where samples were taken with an even distribution (i.e. every 3 days). Summers 2013 and 2014 216 represented ~20 samples in total for each year. Phytoplankton samples were fixed on-site after sampling 217 with Lugol solution. Phytoplankton species were identified with an inverted microscope according to 218 the Utermöhl method (Utermöhl, 1958). Phytoplankton manuals for species identification were Geitler (1930-1932), Huber-Pestalozzi (1955), Fott (1968), Ettl et al. (1978, 1985), Komárek and Fott (1983), 219 220 Starmach (1985), Popovský and Pfiester (1990), Komárek and Anagnostidis (1999, 2005). If centric

221 diatom dominance appeared, we prepared permanent slides following the Comité Européen de Normalisation (2003) standard. Diatoms were identified using Krammer and Lange-Bertalot (1986, 222 223 1988, 1991a, 1991b, Lange-Bertalot (2001) and Krammer (2002). The phytoplankton concentration was 224 first approximated in cells mL<sup>-1</sup>, and then expressed as wet-weight biomass (mg L<sup>-1</sup>) using speciesspecific biovolumes obtained by geometrical approximation according to (Lund & Talling, 1957; Rott, 225 1981). We converted wet weight biomass into carbon biomass (mgC  $L^{-1}$ ) based on Eppley et al. (1970). 226 227 Phytoplankton species were assigned to Reynolds functional groups (RFG) according to Borics et al. (Borics et al., 2007), Padisák et al. (2009), and Abonyi et al. (2021). We quantified the dominance 228 relationships based on the inversed Simpson using the *diversity* function of the 'vegan' library in R 229 230 (Oksanen, 2013).

#### 231 Identification of hydrological events and associated periods

During monitoring of river discharge, a river flow peak was identified when the daily river flow F increased rapidly over a few days, *i.e.* when the slope dF/dt became positive between two consecutive days. This is noticed as \* on Fig. 1. The use of the daily average flow rate allows to smooth the curve, but it is possible to perform a smoothing over a few days if the inter-day variations are too big. Minor peaks (even small as 5 m<sup>3</sup> s<sup>-1</sup> day<sup>-1</sup>) were also identified when a slight increase in flow (even over two days) was coupled with changes in biochemical parameters such as POC:chl-a ratio.

To highlight change in phytoplankton assemblage associated with hydrological events, we identified

239 different periods according to their timing before and after peak flow events (Fig. 1). The 'After'

240 period begins when water discharge start increasing (t\*). 'Before' and 'After' thus correspond to the

241 increase and the beginning of the decrease in water discharge, respectively. It ends at the end of the

recession curve of the river flow at t<sub>2</sub> (Fig. 1). We termed periods outside 'Before' and 'After'

243 'Stable'.

244 We then tested if these periods corresponded to distinct phytoplankton assemblages through

245 multivariate analyses, for each year independently. Three assemblages per year were highlighted,

called 'stable' assemblages (during stable periods), 'disturbed' assemblages (from t\* to t<sub>2</sub>) and

247 'repeatedly disturbed' assemblages (at least two successive disturbed periods).

248 Multivariate analysis

9

249 About 150 phytoplankton species have been observed with many rare species. To characterize the short-250 term relationships between environmental variables (explanatory variables) and taxa biomass (response 251 matrix), a canonical correspondence analysis (CCA) (Legendre & Anderson, 1999) has been performed 252 for each year, using 'ade4' and 'vegan' R libraries (Oksanen, 2013). This method is based on both multiple regression and ordination methods. Prior to these canonical analysis, phytoplankton (biomass) 253 254 community data were Hellinger-transformed to remove the effects of both rare (many zeros) and 255 dominant taxa (Legendre & Gallagher, 2001). The environmental parameters were centered and scaled and collinearity was checked. The link between explanatory and responses variables was tested through 256 a Monte Carlo permutation test with 999 replications (Blanchet et al., 2008) using CCA. CCA was used 257 to identify characteristic phytoplankton assemblages associated with hydrological forcing and to identify 258 259 environmental parameters. A taxa contributes to a canonical axis if its variability in relative frequency among samples contributes to the variability of the assemblage. 260

A correspondence analysis (CA) has also been performed on Hellinger-transformed RFG biomass. A discriminant analysis (DA) was performed on the CA with the function *discrimin* to identify RFG associated with 'stable', 'disturbed' and 'repeatedly disturbed' assemblages. Instead of maximizing the total variance carried by axes 1 and 2 as in a classical CA or CCA, the analysis focused on maximizing the explained inter-group variance depending on a discriminating factor. The level of significance of the discriminating factor has been tested by a Monte Carlo permutation test (based on 999 permutations). All analyses were performed with 'ade4' library in Rstudio.

To link individually taxa and RFG with environmental parameters, a Pearson's correlation matrix wasperformed with the R library 'corrplot' (Wei & Simko, 2017).

270

#### 271 **Results**

#### 272 Overall variability of hydrology, physical and chemical parameters and phytoplankton

#### 273 assemblage among years

274 Hydrology was contrasting among studied years, with water discharges ranging from 50 to 1,900 m<sup>3</sup> s<sup>-</sup>

275 <sup>1</sup> (Fig. S1), and up to 3,580 m<sup>3</sup> s<sup>-1</sup> downstream. The winter high-flow periods in 2013 and 2014 were

wet, with a seasonal maximum of 1900 and 1400 m<sup>3</sup> s<sup>-1</sup>, respectively. Minimum summer discharge 276 was 110 m<sup>3</sup> s<sup>-1</sup> in 2013 and 90 m<sup>3</sup> s<sup>-1</sup> in 2014. Total suspended solids (SM) concentrations ranged 277 between 0.2 and 137 mg L<sup>-1</sup>. SM was in general higher in winter due to major hydrological events, 278 279 though it was also relatively high ( $\approx 50 \text{ mg L}^{-1}$ ) in summer during hydrological events. Nitrates concentration was highly seasonal and synchronous with discharge. DSi showed the lowest values 280 between April and June with 1.4 and 1.1 mg  $SiO_2 L^{-1}$  in 2013 and 2014, while P-DIP was lowest in 281 July and August, with less than 1  $\mu$ g P L<sup>-1</sup> (Fig. S1). Chlorophyll-a values indicated some extensive 282 phytoplankton growth periods, with similar maximum values every year near 60 µg L<sup>-1</sup>. Centric 283 284 diatoms systematically dominated the phytoplankton assemblage in spring and then decreased in dominance towards summer, when green algae started to grow (Fig. S1). Phytoplankton at the end of 285 286 the summer were either dominated by green algae (2012) or diatoms (2013 and 2014).

#### 287 Physical and chemical parameters in summer 2013

From early July to early September 2013, three discharge peak events with increasing intensity were 288 observed, more than eight days apart (events number 1 to 3, Fig. 2a). SM content increased 289 290 simultaneously with river discharge, each time with an increasing intensity, consistent with the 291 dynamic of river discharge (Fig. 2b). The third event resulted in a sharp increase in the POC:chl-a ratio of SM in the river, which increased by a factor of 9, whereas this ratio did not fluctuate during 292 the first two events (Fig. 2a). This sharp increase was partly related to the drop of chlorophyll-a 293 concentration in the water (Fig. 2e). Three days after this event, DIP exceeded 40  $\mu$ g P L<sup>-1</sup> (Fig. 2c). 294 On 19/08/2013, six days after this third discharge event, DIP decreased to near pre-disturbance values 295  $(3 \mu g P L^{-1})$ , while chlorophyll-a increased to its pre-disturbance maximum value (44.2  $\mu g L^{-1}$ ). It is 296 297 noteworthy that overall, DIP and chlorophyll-a concentration correlated negatively (Pearson's correlation, r = -0.57; p = 0.003), indicating DIP consumption when phytoplankton biomass was high. 298 299 DOC increased steadily over 9 days following the third discharge event (Fig. 2d). A slight increase in 300 DOC was also observed on 04/08/2013, following event 2. Phaeopigment concentration remained 301 unchanged by the discharge events.

- A minor increase of river discharge could also be observed at the end of the period, on 03/09/2013
- 303 (number 4, Fig. 2a). This event doubled the POC:chl-a ratio of suspended matter in the river and led to

11

a 50% increase in DOC (Fig. 2d). Two 'stable' periods (outside the before/after periods) could be
identified during the summer 2013.

306 The inverse Simpson index increased during stable hydrology periods, and decreased during peak

discharge events (Fig. 2f). The strongest event 3 resulted in the minimum value of the index, with 2.3

308 on 19/08/2013.

309 Short-term responses of phytoplankton to hydrology in summer 2013

310 To characterize the changes in assemblages and relate them to physical and chemical parameters

311 (temperature, pH, oxygen, nutrients, discharge, DOC), a CCA was performed on the biovolumes data

312 (Fig. 3). Environmental variables explained 62.7% of the total variance in phytoplankton community.

313 The sampling dates grouped together following the hydrological periods specified above (Fig. 2).

314 While some periods showed overlaps (e.g. Before1, After3 and Stable2), others remained highly

separated along with axis 1 and axis 2 (Fig. 3c).

Along axis 1 of the CCA the consecutive periods Stable1 and Before2, on the right of the axis,

317 contrasted with the initial assembly (Before1) and the ones after event 3 (After3, Stable2 and After4).

318 After1 and After2 had similar 'mean' assemblages, i.e. they were located in the centre of the

ordination. Temperature was the most significant variable correlating positively with axis 1 (r = 0.87),

while nitrates (r = -0.53), DIP (r = -0.41) and DOC (r = -0.43) correlated negatively along with the

321 same gradient (Fig. 3b). Considering the weight of these parameters in the prediction of taxa biomass

322 (i.e. the canonical coefficients of environmental parameters in the CCA), temperature changes had a

weight more than twice that of nitrates (-0.17 vs. + 0.40) and four times that DIP (-0.09 vs. + 0.4).

324 Stable1 and Before2 were characterized by higher water temperatures, but lower concentrations of

nitrates, DIP and DOC compared to the other periods. Stable2, Before1, After3 and After4 had higher

326 concentrations and lower water temperature. Along axis 2 of the CCA, After4 was separated from the

other periods (Fig. 3c). pH (r = 0.61) and temperature (r = 0.40), with respective weights of +0.33 and

+0.22 were the main variables explaining these differences. After 4, which corresponded to the

329 assembly of minor event 4, was thus characterized by lower temperatures and pH.

330 The most important contributing taxa along with the canonical axes could be separated into three

assemblages (Fig. 3a). Assemblage A1 (Fig. 3a), which included the initial assemblage, was

332	characterized by a higher biomass of the diatoms Skeletonema potamos (Weber) Hasle (14.4% of
333	contribution to axis 1 and 13.1% to axis 2), centric diatoms (9.0% to axis 1 and 1. 6% to axis 2) and
334	Nitzschia acicularis (Kützing) W.M. Smith (4.77% to axis 1), and the green algae Chlamydomonas sp.
335	Ehrenberg (8.0% to axis 1 and 3.7% to axis 2), Coelastrum microporum Nägeli (1.2% to axis 1 and
336	2.0% to axis 2) and Kirchneriella spp. (1.8% of axis 2 with K. contorta (Schmidle) Bohlin, K. obesa
337	(W. West) Schmidle and K. irregularis (GM Smith) Korshikov). Assemblage A2, corresponding to the
338	Stable1 and Before2 periods, showed higher biomasses of the green algae Actinastrum hantzschii
339	Lagerheim (10.4% to axis1), Crucigenia lauterbornii (Schmidle) Schmidle (10.7% to axis 1),
340	Treubaria triappendiculata Bernard (6. 6% to axis1), Golenkinia radiata Chodat (3.9% to axis1),
341	Lobomonas ampla Pascher (3.6% to axis1), Dictyosphaerium pulchellum Wood (4.4% to axis1),
342	Oocystis spp. Nägeli ex A.Braun (2.6% to axis1), and Didymocystis spp. (2.7% to axis1 with D. fina
343	Komárek, D. bicellularis (Chodat) Komárek and D. planctonica Korshikov). Assemblage A3
344	corresponds to After4, with a higher biomass of the benthic diatom Diatoma vulgaris Bory (4.1% to
345	axis 1 and 36.7% to axis 2) and Scenedesmus spp.: S. spp. 3 Meyen with for instance S. group
346	Scenedesmus sensu stricto sensu Komárek & Fott (3% to axis 1 and 12.7% to axis 2), S. spp.1 Meyen
347	with for instance S. group Abundantes / Spinosi sensu Komárek & Fott (7.8% to axis 2),
348	Acutodesmus spp. (Hegewald) Tsarenko (5% to axis 2), Desmodesmus spp. (R. Chodat) S.S. An, T.
349	Friedl & E. Hegewald (4.5% to axis 2) and S. spp. 2 Meyen with for instance S. group Armati
350	sensu Komárek & Fott (2% to axis 2). The benthic cyanobacterium Merismopedia spp. also
351	contributed to axis 2 (1.16% with <i>M. glauca</i> (Ehrenberg) Kützing and <i>M. tenuissima</i> Lemmermann).
352	This event-driven short-term succession was also found in the time series of species biomasses (Fig.
353	4). S. potamos (Fig. 4b) and Chlamydomonas sp. (Fig. 4c) were positively correlated (Pearson's
354	correlation, $r = 0.52$ ; $p = 0.008$ ), with higher biomasses in before1, after2, after3 and stable2
355	(assemblage 1). In contrast, G. radiata (Fig. 4d) and A. hantzschii (Fig. 4e), also correlated with each
356	other (Pearson's correlation, $r = 0.62$ ; $p < 0.001$ ), and were almost only present during stable1 and
357	Before2 (assemblage A2), i.e. during the period of the lowest discharge fluctuation (Fig. 2a). D.
358	vulgaris (Fig. 4g) showed an increase in biomass at the end of the period (After4), as do Scenedesmus
359	spp. that are previously punctually present (Fig. 4f).

#### 360 Physical and chemical parameters in summer 2014

The 2014 study period began with a long stable period in terms of discharge, followed by four 361 362 temporally close discharge peak events, with Q successively peaking on 16/07/2014, 27/07/2014, 9/08/2014 and 15/08/2014, preventing a return to initial conditions (Fig. 5a). The first discharge event 363 started with a gradual increase on 05/07/2014 (100 m<sup>3</sup> s<sup>-1</sup>) and accelerated on 10/07/2014 (210 m<sup>3</sup> s<sup>-1</sup>) 364 to reach a maximum on 16/07/2014 (402 m<sup>3</sup> s<sup>-1</sup>), corresponding to the highest peak of SM observed 365 over summer 2014 with a concentration of 55 mg  $L^{-1}$  (Fig. 5b). The following events 3 and 4 resulted 366 in slight increases in SM. A peak in POC:chl-a was observed on 06/07/2014, mainly due to the 367 decrease in chlorophyll-a concentration, potentially indicating an early effect of the first disturbance 368 369 (Fig. 5e). We considered this period to be isolated from the initially stable period and referred to as 370 Before1. Event 1 was therefore the major event of summer 2014. It corresponded with a peak of DIP (37 µg P 371  $L^{-1}$ ), which continued to increase with the following discharge disturbances up to 60 µg P  $L^{-1}$ , reached 372

on 9/08/2014 (Fig. 5c). DOC increased after event 1, with a doubling of its concentration from 2 to 4

 $mg C L^{-1}$ , and then remained above  $3 mg L^{-1}$  during the subsequent events (Fig. 5d). When data from

both 2013 and 2014 were taken into account, river discharge correlated positively with DOC

376 concentration (Pearson's correlation, r = 0.62; p < 0.001) and SM (Pearson's correlation, r = 0.71; p < 0.001).

- 378 Chlorophyll-a concentration showed a decreasing trend during the first (Stable) period from 35 to 25 379  $\mu$ g L<sup>-1</sup> (Fig. 5e), likely due to depleted phosphorus availability (Fig. 5c). Chlorophyll-a peaked after 380 event 2 (48  $\mu$ g L<sup>-1</sup>) and then remained under 20  $\mu$ g L<sup>-1</sup> until the end of the summer. Water discharge 381 events as disturbances decreased chlorophyll-a concentration and increased DIP concentrations.
- 382 Similarly to 2013, a significant negative correlation between chlorophyll-a and DIP was observed

383 (Pearson's correlation, r = -0.65; p < 0.001).

- 384 Diversity of the phytoplankton assemblage, monitored with the inverse Simpson index, was maximal
- during the stable period (Fig. 5f), despite declining biomass and limiting dissolved inorganic
- 386 phosphorus concentrations. Diversity gradually increased again at the end of the disturbed period.

- 387 Seven periods could be delineated based on the variations in physical and chemical parameters
- described above (Fig. 5).
- 389 Short-term responses of phytoplankton to hydrology in summer 2014
- Environmental variables explained 42.6% of variance in the phytoplankton community in 2014 (Fig.
- 6). pH was removed from the CCA to respect the rule of low collinearity between explanatory
- 392 variables. Axis 1 was positively correlated with discharge Q (r = 0.88), DIP (r = 0.78), DOC (r = 0.78)
- 393 0.71), silicon (r = 0.60) and SM (r = 0.55) and negatively correlated with nitrates (r = -0.65) and DO
- (r = -0.57) (Fig. 6b). DOC was the parameter with the highest weight (canonical coefficients), with a
- weight of +0.22, followed by nitrates (- 0.19) and DSi (+0.17). Axis 2 was correlated with SM (r =
- 396 0.59), temperature (r = 0.45) and DO (r = -0.34).
- 397 The CCA performed on the 2014 taxa biomass data separated the Stable period to the left of the
- ordination from the other periods, including Before1, the first pre-disturbed period (Fig. 6c). Axis 2,
- meanwhile, separated periods chronologically, with After1 at the top right of the ordination, After2,
- 400 After3 and Before2 in the middle and After4 at the bottom. The post-disturbance assemblages (After1,
- 401 2/3 and 4) were thus located in the periphery, while the two transitional assemblages (Before1 and 2)
- 402 were found near the centre of the ordination, indicating an average composition.
- 403 The phytoplankton taxa contributing the most significantly to the canonical axes clustered around 3
- 404 assemblages along axes 1 and 2 (Fig. 6a). Assemblage B1 corresponded to the initial assemblage
- 405 (Stable), with a higher biovolume of diversified green algae: A. hantzschii (8.4% to axis1), D.
- 406 pulchellum (6.5% to axis 1), Dictyosphaerium sp. Nägeli (5.4% to axis 1 and 6% to axis 2), C.
- 407 *lauterbornii* (5.8% to axis 1 and 3.4% to axis 2), *Micractinium pusillum* Fresenius (6.2% to axis 1),
- 408 Plagioselmis nannoplanctica (H. Skuja) G. Novarino, I.A.N. Lucas & S. Morrall (3.1% to axis 1), G.
- 409 radiata (2.9% to axis 1 and 2.3% to axis 2), Actinastrum aciculare Playfair (2.3% to axis 1 and 4.1%
- 410 to axis 2), Monoraphidium spp. (2.5% to axis 1 with M. arcuatum (Korshikov) Hindák, M. circinale
- 411 (Nygaard) Nygaard, *M. contortum* (Thuret) Komárková-Legnerová, *M. griffithii* (Berkeley)
- 412 Komárková-Legnerová, and *M. tortile* (West & G.S.West) Komárková-Legnerová), *Acutodesmus* spp.
- 413 (2.3% to axis 1 and 11.3% to axis 2), *Franceia ovalis* (France) Lemmermann (1.3% to axis 1). Only
- 414 one diatom contributed to the assemblage 1, with *N. acicularis* (3.1% to axis 1 and 5.2% to axis 2).

415 Assemblage B2 associated with the period following the first major disturbance (After1) was 416 characterized by an increase in Tetraedron spp. (3.3% to axis 2, with T. caudatum (Corda) Hansgirg, T. incus (Teiling) G.M. Smith and T. minimum (A.Braun) Hansgirg), small indeterminate flagellates 417 (1.8% to axis 2), Acutodesmus spp. (11.3% to axis 2), and Scenedesmus spp. 1 (4.2% to axis 1 and 418 419 5.5% to axis 2). This B2 assemblage was also characterized by decreasing biovolumes of the diatom 420 N. acicularis (3.1% to axis 1 and 5.2% to axis 2), and the green algae Dictyosphaerium subsolitarium 421 Van Goor (6.1% to axis 2), C. microporum (4.5% to axis 2), A. aciculare (2.3% to axis 1 and 4.1% to axis 2), and Chroococcus sp. Nägeli (5.1% to axis 2). Assemblage B3 observed at the end of the 422 423 disturbed period (After4) was characterized by a higher biovolume of the benthic cyanobacterium 424 Merismopedia spp. (9.6% to axis 1 and 3.3% to axis 2), the diatom S. potamos (25.3% to axis 1 and 425 5.0% to axis 2) and the green algae Scenedesmus spp. 3 (3.1% to axis 1). 426 This succession with a calm period in terms of river discharge, followed by a period with several discharge events was observed in the time series of species biovolumes (Fig. 7), with a bell-shaped 427 curve of colonial green algae during the stable period (Fig. 7a and b). Tetraedron spp., a small green 428 429 alga, although remaining in low biomass, was only observed during the transition period (Fig. 7c). 430 Some genera such as *Scenedesmus* also took advantage of this transition period (Fig. 7f). The benthic cyanobacteria Merismopedia spp. and the diatoms S. potamos and N. acicularis increased temporarily 431 432 following the major first disturbance (Fig. 7d, e and g). 433 Phytoplankton assemblage and functional trait compositions between hydrologically contrasting

# Phytoplankton assemblage and functional trait compositions between hydrologically contrasting summers

435 Six characteristic phytoplankton assemblages were identified in 2013 (Fig. 3a) and 2014 (Fig. 6a). We 436 could group them in pairs based on their location in relation to the river discharge disturbances (Fig. 437 8): two were in stable condition (A2 and B1 grouped as stable assemblage), two following a strong 438 disturbance (A1 and B2, grouped as disturbed assemblage) and two after a series of disturbances, 439 called here 'repeatedly disturbed' assemblage (A3 and B3). Common species could be identified with at least four green algae during stable conditions (A. hantzschii, D. pulchellum, C. lauterbornii, G. 440 radiata) and one benthic cyanobacterium genus in the repeatedly disturbed period (Merismopedia 441 spp.) (Fig. 8). Some diatoms were observed over two periods, such as the needle-shaped species N. 442

443 acicularis during stable and disturbed periods, and the flexible filaments of S. potamos during disturbed and repeatedly disturbed periods. Several Scenedesmus taxa were observed in the samples, 444 445 in particular in the repeatedly disturbed periods, in B1 and B2, but the specificity of the species to a 446 period was less markable. Three typical assemblages of the Loire River could thus be specified here (Fig. 8): 447 448 1) The 'stable' assemblage occurred each time in phosphorus-depleted condition, with a diverse 449 assemblage, but under the dominance of colonial green algae and diatoms. The stable assembly could be characterized by 16 species: four species have spine-like morphology (n°5 450 G. radiata; n°8 F. ovalis; n°12 M. pusillum; n°13 T. triappendiculata), six species are 451

452 colonial green algae (n°2 *D. pulchellum*; n°3 *Tetrastrum triangulare* (Chodat) Komárek; n°4

- 453 A. hantzschii and A. aciculare; n°11 Oocystis spp. ; n°12 M. pusillum ; n°14 Scenedesmus
- 454 spp., *Desmodesmus* spp. and *Acutodesmus* spp. and two are mobile, with two flagella per cell

455 (n°7 *L. ampla;* n°9 *P. nannoplanctica*). Even if several species were colonial, their size was

456 rather medium compared to large colonial cyanobacteria or Pennate diatoms. The single

457 diatom taxon that regularly occurred during the stable phase was the needle-shaped *N*.

458 *acicularis*. One third of the 'stable' group species belonged to the co-occurring functional

459 groups **F** and **J**, which correspond to middle and lower sections of large rivers and indicate

- 460 low turbulence with enhanced residence time and light availability (Abonyi et al., 2021).
- 461 Finally, these 16 taxa observed in stable conditions belonged to only 5 Reynolds FGs (**D**, **F**, **J**,
  462 **X1**, and **X2**).

2) The 'disturbed' assemblage contained 10 taxa, observed either during A1 or during B2. Three
species were colonial green algae (n°15 *Kirchneriella* spp., n°14 *Scenedesmus* spp. and n°16 *C. microporum*) and one was mobile, with two flagella per cell (n° 18 *Chlamydomonas*). The
flexible filaments of *S. potamos* (n°19) were also observed. The needle-shaped *N. acicularis*(n°1) already observed during Stable period showed a contrasting response to disturbance,
with an increase during A1 and a decrease during B2. The group J of Reynolds was still the
dominant functional group. The characteristic taxa of the 'disturbed' assemblages also

belonged to four Reynolds FGs (**D**, **F**, **J**, **X2**, thus loosing **X1** compared with the 'Stable'
assemblage).

The 'repeatedly disturbed' assemblage had only four characteristic taxa, but all representing different Reynolds FGs (still D and J, with T<sub>B</sub> and Lo more; Fig. 8). The flexible filaments of *S. acicularis* (FG D) were still observed. The benthic cyanobacterium *Merismopedia* spp. (n°20), specific of that period, was observed during both A3 and B3 and was the only representative of the group Lo. A new benthic diatom was observed, *D. vulgaris* (n°21; FG T<sub>B</sub>). Finally, half of the four species were benthic, half were diatoms and only one a small colonial green (*Scenedesmus* spp., FG J).

The discriminant analysis (DA) performed on the CA of RFG biomass shows that 19% of total 479 480 variance of the RFG data was explained by three assemblages' types (Fig. 9). The DA separated the stable period on the left from the disturbed period on the upper right and the repeatedly disturbed 481 period on the lower right of the plot (Fig. 9). The FG F (cov. -0.83 with axis 1; clear epilimnetic 482 habitat) and X1 (cov. -0.79; shallow mixed layer in enriched conditions) characterized the stable 483 484 assemblage. The FG  $T_B$  (cov. 0.55 with axis 1; highly lotic environments, streams and rivulets) characterized both disturbed and repeatedly disturbed assemblages. The FG **D** pointing to the upper 485 right (covariance of 0.55 with axis 1 and 0.32 with axis 2; shallow enriched turbid water including 486 487 rivers) characterized the disturbed assemblage. The FG Lo pointing to the lower right (cov. -0.77 with 488 axis 2; summer epilimnion) characterized the repeatedly disturbed period.

Most RFG and taxa positively correlated with water temperature and solar radiations, while they negatively correlated with river discharge and DIP, irrespective of life form and taxonomic position (diatoms, benthic taxa or colonial green algae, Fig. 10). Chlorophyceae biomass thus correlated positively with solar radiation (Pearson's correlation, r=0.30; p=<0.01) and negatively with river discharge (Pearson's correlation, r=-0.30; p<0.01). Only the benthic diatom *D. vulgaris* correlated negatively with temperature, and positively with nitrates.

495 Variance partitioning showed that time generally explained a low proportion of total variance in 496 phytoplankton, with less than 5% (except the 22% for *Chlamydomonas* sp. in summer 2013; see X1 on 497 Fig. S2). DIP concentration explained between 10 and 40% of the variance at both the seasonal and

short-term time scale for centric diatoms and *N. acicularis* (codon D – see X3 on Fig. S2). Interestingly, water temperature appeared as the most significant predictor for many taxa if data were pooled, but also when analysed separately (see X2 on Fig. S2).

501

#### 502 **Discussion**

503 Here, we asked how short-term alterations in water discharge affected the physical and chemical 504 environment and corresponding changes in phytoplankton community composition and diversity in the 505 Middle Loire. In two contrasting summers, our data suggested that hydrological events systematically 506 affected phytoplankton composition, along with three assembly types: a stable assemblage type with 507 many colonial green algae; a disturbed assemblage type that decreased in biomass but benefited from 508 new taxa such as S. potamos; and a repeatedly disturbed assemblage type that benefited from the 509 resuspension of benthic species, temporarily replacing colonial greens. These short and long responses to disturbance allowed for the maintenance of greater taxonomic and functional diversity on a seasonal 510 511 scale in the Loire River. Taxa richness is mainly driven by green algae and diatoms in accordance with literature (Wehr & Descy, 1998; Wang et al., 2014). This suggests that community dynamics of 512 513 phytoplankton are predictable in the Middle section of large rivers, if hydrology and hydrology-related parameters are monitored with high resolution. 514

515 The biogeochemical functioning of large rivers differs strongly between high and low flow, the latter 516 corresponding to a growing period for phytoplankton and so represents strong biological activity. 517 Variations in discharge, e.g. "water flush", can occur over a few hours that dilutes phytoplankton 518 biomass, increases the mortality rate, and changes a full set of environmental conditions (e.g. turbulence, 519 light climate, nutrients availability). However, community dynamics in phytoplankton has rarely been 520 studied at a sub-weekly time scale – but chlorophyll-a (Bowes et al., 2016) and community biomass 521 (Pathak et al., 2021) - in response to river hydrology (Istvànovics & Honti, 2011), in particular at a fine-522 scale taxonomic and functional levels.

523 Here, we could identify eight water discharge events that have affected the Loire River phytoplankton.

524 Even minor discharge increases modified the physical and chemical parameters with an increase in the

525 POC:chl-a ratio, a decrease in phytoplankton biomass, and a change in assemblage structure. Pickett 526 and White (1985) defined disturbance as "any relatively discrete event in time that disrupts ecosystem, 527 community, or population structure, and that changes resources, availability of substratum, or the 528 physical environment". The hydrological events thus represented disturbances for the phytoplankton, leading to a dilution of the bulk community but with alterations in community composition based on 529 species functional characteristics before, during, and after disturbance. Wang et al. (2014) highlighted 530 531 distinct assemblages driven by hydrological forcing in a large subtropical river, with after-flood assemblages associated with higher abundance of green algae and tychoplanktic taxa. While 532 disturbances in rivers are generally associated with high floods, which then mobilize nearby sediments, 533 here, we refer to a disturbance as soon as the river discharge increases even slightly, such as a daily 534 535 increase of 5 m<sup>3</sup> s<sup>-1</sup> day<sup>-1</sup>. Our study highlights that hydrology-related disturbance is not necessarily related to any threshold, rather, the disturbance can even under slight changes in hydrology. 536

The rate of change of the river discharge appears more important than its intensity to constrain the phytoplankton biomass and assemblage in the Loire River. The discharge rate of change with the previous day was overall mainly negative: river discharge decreased compared with the previous day 62% of days (all seasons considered), while it increased during 35% of the days or remained stable for 3% of the days. When we focused on the period of May to September, river discharge decreased 67% of the days. Hydrological recession thus represents the most common condition in the Middle Loire River in summer.

#### 544 Short-term changes in physical and chemical parameters

545 Temperature, solar radiations, nutrients and river discharge have been identified as main drivers of 546 river phytoplankton at the seasonal scale (Descy et al., 2013). We had expected that during summer 547 high temperature influenced less phytoplankton production and assemblages, being already high 548 compared with spring and fall. In summer, mean daily water temperature varies by a few degrees over 549 a few days, following meteorological forcing and high/low pressure events. Water temperature was a 550 stronger predictor for the biomass of individual phytoplankton taxa than solar radiation. Most of the 551 variance in taxa biomass was also explained by water temperature. It is therefore likely that water 552 temperature indirectly affected phytoplankton growth, e.g. jointly with other meteorological

553 conditions such as calm and sunny versus disturbed periods. River temperature follows a diurnal cycle 554 and a seasonal cycle, associated with meteorological forcing and depending on the hydrological 555 context. Water temperature is sensitive to air temperature, relative humidity, solar radiation and river 556 width, but also river discharge, upstream temperature (Gu & Li, 2002), and lateral groundwater inputs (Wawrzyniak et al., 2017). More than a controlling factor, water temperature seems to be a good proxy 557 558 for short-term meteorological and hydrological forcings. Changes in the similarity of pelagic diatoms 559 in a large German river have been successfully linked to the antecedent precipitation index, which considers rainfall and wetness of the basin, and so remobilisation on a wider scale (Wu et al., 2016). 560 561 At the short time scale, CCA showed that DIP and DOC were associated with discharge peak events 562 for both 2013 and 2014, where suspended matter (SM) also increased nearly simultaneously with 563 discharge peaks. In 2014, SM strongly increased just after the first main disturbance, while next events, despite being higher in intensity, did not lead to higher SM concentration. Remobilization of 564 organic and inorganic particles from the benthos in the Middle Loire seems to depend on the delay 565 between discharge disturbances, with a small remobilization if time for settling of suspended particles 566 567 from one event to the next is too short.

In conjunction with peaks in SM, most of the discharge disturbances resulted in simultaneous drops in 568 chlorophyll-a concentration, likely by dilution. It is known that phytoplankton biomass in large rivers 569 570 varies inversely with discharge, and is linked to water residence time and hydrology-mediated light 571 availability (Wehr & Descy, 1998; Várbíró et al., 2018). However, on our semi-weekly time series, we observe that chlorophyll-a tends to increase rapidly 2-6 days after a discharge event, probably due to 572 573 both the enrichment by meroplankton and the phytoplankton growth stimulated by the mobilisation of 574 nutrients (Minaudo et al., 2018). The increase of SM in the river recharges phosphorus, by desorption 575 processes from the particulate phase to the dissolved one and diffusion of pore water from the 576 underlying sediment (Reddy et al., 1995). Discharge events in summer also flush DIP-repleted water 577 with direct and diffuse inputs from the upper catchment. Disturbances have, therefore, first a negative 578 impact on the phytoplankton biomass (dilution effect combined with light limitation due to 579 resuspended solids), then a positive impact by recharging the ecosystem in potentially bio-assimilable

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580 phosphorus. The response of the assemblages to short-term forcing in P-limited conditions should

581 differ from the response in eutrophic conditions.

582 Short-term nutrients dynamic in summer can be related here to short-term dynamics in phytoplankton and the uptake of nutrients for its biomass. The summer concentration of DIP was lower than 10 µg P 583  $L^{-1}$  in 50% of the samples, while nitrates concentration, with a minimum concentration of 950 µg N  $L^{-1}$ 584 <sup>1</sup>, was sufficient throughout the year for the growth of phytoplankton. A mean biomass of 60  $\mu$ g L<sup>-1</sup> (we 585 had 40 to 50  $\mu$ g L<sup>-1</sup> during this study) indeed requires ~ 100  $\mu$ g P L<sup>-1</sup> (Dillon & Rigler, 1974) and 722 586 µg N L<sup>-1</sup>, based on the Redfield ratio. The concentrations of silicon remained also high enough most of 587 the year, with values >1 mg SiO<sub>2</sub>  $L^{-1}$  in 99% of samples. The lack of correlations between DSi or nitrates 588 589 concentrations with the phytoplankton biomass can be explained by the fact that nutrient uptake by 590 phytoplankton remained low compared to the nutrient stock in the river. On the contrary, concentrations of DIP and chlorophyll-a were negatively correlated. Short-term DIP concentration in the Loire River 591 is thus controlled by phytoplanktic uptake, with maximum DIP concentrations reached when 592 chlorophyll-a concentration is minimal. Descy (2012) also observed concentrations <10 µg P L<sup>-1</sup> from 593 594 June to October in the Loire River. The molar N:P ratio calculated on biologically available nutrients (Ptacnik et al., 2010) was always > 64 during that study, clearly above the 16:1 Redfield ratio, also 595 confirming that P was a limiting nutrient for growth, in accordance with literature (Minaudo et al., 2015). 596 597 Over the last 30 years, the Loire River has indeed undergone changes in its phytoplankton assemblages, 598 with a 90% biomass decrease, which has been related to different factors: P reduction, altered discharge 599 and the invasion of the Asian clam Corbicula sp. (Minaudo et al., 2021). Planktic cyanobacteria always remained a minor fraction of the total phytoplankton biomass, irrespective of the study period (Descy et 600 al., 2012; Minaudo et al., 2021). 601

602 Short-term DOC dynamics were also related to river discharge, with increases of 1 to 5 mg C L<sup>-1</sup>

following discharge disturbances. Most of the total organic carbon (TOC) was in the dissolved form,

- 604 with DOC representing ~70% of the TOC, in accordance with literature (Peterson et al., 1994) and
- other stations on the Loire River (Minaudo et al., 2015). The highest DOC concentrations were
- observed during the main hydrological events, indicating a remobilization of DOC pool during
- discharge peak events. DOC originates from both terrestrial sources (hydrologically connected soils

608 and wetlands) and local production by phytoplankton, microphytobenthos and the microbial loop. 609 Upstream terrestrial DOC inputs thus fuels a semi-labile pool, more conservative, while local 610 production fuels a labile pool, rapidly lost through photolysis, flocculation, consumption or adsorption (Del Giorgio & Pace, 2008). A higher local production or lower lost processes may explain that DOC 611 concentration generally increases after a discharge disturbance. DOC could thus represent an excellent 612 613 chemical proxy for river discharge disturbances in the Loire River, and also elsewhere. 614 In recent years, the use of *in situ* high-frequency spectrophotometric and electrode technologies, particularly of chlorophyll fluorescence probes, represents a good opportunity to go further in the 615 616 study of short-term dynamics with time series data at a scale down to the minute (Wade et al., 2012). 617 Thresholds of environmental drivers can be highlighted, as for water discharge and temperature whose 618 thresholds for cyanobacteria blooms could have been calculated in a eutrophic river (Bowes et al., 619 2016). The main limitation of new technologies is that nutrients are by now rarely measured at such a fine time scale. Moreover, the simultaneous acquisition of assemblage data remains scarce, although 620 sensor development is progressing. The coupling of high frequency physical and chemical 621 622 measurements with daily monitoring of species by flow cytometry or sequencing for instance would be a real step forward in the near future to increase our understanding of assembly rules and short-term 623 624 drivers of river phytoplankton.

625 Short-term functional changes in phytoplankton in response to hydrological disturbance

626 The phytoplankton assemblage measured semi-weekly was in agreement with the seasonal succession already described in the Loire River (Abonyi et al., 2012, 2014; Descy et al., 2012) and more generally 627 in large rivers, with a spring bloom of diatoms, followed by small centric diatoms in May-June, and by 628 629 Chlorophyceae during summer (Descy et al., 2012, 2013). Diatoms represented ~10% of the total 630 biomass in summer, which indicates stable low flow conditions that can trigger the sedimentation of 631 large diatoms. It is known that centric diatoms develop successfully in well-mixed, nutrient-replete and turbid rivers, which are not so different from shallow lakes (Reynolds et al., 1994). Here also, 632 small centric diatoms dominated the community over large diatoms when present. Summer coccal 633 green algae, some with ornamentations and/or mucilage, were typical of the summer highly diverse 634 Chlorophyceae assemblage (Descy et al., 2012). Their dominance has been associated to high light 635

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636 and shallow depth as main environmental drivers (Abonyi et al., 2012). Coccal green algae took 637 advantage of stable periods between disturbances, in accordance with the benthic retention hypothesis 638 (Istvanovics & Honti, 2011). Many species of the planktic genera Scenedesmus, Acutodesmus and Desmodesmus were also observed during all periods from stable to repeatedly disturbed periods. These 639 genera are among the commonest green algae and one of the most diverse genera (John et al., 2002), 640 641 and have already been observed with high biomass in many rivers elsewhere (Gosselain et al., 1994). Their small size and rapid growth rate make these algae very competitive in rivers and water bodies, 642 allowing a good sustainability of populations in high flushing environments where dilution effects are 643 644 likely. Scenedesmus spp. have already been related to hydrological factors and temperature in a 645 German lowland river (Wu et al. 2011). Also, coccal greens represent a group that is favoured by 646 sedimentation and resuspension during hydrological events (Reynolds & Descy, 1996). In nutrient 647 depleted conditions, small cells are also more competitive for nutrients uptake than large ones at low concentrations, thanks to a higher surface/volume ratio. While these taxa are especially sensitive to 648 649 grazing by zooplankton, top-down effects remain largely understudied in the Loire River (Lair & 650 Reyes-Marchant, 1997), but also elsewhere.

Despite the discharge disturbances, colonial green algae, such as *Kirchneriella* spp. or the coccal C. 651 652 microporum, remained an important part of the total biomass. On the contrary, colonial green algae 653 with spine-like ornamentations decreased and raised the question of a counter-selection driven by the 654 turbulent regime. The presence of spines may be useful against grazing and to regulate sinking velocity. Two main strategies against grazing were observed here (Lürling, 2021), the presence of 655 656 spines and the colonial form, sometimes mucilaginous. Grazing pressure in large river is, however, 657 less important than in lakes, because zooplankton is dominated by small-sized fast growing species 658 because of flushing (Wehr et al., 2015). However, our knowledge on how long-term changes in 659 hydrology with extended low flow conditions and re-oligotrophication affected zooplankton 660 community composition remains largely elusive. Spines and protuberances can also increase the 661 resistance to sedimentation, by increasing fluid drag coefficient, and stabilization of the horizontal 662 position with the highest form resistance (Padisák et al., 2003). Spines can also increase the effects of

viscosity at an individual scale like mucilage (Naselli-Flores et al., 2021) and help regulate populationdensity to avoid collapse.

665 S. potamos, a filamentous diatom, was also observed in the aftermath of summer discharge 666 disturbances. Cells of S. potamos form a chain attached by external tubes organized in one marginal ring, giving flexibility to the filament able to support deformations. The length of the chain depends on 667 river turbulence (Duleba et al., 2014). S. potamos is one of the most important species in some **A** 668 669 eutrophic rivers, with high biomass occurring during the growing season and seems favoured by warmer temperatures and high light availability (Kiss et al., 1994; Abonyi et al., 2018b), including in 670 671 the Loire River (Descy et al., 2012). It is a warm-stenotherm centric diatom, with a relatively recent dispersion across the World (Duleba et al., 2014), which may make it as a good indicator of late 672 673 summer warm conditions among planktic river species. Increase in its relative biomass since the last 40 years has been observed in the Danube River and was linked to the increase in water temperature 674 (Duleba et al., 2014). This species was also observed by Descy in the Moselle River and related to the 675 stable summer conditions and the minimum discharge (Descy, 1993). This centric diatom has a thin 676 677 frustule weakly silicified and small chloroplasts. Silicification intensity varies greatly among diatoms 678 and is a strategy against grazing but with a cost on the competition abilities (Assmy et al., 2013; 679 Pančić & Kiørboe, 2018). S. potamos is thus competitive for silicon, with an optimal quota two to 680 three times lower than other centric and large diatoms (Descy et al., 2012). Large filament well 681 silicified of Aulacoseira granulata (Ehrenberg) Simonsen used to be observed in the Loire River before 1990 when nutrient availability was high, in particular silicon  $>7 \text{ mg L}^{-1}$  (Lair & Reyes-682 Marchant, 1997). Its cell and filament sizes can also change with turbulence in a large river (Wang et 683 al. 2015). High summer temperature, higher turbulence and low DSi requirement may explain the 684 685 dominance of S. potamos following discharge disturbances in the Loire River. Also, while DSi might 686 not be limiting in most of the European rivers, the Si:P ratio has largely been altered due to P decrease 687 and damming, likely affecting phytoplankton (Abonyi et al., 2014; 2018a). 688 The needle-shaped diatom N. acicularis maintained its biomass after the discharge disturbances, and

regularly increased in biomass. This truly planktic species is very common in the Loire River and can

690 be abundant at high river discharge (Descy et al., 2012). As shown by experiments and numerical

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691 simulations, a needle-like shape can lead to aggregation during sedimentation thanks to hydrodynamic 692 interactions at low Reynolds number, independently on fluid shear and Brownian motion, (Botte et al., 693 2013). This can be a strategy to decrease sedimentation rate for needle-like shape taxa, such as 694 Pseudo-nizschia spp., which forms chains of needles in stable periods (low Reynolds number), while it becomes individualized in periods of flooding (high Reynolds number) to be more easily driven by 695 turbulent eddies. It would be interesting to make observations of fresh samples of N. acicularis in 696 697 sedimentation to see if this hypothesis is valid for this species. More than minimizing sinking velocity, phytoplankton traits are selected to maximize the entertainability of cells by turbulent eddies (Naselli-698 699 Flores et al., 2021). Interestingly, the two main diatoms N. acicularis and S. potamos belong to both 700 freshwater and marine genera, suggesting a shape that withstands high turbulence as in the sea. 701 D. vulgaris, a characteristic benthic diatom which normally forms zig-zag colonies attached to the 702 substrate, increased in biomass during repeated discharge events. Living in various habitats in rivers, 703 such as rock and plants, it predominates in eutrophic water with high conductivity and with pH > 7704 (Kelly, 2000). This benthic diatom has been observed in high biomass in upstream reaches of the Loire 705 River (Abonyi et al., 2012), but also in the Middle Loire during high flow periods, with other species 706 like Amphora ovalis (Kütz.) Kütz. or Cocconeis placentula Ehrenberg (Descy et al., 2012). Its 707 presence is, therefore, an excellent indicator of flushing or high discharge events. 708 Benthic cyanobacteria observed following repeated discharge disturbances were M. tenuissima and M. 709 glauca. Both meroplanktic species are often associated with bottom silt and live at the water-sediment 710 interface, even if they can be observed in the water column (John et al., 2002). M. tenuissima is also 711 indicative of eutrophic conditions, with bottom reducing conditions (John et al., 2002). Already widely 712 observed in the Loire River, it probably develops in the main river channel on the riverbanks 713 (Minaudo et al., 2021). As for cyanobacteria, warm summer temperature may have promoted this 714 species. 715 Benthic taxa were thus the most responsive group to repeated disturbances, in accordance with their 716 shear-induced resuspension. The presence of benthic diatoms and cyanobacteria with discharge 717 disturbances has already been highlighted in the literature (Rojo et al., 1994; Abonyi et al., 2012).

718 Antecedent discharge disturbance is thus important to consider for the assembly rules of

719 phytoplankton, as well as the role of meroplankton in sustaining diversity. The maintenance of 720 functional richness is needed to sustain ecosystem functions, while high taxonomic richness with 721 functional redundancy increases the insurance of the community against disturbances (Naeem, 2008). 722 Other biotic and abiotic factors than discharge may have controlled phytoplankton assemblage during 723 stable periods, such as spatial heterogeneity. It has been shown that the Loire River has a very high 724 taxonomic diversity, partly explained by its relative shallowness, braided arms and 'dead zones' 725 occurring in the Middle and Lower river sections (Descy et al., 2012). Spatial heterogeneity in phytoplankton distribution increases as river discharge decreases. Our sampling station, located in the 726 727 Middle Loire River, has no significant upstream tributaries over 272 km, but many disconnected pools can form under low flows and may represent a potential inoculum of planktic species. Grazing 728 729 pressure and selectivity may also have controlled phytoplankton assemblage. Small zooplankton 730 probably developed during stable period of summer months, favoured by low discharge and high temperature, especially because we observed traits against grazing. Such zooplankton data would help 731 increase our understanding of the functional traits of green algae. Mussels, such as the Asian clams 732 733 Corbicula spp., are also known to reduce phytoplankton and meroplankton biomasses in the Loire River (Descy et al., 2012). However, feeding selectivity seems to be low in mussels (Boltovskoy et al., 734 1995; Frau et al., 2016). One can also consider that *Corbicula* spp. grazing should not change on a 735 736 short-term scale in response to discharge except if the mixing regime shifts from laminar to turbulent. 737 Mixing regime of the water column is indeed essential to bring phytoplankton cells in contact with benthic invertebrates (Smith et al., 1998). 738

With climate change, summer conditions of high temperatures and strong solar radiations are likely to favor benthic primary producers, while phytoplankton may remain limited by phosphorus availability. Severe low water levels, such as those observed in 2022 in the Loire River, will likely enhance benthic growth, on the other hand, likely decrease exchanges between the benthic and planktic compartments. The decrease of meroplankton enrichment, i.e., taxa that can potentially live both sedimented and resuspended, could compromise the biodiversity of the Loire River. Meroplankton should be emphasized along with the benthic retention hypothesis (Istvànovics & Honti, 2011) and points at the

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746 importance of hydrological variations in mediating phytoplankton biodiversity, biomass and

747 ecosystem functioning via resuspension.

#### 748 Conclusion

749 This study analyzed the impact of short-term discharge disturbances on physical, chemical and biological parameters and on the Loire River phytoplankton assemblage. Suspended matter increased 750 751 with river discharge, while chlorophyll-a concentration decreased due to the dilution effect. We 752 highlighted three assemblage types along with increasing hydrological forcing: (i) a short community response associated with the dilution and collapse of community dominance, followed by the enrichment 753 754 of benthic species. Discharge disturbances promoted species with specific adaptations to turbulence, while stable periods in terms of hydrology promoted species with adaptations against sedimentation and 755 756 grazing. DIP dynamic was directly linked to its consumption by phytoplankton as it correlated 757 negatively with biomass. Even in the short term, temperature remains a key parameter, which was positively correlated to taxa biomass and is likely an excellent proxy for meteorological and hydrological 758 forcings at the short-term scale. In a context of re-oligotrophication and global warming, meroplanktic 759 760 algae will likely plays a decisive role in maintaining phytoplankton diversity and ecosystem functioning in the Loire River, which may also likely the case in other shallow and large rivers. 761

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#### 774 Author Contributions

- FM, CM and NG collected data on physical and chemical parameters; AA and ML collected data on
- phytoplankton; AP and CM conceptualized the study, and performed the data analyses and statistics.
- AP, CM, NG and AA wrote the manuscript, with equal contributions from all authors.
- 778

#### 779 Data Availability Statement

- 780 The data presented in this study will be available as Supplementary materials.
- 781

#### 782 Conflicts of Interest

783 The authors declare no conflict of interest.

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#### 1088 FIGURE CAPTIONS

- 1089 Figure 1. Identification of hydrological events and associated periods. An increase in the slope dF/dt
- 1090 between two consecutive days highlights the beginning of a disturbance and is noticed t\*. The
- 1091 'Before' period corresponds to the seven days preceding t\* and begins on day  $t_1$  ( $t_1 = t^* 7$  days). The
- 1092 'After' period corresponds to the disturbed period, extending from day t\* to the return to stable
- 1093 conditions on day  $t_2$ .

**Figure 2**. Time series in summer 2013 (a) of the river flow Q and the ratio particulate organic carbon : chlorophyll-a (POC:Chl-a), with the same y-axis scale, (b) of concentration in suspended matter, (c) of dissolved inorganic phosphorus (DIP) concentration, (d) of dissolved organic carbon concentration, (e) of chlorophyll-a and pheopigments concentrations, and (f) of the inverse Simpson's index. Vertical red dotted line shows the main disturbances with their numbering. Periods identified are shown at the top of the figure.

- **Figure 3**. CCA performed on phytoplankton assemblages from 2013 explained by environmental
- parameters (p=0.001). (a) Plot of the taxa. Only taxa contributing to at least 1% to canonical axis are

shown. (b) Plot of the environmental parameters (Q: water discharge; SM: suspended matter; Nitrates:

- 1103 nitrates concentration; DSi: DSi concentration; DIP: dissolved inorganic phosphorus concentration;
- 1104 Temperature; DOC: dissolved organic carbon concentration; pH; DO: Dissolved Oxygen). (c) Plot of
- the sampling dates, grouped afterwards according to the period (Bef1: before 1; Bef2: before 2; Aft1:
- After 1; Aft2: After 2; Aft3: After 3; Aft4: After 4; Stable 1; Stable 2) See Fig. 2 for periods' details.

1107 Scale of x and y axes is given by d value at the top right for each plot.

**Figure 4**. Time series of phytoplankton taxa biovolumes in mg C L<sup>-1</sup> during summer 2013, with (a)

1109 Nitzschia acicularis, (b) Skeletonema potamos, (c) Chlamydomonas sp., (d) Golenkinia radiata, (e)

- 1110 Actinastrum hantzschii, (f) Scenedesmus spp. 3 and (g) Diatoma vulgaris. Periods identified are shown
- 1111 at the top of the figure.
- **Figure 5**. Time series in summer 2014 (a) of the river flow Q and the ratio particulate organic carbon :
- 1113 chlorophyll-a (POC:Chl-a), with the same y-axis scale, (b) of concentration in suspended matter, (c) of
- dissolved inorganic phosphorus (DIP) concentration, (d) of dissolved organic carbon concentration, (e)
- 1115 of chlorophyll-a and pheopigments concentrations, and (f) of the inverse Simpson's index. Vertical red

- 1116 dotted line shows the main disturbances with their numbering. Periods identified are shown at the top1117 of the figure.
- 1118 Figure 6. CCA performed on phytoplankton assemblages from 2014 explained by environmental
- parameters (p=0.029). (a) Plot of the taxa. Only taxa contributing to at least 1% to canonical axis are
- shown. (b) Plot of the environmental parameters (Q: water discharge; SM: suspended matter; Nitrates:
- 1121 nitrates concentration; DSi: DSi concentration; DIP: dissolved inorganic phosphorus concentration;
- 1122 Temperature; DOC: dissolved organic carbon concentration; DO: Dissolved Oxygen). (c) Plot of the
- sampling dates, grouped afterwards according to the period (Bef1: before 1; Bef2: before 2; Aft1:
- After 1; Aft2: After 2; Aft3: After 3; Aft4: After 4; Stable) See Fig. 2 for periods' detail. Scale of x
- and y axes is given by d value at the top right for each plot.
- **Figure 7**. Time series of phytoplankton taxa biovolumes in mg C L<sup>-1</sup> during summer 2014, with (a)
- 1127 *Monoraphidium* spp., (b) *Kirchneriella* spp., (c) *Tetraedron* spp., (d) *Skeletonema potamos*, (e)
- 1128 *Merismopedia* spp., (f) *Acutodesmus* spp. and (g) *Nitzschia acicularis*. Periods identified are shown at
- the top of the figure.
- **Figure 8**. Illustration of species representative of the three periods, showing their morphometry and
- traits: stable assemblages with A2 and B1; disturbed assemblage with A1 and B2; repeatedly disturbed
- 1132 assemblage with A3 and B3. RFG are indicated near each draw. Taxa framed with a dotted blue line
- are observed at least over two periods. Taxa framed in a red line are observed in the same period both
- 1134 years. 1. *Nitzschia acicularis* (Kützing) W.M.Smith and *N. dissipata* (Kützing) Grunow; 2.
- 1135 Dictyosphaerium pulchellum Wood; 3. Tetrastrum triangulare (Chodat) Komárek ; 4. Actinastrum
- 1136 *hantzschii* Lagerheim; 5. *Golenkinia radiata* Chodat; 6. *Monoraphidium contortum* (Thuret)
- 1137 Komárková-Legnerová; 7. Lobomonas ampla Pascher; 8. Franceia ovalis (France) Lemmermann; 9.
- 1138 Plagioselmis nannoplanctica (H. Skuja) G. Novarino, I.A.N. Lucas & S. Morrall; 10. Didymocystis
- 1139 planctonica Korshikov; 11. Oocystis lacustris Chodat; 12. Micractinium pusillum Fresenius; 13.
- 1140 Treubaria planctonica (G.M.Smith) Korshikov; 14. Scenedesmus spp. Meyen (with for
- 1141 example Scenedesmus group Armati sensu Komárek & Fott), Acutodesmus spp. (Hegewald) Tsarenko
- 1142 and Desmodesmus spp. (R.Chodat) S.S.An, T.Friedl & E.Hegewald ; 15. Kirchneriella lunaris
- 1143 (Kirchner) Möbius; 16. Coelastrum microporum Nägeli; 17. Tetraedron minimum (A.Braun)

1144 Hansgirg; 18. Chlamydomonas sp. Ehrenberg; 19. Skeletonema potamos (Weber) Hasle; 20.

1145 *Merismopedia tenuissima* Lemmermann; 21. *Diatoma vulgaris* Bory.

1146 Figure 9. (a) Plot of the two axes of the discriminant Analysis (DA) performed on phytoplankton

1147 RFG. The discriminant factor is one of the three assemblages (stable, repeated, repeatedly disturbed)

and each sample is attached to the gravity center of the group it belongs to. (b) Correlation circle plot

showing the phytoplankton RFG. Scale of x and y axes is given by d value at the top right for each

1150 plot.

1151 Figure 10. Pearson's correlation matrix between log10-transformed RFG and taxa and environmental

parameters: Q river discharge, T° daily mean water temperature, DO daily mean dissolved oxygen,

daily mean pH, Cd daily mean conductivity, SM suspended matter, Ry solar radiations, Nitr. Nitrates,

1154 DIP, DSi and COD concentrations. Significativity of the Pearson's correlation is shown: \*means

1155 p<0.05, \*\* p<0.01 and \*\*\* p<0.001. Colored ellipses indicate the positive (blue) or negative (red)

1156 correlations.

1157 Figure S1. Time series of environmental and biological parameters at the station Cinq-Mars-La-Pile.

1158 Q: water discharge; SM: suspended matter; Nitrates: nitrates concentration; TP: total phosphorus

1159 concentration; DSi: silicate concentration; POC: particulate organic carbon; Temp.: temperature;

1160 radiation: solar radiation; Chl-a: total pigment concentration; DIP: dissolved inorganic phosphorus

1161 concentration; DOC: dissolved organic carbon concentration; Phy biom: phytoplankton biomass

- 1162 calculated from cell counts and grouped by class; Tot biom: total phytoplankton biomass calculated
- 1163 from cell counts; Turnover: turnover rate per day.

**Figure S2.** Variance partitioning of taxa biomass depending on time with julian day (X1), water

temperature (X2) and dissolved inorganic phosphorus (X3), performed with the function *varpart* of the

1166 package vegan. The analysis was performed on all data, on summer 2013 and on summer 2014.

1167







Figure 3.



First CCA axis (38.8 % of constrained inertia - 24.3% of total variance)

Figure 4.





05/06/2014 15/06/2014 25/06/2014 05/07/2014 15/07/2014 25/07/2014 04/08/2014 14/08/2014 24/08/2014

Figure 6.





05/06/2014 15/06/2014 25/06/2014 05/07/2014 15/07/2014 25/07/2014 04/08/2014 14/08/2014 24/08/20

Figure 8.



@drawings from Maria Leitao

# Figure 9.



First DA axis (63.8 % of constrained variance)

## Figure 10.

Q T° DO pH Cd SM Ry NO3 PO4 Si COD

1 D \*\* \*\*\* \* F \*\*\* \*\*\* \*\*\* \*\*\* 0 \* \* \*\*\* \* J 0.8 \*\*\* \*\*\* \*\*\* \* TΒ \* X1 \*\*\* \*\*\* \*\*\* \*\* \*\*\* \*\*\* \*\* 0.6 Х2 \*\*\* \* \*\*\* \*\* \* TD \*\* \*\* \*\* \* 0.4 Lo \*\* \*\* \*\*\* \*\* \*\* \*\*\* \*\*\* \* \*\*\* \*\*\* \*\* \*\*\* \*\* 0.2 \*\*\* \*\* \*\* \*\*\* \*\*\* \*\*\* \*\*\* \* \* \*\*\* \*\*\* \*\* \* \*\*\* \* 0 \*\*\* \*\*\* \*\* \*\* \*\* \* \*\* \* -0.2 \*\*\* \*\* \*\*\* \*\* \* \* \*\* \*\* \* \*\*\* \* \* \* -0.4 \*\*\* \*\*\* \*\*\* \*\*\* \*\*\* 0 0 \*\*\* \* \*\* \*\*\* \*\* \* \* \* -0.6 \*\* \*\* \* \*\*\* \* -0.8 \*\*\* \*\*\* \*\* \*\* \* \* \*

Monoraphidium contortum Desmodesmus spp. Nitzschia acicularis Dictyosphaerium subsolitarium Acutodesmus spp. Nitzschia sp. Nitzschia sp. Micractinium pusillum Oocystis sp. Merismopedia tenuissima Actinastrum hantzschii Golenkinia radiata Coelastrum microporum Dictyosphaerium pulchellum Diatoma vulgaris Kirchneriella irregularis Figure S1.



