

Phytoplankton community responses to temperature fluctuations under different nutrient concentrations and stoichiometry

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Citation: Gerhard, M., A. M. Koussoroplis, H. Hillebrand, and M. Striebel. 2019. Phytoplankton community responses to temperature fluctuations under different nutrient concentrations and stoichiometry. *Ecology* 100(11):e02834. 10.1002/ecs.2834

Abstract. Nutrient availability and temperature are important drivers of phytoplankton growth and stoichiometry. However, the interactive effects of nutrients and temperature on phytoplankton have been analyzed mostly by addressing changes in average temperature, whereas recent evidence suggests an important role of temperature fluctuations. In a laboratory experiment, we grew a natural phytoplankton community under fluctuating and constant temperature regimes across 25 combinations of nitrogen (N) and phosphorus (P) supply. Temperature fluctuations decreased phytoplankton growth rate (r_{\max}), as predicted by nonlinear averaging along the temperature–growth relationship. r_{\max} increased with increasing P supply, and a significant temperature \times P \times N interaction reflected that the shape of the thermal reaction norm depended on nutrients. By contrast, phytoplankton carrying capacity increased with N supply and in fluctuating rather than constant temperature. Higher phytoplankton N:P ratios under constant temperature showed that temperature regimes affected cellular nutrient incorporation. Minor differences in species diversity and composition existed. Our results suggest that temperature variability interacts with nutrient supply to affect phytoplankton physiology and stoichiometry at the community level.

Key words: carrying capacity; growth rate; N:P ratios; phytoplankton composition; temperature variability; thermal performance.

INTRODUCTION

Temperature and nutrients are among the strongest drivers of primary production, both at regional and global scales (Tadonl  k  2010, Kraemer et al. 2017). In lentic ecosystems, phytoplankton biomass and growth can respond very rapidly to changes in nutrient and temperature conditions (Moss et al. 2011, Cross et al. 2015). Previous studies have either analyzed the temperature–nutrient interactions with regard to constant temperatures (Thomas et al. 2017, Thrane et al. 2017, Verbeek et al. 2018) or the effect of temperature variability independently (Vasseur et al. 2014, Bernhardt et al. 2018b). We provide an experimental test using a natural phytoplankton community subjected to 25 nutrient regimes, each under constant and fluctuating temperature showing that temperature fluctuations have

nonneutral consequences for resource uptake, growth and standing stocks.

As for all ectotherms, temperature determines metabolic rates of phytoplankton, including growth rates (Brown et al. 2004). Phytoplankton thermal reaction norms (TRN) for growth are unimodal left-skewed functions of temperature (Thomas et al. 2012). Because TRN are skewed, small rises in average temperature above the thermal optimum can have strong detrimental effects on phytoplankton growth (Deutsch et al. 2008). Most studies on the thermal dependence of biological rates focus on changes in mean temperature, but a steadily increasing number of studies demonstrates the importance of considering thermal variability around the average (Ruel and Ayres 1999, Estay et al. 2011, Vasseur et al. 2014, Dowd et al. 2015, Schaum et al. 2018). Because of the nonlinearity of the TRN, population growth in thermally fluctuating environments differs from estimates solely based on mean temperature (Estay et al. 2014, Bernhardt et al. 2018b), a mathematical phenomenon known as Jensen’s

Manuscript received 26 April 2019; accepted 25 June 2019.
Corresponding Editor: Shelley E. Arnott.

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inequality (Jensen 1906, Ruel and Ayres 1999). The direction (positive or negative) of such variability effects depends on the curvature of the TRN at a given temperature mean value: concavity decreases population growth, whereas convexity increases it. Hence, under fluctuating temperature conditions, the “realized” TRN for growth should strongly differ (e.g., lower maximum growth, thermal optimum or thermal breadth) from that inferred under constant conditions (Bernhardt et al. 2018b).

Currently, our knowledge on thermal variability effects on phytoplankton is solely based on a single species studies at high nutrient supply (Bernhardt et al. 2018b). Nutrient supply has empirically been shown to interact strongly with temperature (Thrane et al. 2017, Marañón et al. 2018, Verbeek et al. 2018). Fundamentally, this means that at a given temperature, nutrients can change the height and the curvature of the population growth TRN (Thomas et al. 2017). In a thermally fluctuating environment, such changes of the TRN imply that the effect size of thermal variance might strongly depend on the nutritional context (Koussoroplis

et al. 2017; Fig. 1), a prediction that remains to be verified. Furthermore, it is necessary to expand our understanding of thermal variability effects from single species to natural phytoplankton communities, because species rarely occur alone in nature.

In addition to growth rate, other important traits such as phytoplankton maximal biomass (or yield) might also be affected by interactions between temperature regimes and nutrient availability. It has been demonstrated that in phytoplankton populations, the carrying capacity decreases with increasing temperature, presenting a convex relationship (Savage et al. 2004, Bernhardt et al. 2018a). In addition, combined effects of enhanced temperature and nutrient enrichment on phytoplankton communities show an increase in algal biomass, whereas this response is constrained when only one factor is present (de Senerpont Domis et al. 2014, Verbeek et al. 2018). Using worldwide time series data, Kraemer et al. (2017) found that the correlation of phytoplankton biomass and temperature in lakes may be negative or positive, depending on the lake trophic state. Nevertheless,

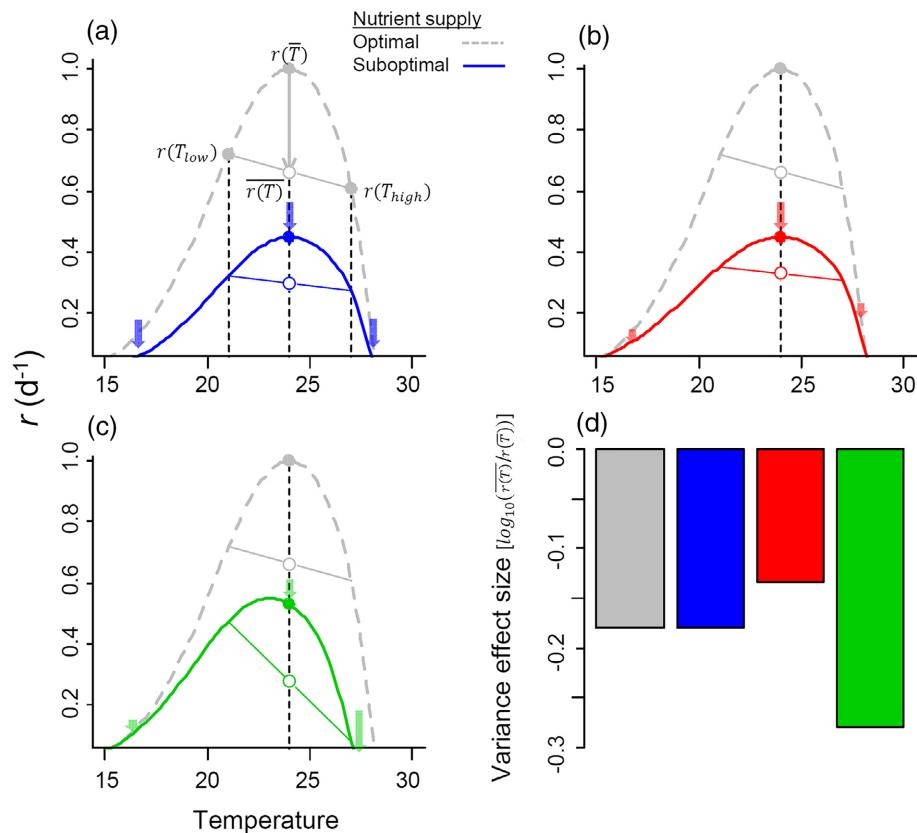


FIG. 1. Hypothetical community thermal reaction norms (TRN) and nutrient-dependent variance effect. Fluctuating temperatures (e.g., T_{low} , T_{high}) around the thermal optimum should decrease the average growth rate (r) of an algal community, $r(\bar{T}) = (r(T_{low}) + r(T_{high}))/2$, relative to that under constant temperature, $r(\bar{T})$ (Jensen’s inequality). Suboptimal nutrient supply (concentrations, ratios) could change the shape of the TRN in various ways (a)–(c). Depending on whether nutrient supply affects the different regions of the TRN proportionally or not (colored arrows), the effect size of variance on r (d) can either (a) remain unchanged (all regions of the TRN decrease proportionally), (b) decrease (proportionally higher decrease around the optimum than near the tolerance limits), or (c) increase (proportionally higher decrease of the upper tolerance limit).

how thermal fluctuations may affect maximal biomass under different nutrient conditions remains unknown and difficult to predict (as for growth).

The balance between nutrients, in addition to concentrations, is fundamental to primary producer's development and plays an important role describing how organisms use the nutrients present in the environment (Sterner and Elser 2002). Phytoplankton stoichiometry (elemental nitrogen: phosphorus, hereafter N:P ratio) generally reflects the environmental nutrient availability (Klausmeier et al. 2004a, b). However, a higher correspondence between phytoplankton N:P and supply N:P ratios is expected under slow-growing conditions (stationary growth phase), whereas in fast-growing conditions the phytoplankton stoichiometry tends to be more restricted to a range of lower N:P ratios (Hillebrand et al. 2013). Thus, the growth rate hypothesis (GRH; Sterner and Elser 2002) links growth rate with stoichiometry, proposing that there is a greater allocation of resources to P-rich ribosomal RNA under high-growth-rate conditions (exponential growth phase) generating a decrease in cellular N:P ratios (Elser et al. 2000). Moreover, recent studies have analyzed the effect of temperature on the phytoplankton resource use and stoichiometry, showing that higher temperatures may cause an increase in the phytoplankton optimal N:P supply ratio (Thrane et al. 2017) as in the phytoplankton intracellular N:P ratios (Yvon-Durocher et al. 2015). It has been proposed that fewer P-rich ribosomes are needed to maintain the protein synthesis under warmer temperatures, generating a decrease in demand for P relative to N (Toseland et al. 2013). Hessen et al. (2017) found higher RNA content in phytoplankton under lower temperature treatments, indicating a compensatory mechanism to maintain growth in colder ecosystems. However, the RNA content was not reflected in the phytoplankton N:P ratios (Hessen et al. 2017). Hence, mechanisms explaining phytoplankton stoichiometric responses to temperature are still not clear, but these responses might reflect the effect of nutrients and temperature regimes on the growth rate and biomass. Furthermore, species composition may change according to their optimal temperature (Paerl and Huisman 2008) and optimal nutrient conditions to grow (Dokulil et al. 2007, Hillebrand et al. 2013), influencing the community physiological responses to environmental conditions.

The aim of this study was to evaluate the responses of a natural phytoplankton community to different nutrient concentrations and ratios under constant and fluctuating temperature conditions. To address this topic, we tested separate hypotheses for maximum growth rate during the exponential phase (H1) and maximum biomass achieved in the stationary phase (H2).

H1: The nonneutral effect of temperature fluctuations on phytoplankton growth rate as expected based on Jensen's inequality depends on the interactions with nutrient supply concentrations and ratios. We test this hypothesis at the community level, where we propose three possible scenarios based on different effects of

resource supply on the phytoplankton TRN (Fig. 1). Depending on how suboptimal nutrient supply alters the TRN, the effect of thermal variability can increase, decrease, or stay neutral across a range of resource supply gradient.

H2: Phytoplankton maximum biomass (K) reflects the interactive effect of nutrient availability and temperature regime. K will increase with increasing nutrient supply, but also with fluctuating temperature, reflecting the convex relationship between K and temperature. As for growth, we expect nutrient supply to alter the effect size of fluctuations.

METHODS

Experimental setup

A laboratory experiment was conducted using a natural freshwater phytoplankton community. The community was collected from the Graftschafsee lake (Germany, 53°33'05" N; 7°58'49" E) in January 2017. We used a factorial design including 25 nutrient treatments and two temperature conditions (constant and fluctuating). Two replicates were generated for each treatment combination, resulting in a total of 100 experimental units. Nutrient treatments were performed using five N and five P levels, generating a wide gradient of N:P ratios (Table 1). N and P were added to the lake water as a unique pulse of K_2HPO_4 and $NaNO_3$ at the beginning of the experiment. The total dissolved nutrients (lake nutrient concentrations plus added nutrients) were considered as the nutrient supply and ranged from 0.76 to 3.33 $\mu\text{mol P/L}$ and 22.7 to 62.3 $\mu\text{mol N/L}$ (Table 1). Nutrients (except N and P) were added using growth medium stock solutions to avoid the limitation by other elements (Guillard and Lorenzen 1972). The experimental design involved a constant temperature treatment and a day–night fluctuating temperature treatment. The experimental units were incubated in a fridge under controlled conditions and a set of LED lamps was added to simulate the sunlight spectrum in a day–night cycle. The lamp cycle generated fluctuations in the air temperature,

TABLE 1. Nutrient treatments.

N ($\mu\text{mol/L}$)	P ($\mu\text{mol/L}$)				
	0.76	1.37	1.99	2.69	3.33
62.3	82	45	31	23	19
52.7	69	38	26	20	16
43.3	57	32	22	16	13
33.1	43	24	17	12	10
22.7	30	17	11	8	7

Notes: Supplied concentrations of N (rows) and P (columns), and the resulting N:P supply ratios (in bold). The concentrations and ratios correspond to the lake dissolved nutrient content plus the nutrient addition.

influencing the temperature in the treatments. These fluctuations established the fluctuating temperature treatments. The constant temperature conditions were assured using a heat plate that maintained the treatment's temperature at $9^{\circ} \pm 1^{\circ}\text{C}$. Hence, all the treatments had the same average temperature at $9^{\circ} \pm 1^{\circ}\text{C}$, but the fluctuating treatments oscillated by $4^{\circ} \pm 1^{\circ}\text{C}$ around this average (around 5°C during the night and around 13°C during the day). This average temperature was chosen to allow phytoplankton growth but also to minimize potential effects of a very high sudden increase in temperature on the natural community (lake in situ temperature at the sampling moment was 2°C).

Cell culture flasks (250 mL) were used as experiment units and filled with 200 mL lake water, previously filtered through a $53\text{-}\mu\text{m}$ mesh to remove zooplankton grazers. The flasks were daily shaken by hand and randomly rearranged to avoid sinking losses and possible location effects. Light conditions were the same for all treatments with a 16:8 day–night cycle and a light intensity of about $250 \mu\text{mol photon} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$. Temperature and light conditions were monitored with continuous data loggers (HOBO Pendant[®], Onset, Bourne, MA, USA) and optical density measurements (wavelength 440–450 nm) were performed daily during the experiment to monitor the phytoplankton growth. The experiment was interrupted at day 18 when the phytoplankton communities were expected to be in stationary state, but before a decline in biomass occurred.

Samplings and laboratory analysis

Samples were taken before the beginning of the experiment (lake water and added nutrient medium), during the phytoplankton exponential growth phase and at the end of the experiment, during the phytoplankton stationary growth phase. For each sample, we removed 50 mL from each flask without refilling. In vivo chlorophyll *a* was measured with a hand fluorometer (AquaFluor[™]; TURNER DESIGNS, San Jose, CA, USA) and samples for particulate organic carbon (C), N and P were filtered onto acid-washed precombusted glass-fiber filters (Whatman GF/C). Filters for particulate organic C and N were dried at 60°C for 3 days, put in tin capsules, and measured using an elemental analyzer (Flash EA 1112; Thermo Scientific, Waltham, MA, USA). Particulate organic P was measured by molybdate reaction after digestion with potassium peroxydisulfate ($\text{K}_2\text{S}_2\text{O}_8$) solution (Wetzel and Likens 2003). The filtered water was collected and analyzed for dissolved nutrients (PO_4^{3-} , NO_3^-) by a continuous-flow analyzer (Euro EA 3000; HEKAtech GmbH, Weggberg, Germany). To analyze the phytoplankton community, a subsample (7 mL) was taken and fixed with lugol 1%. Counting and identifying was done using an inverted microscope (Leica) based on Utermöhl's method (1958). Phytoplankton was identified to the species level and morphospecies were used when clear assignment of a species name was not possible.

Bioassay

After the phytoplankton communities reached their stationary growth phase, N and P pulses were added to identify the limiting nutrient in each treatment. For that, the two replicates of each treatment were mixed and equally redistributed in two flasks to ensure homogeneous conditions. One replicate received an N (1 mmol per 100 mL), the other a P (50 μmol per 100 mL) pulse. The response of phytoplankton biomass was measured daily (optical density) for 9 d.

Statistical analysis

Particulate organic C (hereafter particulate C), chlorophyll *a* concentrations and optical density measurements were used as phytoplankton biomass proxies. Spearman correlations showed that particulate C was highly correlated with optical density ($r = 0.93$, $P < 0.001$), but the correlation coefficient of particulate C and chlorophyll *a* was low ($r = 0.46$; $P < 0.001$; Appendix S1: Fig. S1). Because of that, we used the daily optical density measurements to determine the phytoplankton growth curves. Repeated-measures ANOVAs using the mean values of the two replicates ($n = 50$) were performed to test the effects of treatments and their changes over time on the phytoplankton N:P ratios, chlorophyll *a*, and particulate C. N:P ratios were calculated as the molar ratios of dissolved and particulate nutrients, corresponding to the N:P supply and phytoplankton N:P ratios, respectively.

The potential maximum phytoplankton biomass in the stationary growth phase (carrying capacity, K) and the maximum growth rate of the community in the exponential growth phase (r_{max}) were estimated for each treatment using the daily optical density measurements of both replicates together ($n = 50$) and following the logistic growth curve: $\text{OD} \sim K/(1 + ((K - N_0)/N_0) * \exp(-r_{\text{max}} * \text{time}))$, where OD is the optical density, *time* is the experiment day, and N_0 the start value (optical density at Day 1 in this case). GLMs including two- and three-way interactions were performed to analyze the effect of the nutrient and temperature treatments on r_{max} and K . *F*-tests and *P* values were calculated to identify significant effects and residuals analyses were used to corroborate the test assumptions. Response surfaces of r_{max} and K to the treatments and the nutrient-dependent variance effects sizes between temperature regimes were visualized as nonparametric thin-plate splines generated with the Tps function in the fields R package. The variance of r_{max} and K effects sizes between temperature regimes were calculated based on the \log_{10} -response ratio. Because not all treatments were totally saturated at the end of the experiment, K was overestimated by the logistic growth curve applied for these cases (Appendix S1: Table S1). We excluded the maximal K value of the fluctuating temperature treatment identified as outlier, because this value was far above the observed optical density

values and might lead to overestimated effect sizes for fluctuations on K due to this point.

To evaluate the effect of P and N pulses in the bioassay we calculated the log-response ratio (LRR) for each treatment as $\ln(\text{OD}_f/\text{OD}_i)$, where OD_f is the optical density at the end of the bioassay and OD_i is the initial optical density (after the nutrient addition). Differences in the LRR between temperature treatments were analyzed performing one-way ANOVAs considering the data for N and P addition separately. Because there were no differences between temperature treatments ($F = 0.0$, $P = 0.994$ for P addition; and $F = 0.0$, $P = 0.891$ for N addition), linear regressions were used to analyze the N and P supply effect on the LRR.

The final experiment sampling was processed to analyze the phytoplankton community (one of the two replicates were randomly selected and counted, $n = 50$). Richness and the inverse Simpson diversity (i.e., effective number of species) were calculated for each sample as diversity indices. GLMs including two- and three-way interactions were performed to analyze the effects of nutrient supply and temperature regimes on diversity. Analysis of similarity (ANOSIM) using Bray-Curtis index with 999 permutations was conducted to test differences in the phytoplankton community composition between constant and fluctuating temperature. A Simper test was applied to identify species contribution to the dissimilarity. As phytoplankton abundance (number of individuals per liter) was positively correlated with particulate C (Spearman correlation, $r = 0.75$, $P < 0.001$) we used the abundance of species for the composition analyses. Rare species, defined as species found in only one sample with less than five individuals, were excluded from the ANOSIM.

P and N supply concentrations were used as continuous variables and were log-transformed for the analyses to avoid an overestimation of the effect at the highest concentrations. All graphs and statistical analyses were performed using R version 3.5.0 (R Development Core Team 2018).

RESULTS

Growth rate

The phytoplankton growth rate (r_{\max}) was affected by supplied nitrogen (N) and phosphorus (P) concentrations as well as the temperature condition (significant three-way interaction; Table 2). Phytoplankton r_{\max} increased with increasing P supply, but decreased slightly with increasing N supply (Fig. 2; significant $\log(\text{N}) \times \log(\text{P})$, $\log(\text{N})$ and $\log(\text{P})$ effects, Table 2; Appendix S1: Fig. S2). Furthermore, temperature fluctuations decreased r_{\max} in comparison with constant conditions (Fig. 2, Table 2; Appendix S1: Fig. S2). The highest r_{\max} values were found under constant conditions at the highest P and intermediate N levels and an N:P supply ratio ~ 13 (optimal nutrient supply; Fig. 2a, significant temperature $\times \log(\text{P})$ interaction; Table 2). In the fluctuating temperature treatments, r_{\max} increased with the P supply level, independent of the N concentrations (Fig. 2b).

Compared to optimal nutrient conditions (where r_{\max} showed the highest values under constant temperature), we found that suboptimal nutrient supply changed the quantitative response of r_{\max} (effect size) to temperature fluctuations. The greatest relative decrease in r_{\max} because of temperature fluctuations was observed for optimal nutrient supply conditions, and remained high under intermediate N:P supply ratios (~ 16 – 30 ; Fig. 3a; Appendix S1: Fig. S3). However, the effect size of temperature fluctuations on r_{\max} declined at extreme N:P supply ratios. Based on the relative change in r_{\max} with temperature regimes across the nutrient conditions, we inferred that in comparison with optimal nutrient supply: (1) suboptimal nutrient supply decreased r_{\max} and thermal breadth, maintaining similar effect size of temperature fluctuations under intermediate N:P supply ratios (Fig. 1a, d); and (2) suboptimal nutrient supply decreased r_{\max} but thermal breadth remained unchanged, reducing temperature fluctuation effect size in extreme N:P supply ratios (Fig. 1b, d). The results did not show any case in which the temperature regime effect size on r_{\max} was

TABLE 2. ANOVA of supply nutrient concentrations (N and P) and temperature (Temp) effects on phytoplankton growth rate (r_{\max}), potential maximum biomass (K), richness (R) and the effective number of species (Inverse Simpson diversity, $1/D$).

Effect	Df	r_{\max}		Log(K)		Log R		$1/D$	
		F	P	F	P	F	P	F	P
Temp.	1	126.5	<0.001*	24.8	<0.001*	25.3	<0.001*	0.1	0.810
Log(N)	1	5.4	0.025*	44.9	<0.001*	1.0	0.332	0.0	0.980
Log(P)	1	128.6	<0.001*	18.5	<0.001*	0.1	0.770	0.4	0.548
Temp \times log(N)	1	1.5	0.228	1.0	0.332	3.7	0.062	0.2	0.624
Temp \times log(P)	1	11.7	0.001*	3.1	0.088	0.0	0.938	6.6	0.014*
Log(N) \times log(P)	1	4.4	0.042*	11.3	0.002*	0.0	0.864	0.2	0.648
Temp \times log(N) \times log(P)	1	7.26	0.010*	2.0	0.163	0.1	0.774	0.0	0.858

Note: N, P, R, and K were log-transformed for the analyses.

*Significant differences ($P < 0.05$).

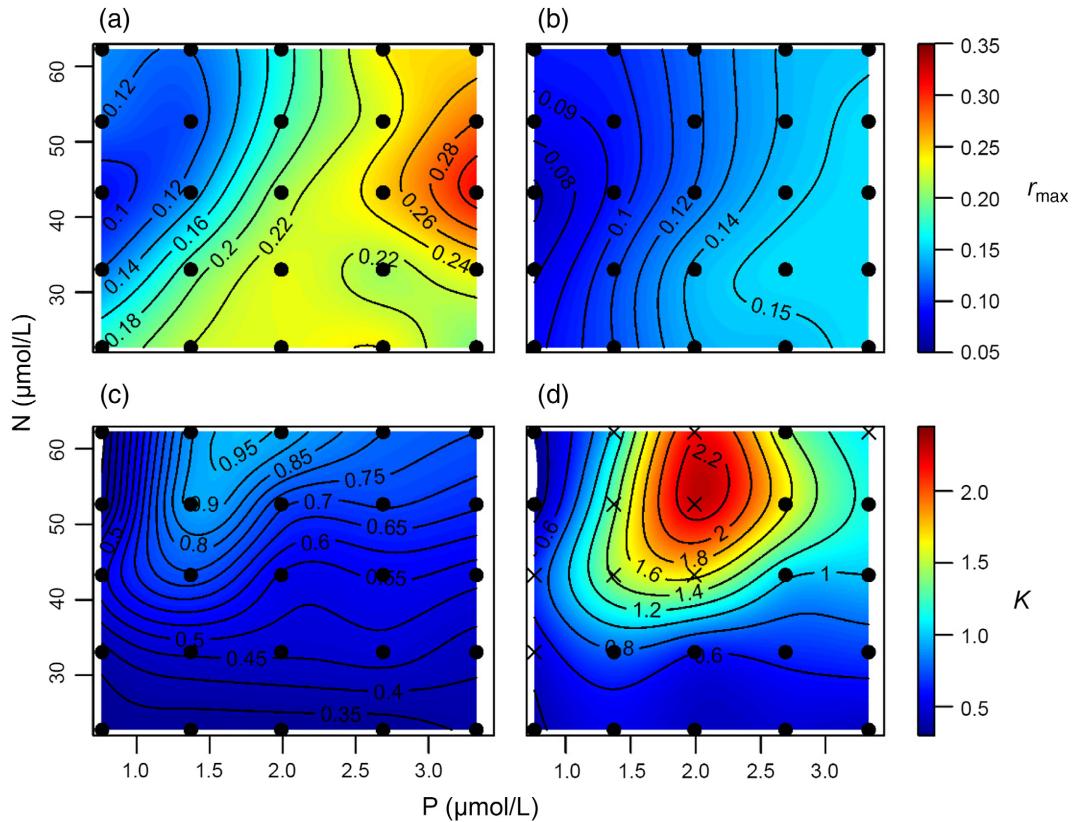


FIG. 2. Thin-plate spline regression response surfaces of the effects of phosphorus and nitrogen on (a), (b) phytoplankton exponential growth rate (r_{\max} , d^{-1}) and (c), (d) potential maximum biomass (K , based on optical density) under (a), (c) constant and (b), (d) fluctuating temperature regimes with equal mean temperature values. Circles: experimentally derived parameter estimates used for the thin-plate spline regressions. Crosses: nonsignificant parameter estimates (see Appendix S1: Table S1).

higher under suboptimal nutrient supply than under optimal nutrient conditions (Fig. 1c, d).

Potential maximum biomass

Increased N and P supply concentrations had a positive effect on the potential maximum phytoplankton biomass (K), with the effect of N supply being more pronounced than P (Fig. 2; significant $\log(N) \times \log(P)$, $\log(N)$ and $\log(P)$ effects, Table 2; Appendix S1: Fig. S2). For both temperature treatments, the highest K was found at similar nutrient concentrations (intermediate P and high N levels), but the N:P supply ratio resulting in maximum K was higher for constant temperature (Fig. 2c, d). Fluctuations in temperature increased K in comparison with constant conditions, and noninteractive effects between temperature regimes and nutrient concentrations were found (Table 2). Greatest effect size of temperature fluctuations on K was found in similar nutrient conditions as for the maximal K values (Fig. 3b; Appendix S1: Fig. S3). Because K was overestimated in some of the treatments, we compared the results obtained for K with the realized biomass during the stationary growth phase measured as particulate C and

chlorophyll a , which resulted in very similar conclusions than obtained for K (see Appendix S1: Table S2; Appendix S1: Fig. S2, and Discussion).

Stoichiometry

The phytoplankton N:P ratios reflected the nutrient supply, increasing with increasing N concentration and decreasing with higher P concentration (Fig. 4; Appendix S1: Table S2, Appendix S1: Fig. S4). However, stoichiometric responses of the phytoplankton community to nutrient concentrations were affected by temperature conditions and the growth phase (significant $\log(N) \times \log(P) \times \text{phase}$ and temperature $\times \log(P) \times \text{phase}$ interactions; Appendix S1: Table S2, Appendix S1: Fig. S4). In comparison to the exponential phase, phytoplankton N:P ratios were higher in the stationary growth phase, where the positive N effect and the inverse P effect were more pronounced (Fig. 4, significant $\log(N) \times \text{phase}$ and $\log(P) \times \text{phase}$ interactions; Appendix S1: Table S2). Consequently, the phytoplankton N:P ratios showed a higher mean and higher variance at the stationary than in the exponential phase (26.2 ± 10.2 and 18.4 ± 6.9 , respectively). During the exponential phase, the phytoplankton

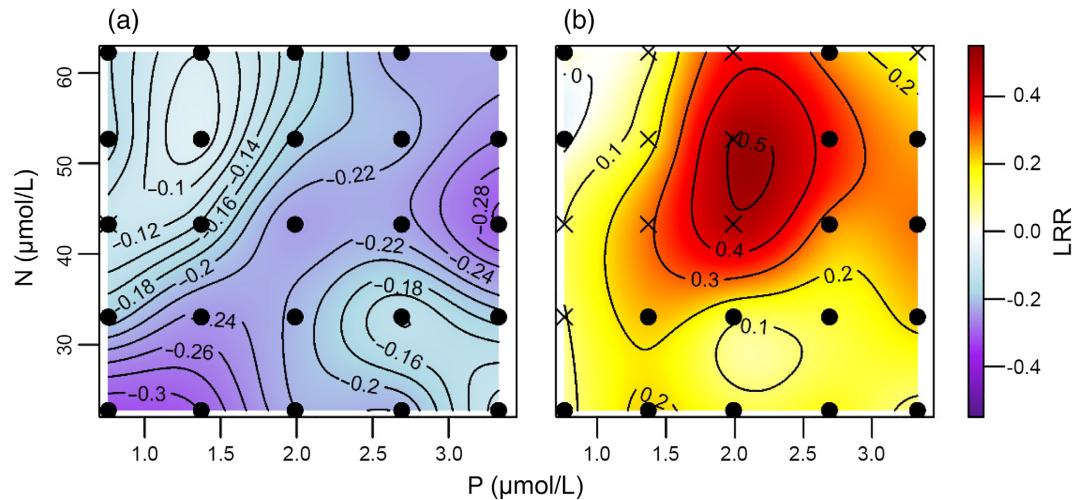


FIG. 3. Thin-plate spline regression response surfaces of nutrient-dependent variance effect sizes (\log_{10} -response ratio, LRR) on (a) phytoplankton exponential growth rate (r_{\max} , d^{-1}) and (b) maximum potential biomass (K , based on optical density). Circles: experimentally derived parameter estimates used for the thin-plate spline regressions. Crosses: nonsignificant parameter estimates (see Appendix S1: Table S1).

N:P ratios increased linearly with the N:P supply ratio; however, the slope of this increase was below the 1:1 line (Fig. 4a). After reaching the stationary phase the phytoplankton N:P equaled the supply ratio until a value of ~ 40 , but remained relatively constant for higher values (Fig. 4b). The temperature fluctuations resulted in lower phytoplankton N:P ratios than at constant temperature in the exponential phase, but this difference decreased in stationary phase (Fig. 4, significant temperature \times phase interaction, Appendix S1: Table S2).

The additional bioassays we performed showed that the phytoplankton communities were N, P, and colimited according to the N:P supply ratios at the beginning of the experiment, but colimitation was present in a wide range of ratios. The effect (LRR) of nutrient pulses did not show differences among temperature conditions ($F = 0.0$, $P = 0.994$ for P addition; and $F = 0.0$, $P = 0.891$ for N addition). In the N-addition treatments, the LRR increased with P (log-transformed; $t = 4.7$; $P < 0.001$) showing the highest values at lowest N and highest P levels (significant $\log(N) \times \log(P)$ effect, $t = -4.7$; $P < 0.001$; Appendix S1: Fig. S5). The opposite was found in the P-addition treatments, where the LRR increased with N (log-transformed, $t = 5.2$; $P < 0.001$) and decreased with P (log-transformed, $t = 3.2$; $P = 0.002$ and significant $\log(N) \times \log(P)$ interaction, $t = -4.0$; $P < 0.001$; Appendix S1: Fig. S5).

Phytoplankton composition and diversity

The phytoplankton abundance was dominated by chlorophytes, which represented between 88 and 98% of the total abundance across all treatments. Although the ANOSIM performed for the species abundance between

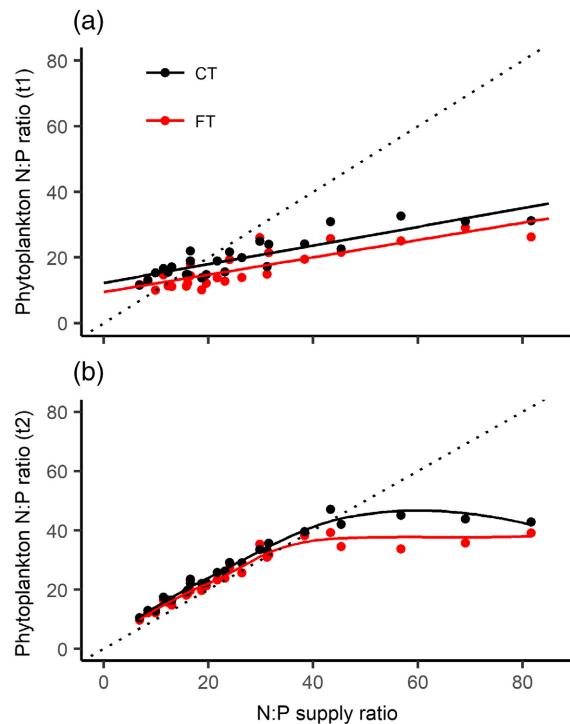


FIG. 4. Phytoplankton cellular N:P ratios across the N:P supply ratios at (a) the exponential (t1) and (b) stationary (t2) growth phase for constant (CT, black) and fluctuating (FT, red) temperature conditions. The dotted lines indicate the 1:1 line in each case.

the temperature conditions showed significant effects, the dissimilarity was low ($R = 0.13$; $P = 0.005$) indicating that the difference in temperature treatments had a

small effect on the species composition. Abundance variations of the three dominant chlorophytes (*Monoraphidium contortum*, *Dictyosphaerium pulchellum*, unicellular chlorophyta) explained 76% of the differences (Simper test; Appendix S1: Fig. S6).

The diversity indices of the phytoplankton community were affected by temperature treatments and P supply (Table 2). The effective number of species decreased with increasing P concentration at constant temperature (significant temperature \times log(P) interaction, Table 2). Richness increased with temperatures fluctuations but was not affected by nutrient concentrations (Table 2). The mean richness was 35.1 (SD = 4.01) for constant temperature treatments and 40.1 (SD = 2.6) for fluctuating temperature treatments, and the effective number of species mean was 3.7 (SD = 0.5) and 3.7 (SD = 0.6), respectively.

DISCUSSION

Our results showed that without changes in average temperature, fluctuating temperature alters phytoplankton growth rate, biomass, and stoichiometry. Phytoplankton community r_{\max} decreased under fluctuating temperature conditions, but the nutrient conditions influenced the effect size of thermal fluctuations on r_{\max} , suggesting alterations of the community-wide TRN shape. Similarly, the nutrient context affected K sensitivity to thermal variability, but contrary to r_{\max} , K was higher under fluctuating conditions. Responses to temperature treatments of phytoplankton stoichiometry supported the idea that cellular nutrient allocation can be adjusted according to the temperature regime. Despite the physiological responses, small changes in phytoplankton species composition suggest that these responses cannot be explained to a large extent by variation in species composition.

The observed decrease in r_{\max} with temperature fluctuations suggests that variability in temperature implies a cost in performance (Vasseur et al. 2014). The characteristic nonlinear shape of TRNs causes temperature fluctuations to decrease an organism's performance when the thermal variance reaches the concave region of the curve, which occurs above the inflection point and becomes stronger closer to and above the optimum (Bernhardt et al. 2018b). Our experiment was conducted at an average temperature of around 9°C, whereas the temperature at the initial sampling location was 2°C. Given the difference between in situ and experimental temperature, we assume that our temperature treatments were above the inflection points of the community growth TRN where the convex low-temperature response switches to a concave response curve. Under this assumption, Jensen's inequality (Jensen 1906, Ruel and Ayres 1999) predicts that temperature fluctuations should decrease the community r_{\max} compared to constant temperatures. Our experimental observations are in line with this prediction, suggesting that Jensen's

inequality can predict the direction of thermal variability effects not only at the population level (Bernhardt et al. 2018b), but also at the community level.

It should be noted that temperature fluctuations do not necessarily have negative effects on r_{\max} . Fluctuations can improve r_{\max} if they lie within the lower convex region the TRN (Bernhardt et al. 2018b). Alternatively, fluctuations might be neutral for r_{\max} if they lie within the linear region around the inflexion point or at the upper end of the TRN (Bernhardt et al. 2018b).

According to H1, the effect size (LRR) of thermal variability on r_{\max} was shaped by nutrient conditions. At the single phytoplankton species level, recent results suggest that low nutrient concentrations strongly decrease thermal optimum and tolerance (Thomas et al. 2017). In this case, nutrient supply limitation would increase sensitivity to fluctuating temperatures (as proposed in Fig. 1c). Our results depict a more complex image at the community level. The growth performance of the studied community was more sensitive to temperature variability when nutrient supply was balanced (moderate N:P supply ratios) across a wide range of nutrient concentrations (Fig. 1a). Contrary, under extreme unbalanced N:P supply ratios the relative response of r_{\max} to temperature fluctuations was small (Fig. 1b). Interestingly, nutrient concentrations per se only marginally affected sensitivity to thermal fluctuation as long as their ratio remained balanced.

Nevertheless, P and N concentrations had an important effect on phytoplankton r_{\max} and K associated with the respective growth phases. Phytoplankton r_{\max} was positively influenced by P supply and K by N supply, indicating growth rate limitation by P and yield limitation by N. This conclusion is in concordance with the GRH (Sterner and Elser 2002) and previous findings (Klausmeier et al. 2004a) which propose an increase of organismal P concentration with higher growth rate for P-rich ribosomes, while greater N allocation to protein synthesis are required when biomass is generated under stationary growth. Consequently, phytoplankton N:P ratios were higher and showed more flexibility in the stationary than in the exponential phase, reflecting the N:P supply (Klausmeier et al. 2004b, Hillebrand et al. 2013). However, even in the stationary phase, phytoplankton stoichiometry seemed to be restricted by physiological limits, becoming saturated with increasing N:P ratios (Hall et al. 2005). This result suggested the presence of an elemental threshold ratio at higher N:P supply ratios (~40) that may reflect strict P-limitation as has been shown in terrestrial plants (Sardans et al. 2012). The bioassay showed that phytoplankton biomass increase was either P, N, or colimited according, in general, to the N:P supply ratios across the nutrient levels and temperature conditions (Appendix S1: Fig. S5). We found that N additions spiked biomass production across a variety of N and P supply rates, which indicates that at least some of the algae in the community were N-limited even under generally more P-limiting conditions. This

reflects previous studies in colimitation, which, however, tended to find more frequent colimitations in systems with low nutrient content (Harpole et al. 2011).

Constant temperature increased phytoplankton N:P ratios in comparison with temperature fluctuations. This suggests that although temperature fluctuations decreased r_{\max} , more P in relation to N was allocated in the phytoplankton cells in comparison with constant conditions. The earlier exponential growth and N incorporation under constant temperature in the exponential phase probably influenced the higher phytoplankton N:P ratios (data not shown).

Previous studies showed increases in phytoplankton N:P ratios with increasing temperature (de Senerpont Domis et al. 2014, Yvon-Durocher et al. 2015). These results support the theory that more P-rich ribosomes are needed in colder systems to compensate for lower efficiency in protein synthesis, decreasing the N:P ratios (Toseland et al. 2013). Nevertheless, the positive relation between temperature and N:P ratios was not always found (Hessen et al. 2017, Skau et al. 2017). The mechanisms and patterns of phytoplankton stoichiometric responses to temperature are not clear, but our results support the idea that cellular stoichiometry reflects the effects and interactions between temperature regimes, nutrient supply, and growth phase.

As expected in H2, fluctuating temperatures increased phytoplankton K in comparison with constant conditions, and the effect size of temperature fluctuations changed across the nutrient gradient. It has been shown in phytoplankton populations that the carrying capacity decreases nonlinearly with increasing temperature (Bernhardt et al. 2018a). According to that, our results suggest that nonlinear averaging could be applied to predict K responses to thermal variability, but more studies are needed to test this. Savage et al. (2004) proposed that warmer temperatures increase maximum population growth rates and reduce the carrying capacity as a consequence of the higher metabolic rates (metabolic scaling theory; Brown et al. 2004). Our data showed similar patterns between temperature regimes, suggesting that similar compensatory mechanisms could operate in this case, causing constant temperature to present lower K but higher r_{\max} than fluctuating temperature conditions. As r_{\max} , K tended to increase the sensitivity to temperature variability when nutrient supply was balanced, but presenting a positive response to temperature fluctuations.

The analyses of K responses in our study were limited because not all the treatments were totally saturated at the end of the experiment generating an overestimation in some K values (Appendix S1: Table S1). For that reason, we compared K (reflecting maximum potential biomass) with the realized phytoplankton biomass measured as chlorophyll a and particulate C in the stationary growth phase. In concordance with the temperature regimes and nutrient supply effects on K , chlorophyll a increased with temperature fluctuations and N supply (more than P supply; Appendix S1:

Table S2; Appendix S1: Fig. S2). But, particulate C was higher under constant conditions and increased with N and P supply (Appendix S1: Table S2; Appendix S1: Fig. S2). These differences likely reflect a faster response of chlorophyll a to environmental changes in comparison to particulate C. This was supported by changes in the chlorophyll a response to temperature regimes and nutrient supply between growth phases in our experiment. Constant temperature and P supply increased chlorophyll a during the exponential growth phase, whereas the opposite was shown in the stationary growth phase. In concordance with previous results, chlorophyll a responses reflected the interactive effects of nutrient concentrations and temperature on phytoplankton biomass (Kraemer et al. 2017; Appendix S1: Table S2; Appendix S1: Fig. S2).

Even though the average temperature did not change in our study, temperature fluctuation had an effect on the physiological responses of the phytoplankton communities. This is in concordance with the idea that phytoplankton species can acclimate to short-term temperature variations by changing biochemical composition (Woods et al. 2003) and the balance between photosynthesis and respiration (Staehr and Birkeland 2006). Our results showed that the phytoplankton biomass responses to temperature conditions depended on the estimator used (chlorophyll a , particulate C, K based on optical density) and on the growth phase, suggesting that these factors should be considered in the interpretation of experimental results. This is relevant, taking into account that experiments are carried out using variable phytoplankton biomass estimations, growth conditions, and considering that some microcosms experiments are sampled only once because they are volume limited.

Small differences in species composition between treatments found in the analyzed communities suggest that the phytoplankton physiological responses are not simply reflecting different community composition. Whereas we conducted this experiment with a single inoculum, the high diversity of the assemblage and the lack of strong shifts in composition reflects a general physiological response rather than one driven by species identity. Similarly, de Senerpont Domis et al. (2014) found changes in phytoplankton biomass and stoichiometry in response to different temperature and P concentration without a change in phytoplankton composition. Related to this, Thomas et al. (2012) showed that adaptation to local temperature conditions is not highly phylogenetically constrained. As our experiment was a short-term experiment, it is reasonable to expect that changes in phytoplankton community composition would need more time to be detected. Also, changes in species composition were restricted in the closed experimental units to the initial species composition (site-dependent effect), which had, for example, low cyanobacteria presence (Striebel et al. 2016, Kraemer et al. 2017).

Higher richness was found under fluctuating temperature in comparison with constant temperature. This is in concordance with the idea that environmental fluctuations may induce species coexistence (Descamps-Julien and Gonzalez 2005). However, the effective number of species did not differ between temperature treatments, suggesting that although temperature fluctuations altered richness, they did not influence the community evenness. Additionally, the decrease in the effective number of species with increasing P supply that occurred only under constant temperature shows that temperature patterns can be influenced by interactions with nutrient concentrations.

Strong interactive effects of temperature regimes and nutrient supply on phytoplankton community growth rate, biomass, and stoichiometry show the relevance of considering these factors to predict phytoplankton development. Our results suggest that in systems with high daily temperature variation phytoplankton community growth and biomass may differ accordingly to the nutrient context. Detected changes in stoichiometry at the community level implies a differential resource allocation according to the temperature regimes. These changes in nutrient ratios can affect the nutrient fluxes into trophic levels. Interactive effects of temperature regimes and nutrient conditions observed in this study may also have implications for the analysis of phytoplankton performance in different climatic regions, and has potentially large consequences under predicted changes in environmental variability under global change.

ACKNOWLEDGMENTS

We thank Dorothee Hodapp for the statistical advice and Patrick Thomas for the comments that helped to improve this paper. We thank Silvia Heim, Alexander Wacker, Heike Rickels, and Andrea Gall for technical support. MG was supported by the Uruguayan Agency of Investigation and Innovation (ANII) and the German Academic Exchange Service (DAAD). MS, HH, and AMK acknowledge funding by the German Research Foundation (DFG: STR 1383/1-1, STR 1383/6-1, HI 848/15-1, HI 848/24-1 within the priority program DynaTrait SPP 1704 and KO5330/1-1).

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