

DIATOM GROWTH RESPONSES TO PHOTOPERIOD AND LIGHT ARE PREDICTABLE FROM DIEL REDUCTANT GENERATION¹

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Light drives phytoplankton productivity, so phytoplankton must exploit variable intensities and durations of light exposure, depending upon season, latitude, and depth. We analyzed the growth, photophysiology and composition of small, *Thalassiosira pseudonana*, and large, *Thalassiosira punctigera*, centric diatoms from temperate, coastal marine habitats, responding to a matrix of photoperiods and growth light intensities. *T. pseudonana* showed fastest growth rates under long photoperiods and low to moderate light intensities, while the larger *T. punctigera* showed fastest growth rates under short photoperiods and higher light intensities. Photosystem II function and content responded primarily to instantaneous growth light intensities during the photoperiod, while diel carbon fixation and RUBISCO content responded more to photoperiod duration than to instantaneous light intensity. Changing photoperiods caused species-specific changes in the responses of photochemical yield (e^-/photon) to growth light intensity. These photophysiological variables showed complex responses to photoperiod and to growth light intensity. Growth rate also showed complex responses to photoperiod and growth light intensity. But these complex responses resolved into a close relation between growth rate and the cumulative daily generation of reductant, across the matrix of photoperiods and light intensities.

Key index words: cell size; diatom; electron transport; growth; photoperiod; photosystem II; RUBISCO; *Thalassiosira*

Abbreviations: ANOVA, Analyses of Variance; C:N, carbon-to-nitrogen; $F_0'_{2s}$, base line fluorescence in the light adapted state; F_0 , base line fluorescence in darkness; $F_M'_{2s}$, maximal fluorescence in the light adapted state; F_M' , maximal fluorescence under growth light illumination; F_M , maximal fluorescence in darkness; FRR, Fast Repetition Rate; F_s , steady state fluorescence under growth light intensity; K_L , saturated growth light intensities; L:D, Light:Dark; PSII, Photosystem II; RbcL, large protein subunit of the Ribulose-1,5-bisphosphate Carboxylase Oxygenase enzyme; RUBISCO, Ribulose-1,5-bisphosphate Carboxylase Oxygenase; μ , cell specific exponential growth rate; μ_{max} , maximum cell specific exponential growth rate; σ_{PSII}' , effective absorbance cross-section serving Photosystem II photochemistry

The effects of light intensity on the photosynthetic responses and growth rates of diatoms and other phytoplankton have been well studied (Platt and Jassby 1976, Falkowski et al. 1985, Falkowski and Raven 2007). Interactions between light intensity and nutrient limitation have also been explored (Shuter 1979, Laws and Bannister 1980, Sakshaug et al. 1989, Geider et al. 1998, Pahlow 2005, Loeb1 et al. 2010, Halsey and Jones 2015). Furthermore, the effects of fluctuating or oscillating light have been studied physiologically (Litchman 2000, 2003, Mackenzie and Campbell 2005, Wagner et al. 2005, Key et al. 2010, Orefice et al. 2016) and through competition models (Huisman et al. 1999a,b, Litchman and Klausmeier 2001, Litchman et al. 2004).

Beyond instantaneous light intensity, the diel light:dark cycle is a significant environmental variable; the diel light:dark cycle can regulate phytoplankton metabolic processes through the influences of photoperiod or of night length, maximum

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irradiance or spectral composition, or the oscillation pattern (Prézelin 1992, Orefice et al. 2016). Photoperiod varies seasonally at temperate to high latitudes, from total darkness during the high latitude winter, to constant illumination during the summer solstice. At mid-latitudes, photoperiod waxes and wanes through the year on top of shorter term changes in light intensity imposed by mixing. Photoperiod length interacts with light intensity (Brand and Guillard 1981) and with the oscillation pattern (Orefice et al. 2016) to modulate photosynthesis and metabolism in diatoms, whose photosynthesis involves tight coupling between chloroplasts and mitochondria (Prihoda et al. 2012, Bailleul et al. 2015). Yet, surprisingly few studies (Chisholm and Costello 1980, Brand and Guillard 1981, Tilzer and Dubinsky 1987, Nielsen 1997, Tang and Vincent 2000, Walter et al. 2015) have systematically examined the effects of photoperiod on marine phytoplankton metabolism and physiology. In contrast, a recent meta-analysis attempted to extrapolate phytoplankton growth responses to light intensity from data derived from studies with different applied photoperiods (Edwards et al. 2015).

Growth, photosynthetic, and adaptive traits are strongly influenced by organism size (McMahon and Bonner 1983, Peters 1983, Kooijman 1986, West et al. 1997, Marañón et al. 2007, 2013). Diatoms span a wide range of sizes (Beardall et al. 2009) and their metabolism and photophysiology show predictable patterns across related species of increasing size (Finkel 2001, Finkel et al. 2004, Key et al. 2010, Wu et al. 2014a,b). At least two broad categories of mechanisms conceptually connect organism size, photoperiod, light intensity, and growth rate. First, small cell volume could limit capacity for carbon storage of carbohydrate and lipid, resulting in limited flexibility in cellular carbon-to-nitrogen (C:N) ratios (Ross and Geider 2009), and limited energy reserves to fuel respiration to support dark synthesis of proteins and structural apparatus (Geider and Osborne 1989, Talmy et al. 2014). Alternately, cell size dependent differences in the coupling of reductant generation to retained carbon assimilation (Suggett et al. 2009, Halsey et al. 2010, 2013) could cause differences in growth rate responses to light intensity and photoperiod.

To explore the interacting effects of diatom cell size and photoperiod on the light responses of growth rate and physiology, we selected two coastal diatom strains differing by ~ 4 orders of magnitude in cell biovolume, *Thalassiosira pseudonana* ($\sim 40 \mu\text{m}^3$), and *Thalassiosira punctigera* ($\sim 300,000 \mu\text{m}^3$), and grew them under a matrix of different photoperiods and light intensities. To explore the underlying mechanisms linking growth rate with diatom cell size, photoperiod, and light intensity, we measured cellular carbon and nitrogen contents, Photosystem II (PSII) electron transport, cellular PSII content, Ribulose-1,5-bisphosphate Carboxylase Oxygenase (RUBISCO) activity and

cellular RUBISCO content. Our experiments led us to ask how cell size and photoperiod interact to influence the coupling of photosynthetically generated reductant to carbon assimilation and to growth.

MATERIALS AND METHODS

Culture protocol and growth rate. We semicontinuously cultured two strains of coastal, temperate, marine centric diatoms, *T. pseudonana* (CCMP 1335; $\sim 40 \mu\text{m}^3$) and *T. punctigera* (CCAP 1085/19) ($\sim 300,000 \mu\text{m}^3$) obtained from the Provasoli-Guillard National Center of Marine Phytoplankton, with f/2 medium (Guillard and Ryther 1962) in 2 L polycarbonate bottles (Nalgene, USA) at 18°C. During cultivation, the culture bottles were manually shaken 2–3 times a day and arbitrarily distributed in a growth chamber. Light in the chamber was provided by fluorescent tubes (Sylvania 17W T8 4100K) automatically turned on at 08:00 and maintained continuously (24:0 Light:Dark (L:D) cycle) or turned off at 12:00 (4:20 L:D cycle), 16:00 (8:16 L:D cycle), or 24:00 (16:8 L:D cycle) in a square-wave pattern (Orefice et al. 2016). Cultures under all four L:D cycles were grown under 75, 150, and 300 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$, measured with a microspherical quantum sensor (US-SQS; Waltz, Germany) submersed in a culture bottle filled with seawater. In addition, 24:0 L:D cultures were grown under 25 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$, to achieve light limitation of growth rate under continuous light, where growth rate was already nearly saturated at 75 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$. 4:20 L:D cultures were also grown under 600 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$, because 300 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ did not saturate growth rate under this short photoperiod. We grew two or three replicate culture bottles of each species under each combination of light intensity and photoperiod, for a total of 60 separate semi-continuous cultures.

To track the growth of small *T. pseudonana*, we measured fluorescence from culture samples with excitation of 440 nm and emission at 650 nm at 10:00 a.m. every day (Molecular Devices SpectraMax Gemini EM), before and after dilutions with sufficient fresh media to maintain cultures between 110 and 220 ng Chlorophyll *a* $\cdot \text{mL}^{-1}$. To track the growth of larger *T. punctigera*, we took duplicate 2 mL samples at 10:00 a.m. and fixed in Lugol's acid solution, and then counted cells with a Sedgwick Rafter chamber under an inverted microscope. The cell density for *T. punctigera* was maintained at 676–1,100 cells $\cdot \text{mL}^{-1}$, equivalent to 60–114 ng Chlorophyll *a* $\cdot \text{mL}^{-1}$. The cell specific exponential specific growth rate was estimated as:

$$\mu = [\ln(N_t) - \ln(N_0)]/\Delta t$$

where N_t is the culture fluorescence or cell number at time t , N_0 is the fluorescence or cell number at time 0, and Δt is the time elapsed from time 0 to time t . At least 15 transfers of semi-continuous dilution were done for cultures growing under each regime of photoperiod and light intensity, resulting in more than 12 cellular generations before sampling.

Sampling protocol. Samples to measure the chlorophyll fluorescence induction and relaxation kinetics, RUBISCO, carbon and nitrogen contents were taken at two hours prior to dawn (06:00), immediately prior to dawn (08:00), then every 2 h within the illuminated period, and again two hours after lights off. In total, nine sample time points were taken from the 16:8 L:D treatment, seven from the 8:16 L:D treatment, and five from 4:20 L:D treatment. Samples were taken at 06:00, 10:00, and 18:00 for the 24:0 L:D treatment. At each sampling point, the 2 L bottle was gently mixed to disperse the cell suspension prior to withdrawal of samples.

Fluorometer measurements. At each time point a 2 mL sample taken from each 2 L bottle was incubated in darkness for 5 min within a cuvette with temperature control (18°C). Chlorophyll fluorescence kinetics was measured using the Fast Repetition Rate (FRR) fluorescence technique (FL 3500; Photon Systems Instruments, Czech Republic) (Kolber et al., 1998). A train of $40 \times 1.2 \mu\text{s}$ flashlets of blue light (455 nm ; $\sim 100,000 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$) were applied over $128 \mu\text{s}$ to progressively close PSII reaction centers. The sample was then allowed to relax over 2 s of darkness, followed immediately by a second flashlet train. This chlorophyll fluorescence induction protocol was then repeated after 2 min exposure to growth light intensity. We analyzed the resulting FRR induction curve with the PSIIWORX script (A. Barnett, sourceforge.net) for MATLAB software, and derived the base line fluorescence in darkness (F_0) or steady state fluorescence under growth light intensity (F_s), the maximal fluorescence in darkness (F_M), or under growth light illumination (F_M'), and the base line and maximal fluorescence in the light adapted state, $F_0'_{2s}$ and $F_M'_{2s}$, which were measured after 2 s of darkness following the fluorescence induction under growth light. We used the magnitude of any increase from F_M' to $F_M'_{2s}$ to apply a proportional correction to $F_0'_{2s}$ to estimate the actual level of F_0' prevailing under illumination (Oxborough and Baker 1997, Ware et al. 2015):

$$F_0' = F_0'_{2s} \times \{1 - [(F_M'_{2s} - F_M')/F_M'_{2s}]\}$$

We followed (Suggett et al. 2009, Huot and Babin 2010) to estimate the rate of reductant generation from PSII under the growth light intensity applied during the photoperiod, as:

$$\text{PSII electron transport} (e^- \cdot \text{PSII}^{-1} \cdot \text{s}^{-1}) = \sigma_{\text{PSII}}' \times I \times q_p$$

where $\sigma_{\text{PSII}}' (\text{A}^2 \cdot \text{quanta}^{-1})$ is the effective absorbance cross-section serving PSII photochemistry at growth light intensity I ($\text{photons} \cdot \text{A}^{-2} \cdot \text{s}^{-1}$), and $q_p, (F_M' - F_s)/(F_M' - F_0')$ is the proportion of PSII instantaneously open and ready to perform photochemistry under light intensity I . We estimated the functional pool of PSII centres competent for photochemistry, $\text{PSII}_{\text{active}}$, using a linear calibration of $F_0'/\sigma_{\text{PSII}}'$ compared with oxygen flash yield measures of $[\text{PSII}_{\text{active}}]$ from culture samples of *T. pseudonana* and *T. punctigera* (Oxborough et al. 2012, Silsbe et al. 2015).

We estimated the electron transport rate per total protein as:

$$e^- \cdot \mu\text{g protein}^{-1} \cdot \text{s}^{-1} = (e^- \cdot \text{PSII}^{-1} \cdot \text{s}^{-1}) \times (\text{PSII}_{\text{active}} \cdot \mu\text{g protein}^{-1})$$

Carbon and nitrogen analyses. At each sampling point 20 mL of culture was filtered onto a pre-combusted (5 h, 450°C) Whatman GF/F glass fiber filter (13 mm in diameter), followed by 10 mL of 50 mmol \cdot L⁻¹ HCl to dispel inorganic carbon on filters, which were then dried in an oven at 55°C for 12 h and stored in a desiccators for later analyses. Cellular carbon and nitrogen contents were measured with a Vario EL III Elemental Analyzer (Elementar, Hanau, Germany). In parallel with culture filtering equivalent 20 mL media volumes were filtered through pre-combusted GF/F filters. These blank filters were interspersed with sample filters during the elemental analyses. C of blank filters accounted for less than 5% of the total C in the culture samples while and N of blank filters accounted for less than 10% of total N in the culture filters. Filter blank determinations were subtracted from culture C & N determinations before further analyses.

Cell count, protein, and Chlorophyll a measurements. At each sampling point duplicate 2 mL samples were fixed in Lugol's

acid solution. The cell suspension density was then measured with a Coulter Z2 counter (Beckman Instruments, Mississauga, Canada) for *T. pseudonana* and with a Sedgewick Rafter chamber under an inverted microscope for *T. punctigera*. At the same time, 50 mL of culture was vacuum-filtered onto a binder-free Whatman GF/F glass fiber filter (25 mm in diameter), which was immediately flash frozen in liquid nitrogen and stored at -80°C until later analyses of proteins and chlorophyll.

Total protein was extracted from the frozen filters into 450 μL of denaturing extraction buffer (0.1375 mol TRIS \cdot L⁻¹ buffer, 0.075 mol Lithium Dodecyl Sulfate \cdot L⁻¹, 1.075 mol glycerol \cdot L⁻¹, 0.5 mmol EDTA \cdot L⁻¹, 0.1 mg Pefabloc \cdot mL⁻¹; Roche) (Brown et al. 2008). The filters were suspended in the extraction buffer and subjected to three cycles of 60 s at $6.5 \text{ m} \cdot \text{s}^{-1}$ to break up cells using a MPBio FastPrep[®]-24 instrument with bead lysing matrix D (MPBio SKU 116913050). We then measured the concentration of total protein in the extracts using a Bio-Rad DC protein assay kit (500-0116) with known BSG standards. We loaded extract containing 0.5 μg of total protein onto 4%–12% acrylamide precast NuPAGE gels (Invitrogen, ThermoFisher, Ottawa, Canada) in parallel with a range of 1,500, 750, 375, 187.5 fmol large protein subunit of the Ribulose-1,5-bisphosphate Carboxylase Oxygenase enzyme (RbcL) standard per standard lane (Agrisera, www.agrisera.se, AS01 017S) to establish a standard curve for determination of RUBISCO large subunit (RbcL) content in the samples. We ran electrophoresis for 25 min at 200 V and transferred proteins to a polyvinylidene fluoride membrane for 60 min at 30 V. After membrane blocking with 2% ($\text{g} \cdot \text{mL}^{-1}$) ECL Advance blocking reagent (GE Healthcare, Mississauga, Canada), we applied a primary antibody (Agrisera, AS03 037, 1:20,000 in 2% ECL Advance blocking dilution), then an anti-rabbit secondary antibody coupled with horseradish peroxidase (Agrisera, AS09 602, 1:20,000). After this, we developed the membranes with chemoluminescence using ECL Advance (GE Biosciences) and imaged under a CCD imager (BioRadVersa-Doc 4000MP). Finally, we quantified RbcL by fitting the sample signal values to the protein standard curves, taking care that all sample signals fell within the range of protein standard curve and that no band signals were saturated (Brown et al. 2008).

To measure chlorophyll *a*, 60 μL of the protein extract was added to 440 μL 90% acetone (v/v) saturated with magnesium carbonate; after 15 min extraction in the dark at 4°C and 2 min centrifugation (13,000 g), we measured the absorbance of the supernatant at 664, 630, and 750 nm using a UV/VIS photospectrometer (UV-1800, Shimadzu, Japan). We estimated chlorophyll *a* content (Jeffrey and Humphrey 1975) as:

$$[\text{Chlorophyll } a] = 11.47 \times (A_{664} - A_{750}) - 0.4 \times (A_{630} - A_{750})$$

Data analysis. We fit the response of growth rate to culture growth light intensity using the equation of (Eilers and Peeters 1988):

$$\mu = I/(a \times I^2 + b \times I + c)$$

where μ is cell specific exponential growth rate (d^{-1}) at the particular light intensity; I is the growth light ($\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$) and a , b and c are parameters to constrain the equation to the plot of measured values for μ across a range of growth light intensity. We calculated the maximum cell specific exponential growth rate (μ_{max}) as:

$$\mu_{\text{max}} = 1/(b + 2\sqrt{a \times c})$$

and saturated growth light intensities (K_L) as:

$$K_L = \sqrt{c/a}$$

Because of our experimental design, we lacked sufficient determinations of μ under low light intensities to reliably

estimate the more familiar K_E light saturation parameter (MacIntyre and Cullen 2005).

We estimated apparent carbon assimilation per RUBISCO active site ($C \cdot \text{RbcL}^{-1} \cdot \text{s}^{-1}$) as:

$$\begin{aligned} \text{RUBISCO turnover rate} (C \cdot \text{RbcL}^{-1} \cdot \text{s}^{-1}) = \\ (C \cdot \text{cell}^{-1}) \times (e^\mu - 1) \times (\text{cell} \cdot \text{RbcL}^{-1}) \\ \times (24\text{H}/\text{Photoperiod H}) \end{aligned}$$

where μ (units of s^{-1}) is cell specific exponential growth rate. ($24 \text{ H}/\text{Photoperiod H}$) assigns 24 h of diel C assimilation to RUBISCO activity during the photoperiod hours only.

We used ANOVA with Bonferroni or Tukey post-tests (Prism 5, Graphpad Software, La Jolla, CA, USA; RStudio 0.99.491) and comparisons of linear and non-linear curve fits to detect significant differences among cultures of each species from each L:D cycle and growth light intensity.

RESULTS

In the smaller diatom *T. pseudonana*, the response of cell specific growth rate (μ) to increasing growth light intensity (Fig. 1A) was strongly influenced by the length of the photoperiod. Under continuous 24:0 L:D, *T. pseudonana* showed a sharp rise in μ from low to moderately low light, followed by growth inhibition at $150 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ and above. As the photoperiod decreased, the response of growth to light intensity became

progressively weaker, and maximum growth rate μ_{max} decreased. Thus, under the shortest photoperiod (4:20 L:D), *T. pseudonana* showed little change in μ across the range of lights from 75 to $600 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ (Fig. 1A), although under this short photoperiod the cells were unable to reliably maintain growth at $25 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ (data not shown). The larger diatom *T. punctigera* showed an opposing pattern of growth responses to photoperiod; μ increased as photoperiod decreased, and μ reached higher values at $150 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ or above (Fig. 1B).

We chose combinations of growth light intensities and photoperiods to achieve equivalent cumulative photons per day from multiple combinations of light intensity and photoperiod. At subsaturating growth light intensities, *T. pseudonana* showed a positive, linear response of μ to cumulative photons per day ($r^2 = 0.90$; $F_{1,12} = 106.4$; slope significantly greater than 0, $P < 0.0001$; Fig. 1C). But at a given cumulative photon dose such as ~ 4 or $\sim 8 \text{ mol photons} \cdot \text{m}^{-2} \cdot \text{d}^{-1}$, μ varied widely, depending upon whether the light intensity was sub- or super-saturating for growth, which in turn depended upon the photoperiod over which, the photons were applied (Fig. 1C). In contrast, in *T. punctigera* μ responded primarily to photoperiod, with only modest growth rate responses to daily cumulative photon dose (Fig. 1D).

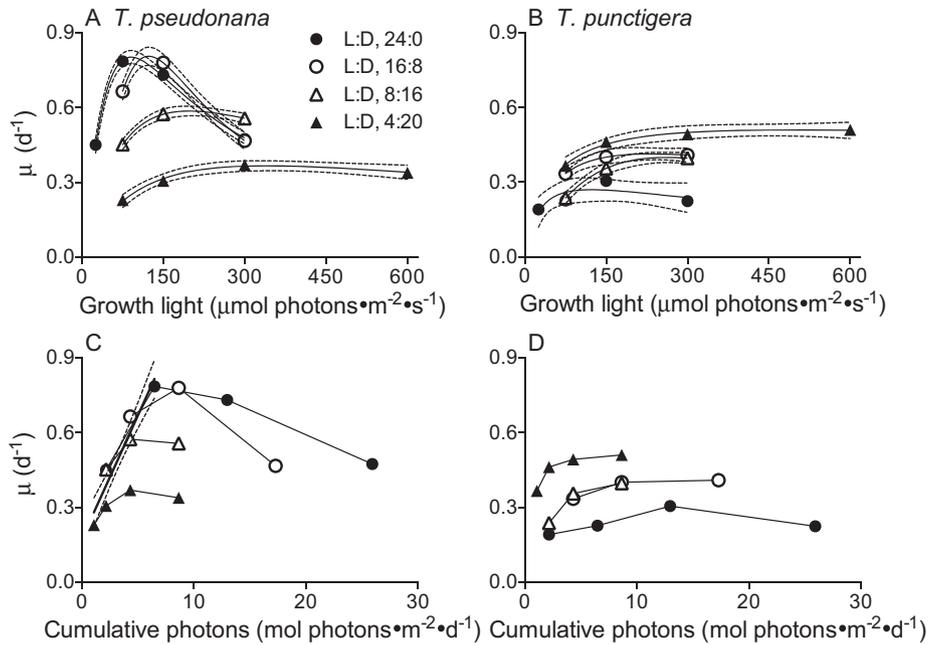


FIG. 1. Cell specific exponential growth rate (μ , d^{-1}) as a function of culture growth light intensity during the photoperiod ($\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$) (A, B) or of daily cumulative photons ($\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{d}^{-1}$) (C, D) for a small diatom *Thalassiosira pseudonana* (A, C) or a large diatom *Thalassiosira punctigera* (B, D) at light:dark (L:D) cycles of 24:0 (filled circles), 16:8 (open circles), 8:16 (open triangles), and 4:20 (filled triangles). Points show average growth rate determinations from two or three independently grown cultures; the range of replicated points falls within the symbol size. Solid lines in A and B show light intensity response curves of growth rate fitted with the equation of (Peeters and Eilers 1978, Eilers and Peeters, 1988). Bold line in C shows linear regression of pooled μ to daily cumulative photons under subsaturating growth light conditions; thin dashed lines show 95% confidence intervals on the fitted curves.

We used the curve fits in Figure 1, A and B to extract estimates of μ_{\max} , and then plotted them with 95% confidence intervals versus photoperiod (Fig. 2A). For *T. pseudonana* μ_{\max} showed a non-linear increase with increasing photoperiod ($r^2 = 0.99$, Absolute Sum of Squares 6.467×10^{-6} , Standard deviation of residuals = 0.002543, $df = 1$). In *T. punctigera* μ_{\max} under the shortest photoperiods overlapped with μ_{\max} from *T. pseudonana*, but as photoperiod increased μ_{\max} decreased for *T. punctigera* ($r^2 = 0.86$; $F_{1,2} = 12.24$; slope not significantly less than 0, $P < 0.09$) to levels well below the confidence intervals for μ_{\max} for *T. pseudonana* measured at equivalent photoperiods (Fig. 2A). Thus, the general pattern of decreasing maximal growth rates with increasing cell size within a taxonomic group (Finkel et al. 2004, Beardall et al. 2009) can actually reverse, depending upon the photoperiod. In both diatoms, the light saturation parameter K_L decreased with increasing photoperiod (two-way ANOVA, $F_{3,8} = 496.5$, $P < 0.0001$), but across the photoperiods *T. punctigera* showed consistently higher K_L (two-way ANOVA, $F_{1,8} = 234.2$, $P < 0.0001$) and thus required a higher light intensity to saturate growth rate. The smaller and larger diatoms thus show contrasting growth responses:

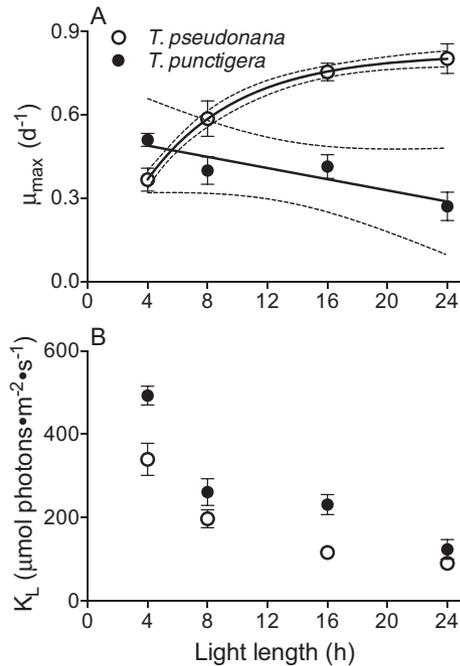


FIG. 2. Maximum cell specific exponential growth rate (A, μ_{\max} , d^{-1}) and saturation growth light (K_L) (B, $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$) derived from the growth versus irradiance curves (Fig. 1, A and B), plotted as a function of photoperiod for *T. pseudonana* (open circles) and *T. punctigera* (filled circles). Solid lines in A show single phase exponential rise to a plateau for *T. pseudonana*, and a linear decrease for *T. punctigera*; thin dashed lines show 95% confidence intervals on the fitted curves. Error bars on points show the 95% confidence intervals on the fitted parameters, derived from the curves in Figure 1.

T. pseudonana showed growth rate advantages under long photoperiods and light intensities below $150 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ (Fig. 1A); whereas *T. punctigera* showed growth advantages under short photoperiods and light intensities at or above $150 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ (Fig. 1B).

We sought to understand the mechanisms for these contrasting responses to light intensity and photoperiod by analyzing cell C and N, PSII, and carbon metabolism. The cell volume of *T. pseudonana* showed only limited responses to instantaneous growth light intensity or to photoperiod (Fig. S1A in the Supporting Information), but in *T. punctigera* cell volume increased under 24:0 L:D when growth was slowest (Fig. S1B). In *T. pseudonana*, average cellular carbon content increased at the lowest light intensity (Fig. S1C), but there was no obvious pattern of changing carbon content with instantaneous growth light intensity, nor with photoperiod in *T. punctigera* (Fig. S1D). These determinations of carbon contents represent averages over multiple determinations made across the diel cycles (Table 1). Interestingly, the combination of treatments that provoked the largest variation in diel C: N was the 4:20 L:D $600 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ treatment of *T. pseudonana* (Fig. S2G in the Supporting Information), the highest growth light intensity condition for the smaller cells.

We next examined the function and content of PSII, to assess the generation of reductant in the cells across diel cycles and light intensities. In *T. pseudonana*, the reductant generation rate per PSII showed a smooth response to instantaneous light intensity, peaking under $300 \mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$, with only modest effects of photoperiod on the pattern (Fig. 3A). *T. punctigera*, in contrast, showed a strong response of the reductant generation rate per PSII to photoperiod with suppression of reductant generation per PSII across all growth light intensities under a 24:0 L:D cycle, and accelerated reductant generation per PSII under the 4:20 L:D cycle and high light (Fig. 3B). The content of $\text{PSII}_{\text{active}}$ generally decreased with increasing growth light in *T. pseudonana* (Fig. 3C), but not in *T. punctigera* (Fig. 3D), with scattered effects of photoperiod in each species. Thus, PSII function and content respond primarily to instantaneous growth light intensity in *T. pseudonana*, and to both growth light intensity and photoperiod in *T. punctigera* (Fig. 3).

To assess the performance of assimilation of carbon into biomass we used cellular carbon content, cell specific growth rate (s^{-1}) and RbcL content to estimate a lower bound on achieved RUBISCO turnover rate. This lower bound estimator neglects any carbon that is fixed through RUBISCO but then respired or released to the media as dissolved organic carbon, which we did not measure. In both *T. pseudonana* (Fig. 4A) and particularly in

TABLE 1. Cellular contents (mean \pm SD) of carbon (C, $\text{pg} \cdot \text{cell}^{-1}$ or $\text{ng} \cdot \text{cell}^{-1}$), nitrogen (N, $\text{pg} \cdot \text{cell}^{-1}$ or $\text{ng} \cdot \text{cell}^{-1}$), mole C:N, chlorophyll a (Chl a, $\text{pg} \cdot \text{cell}^{-1}$ or $\text{ng} \cdot \text{cell}^{-1}$) and proteins ($\text{pg} \cdot \text{cell}^{-1}$ or $\text{ng} \cdot \text{cell}^{-1}$) for *T. pseudonana* and *T. punctigera*. These determinations represent averages over multiple determinations made across the diel cycles from two separate culture replicates for each condition. $n = 4-8$ depending upon condition.

Species	L:D cycles	Growth light		N ($\text{pg} \cdot \text{cell}^{-1}$)	C ($\text{pg} \cdot \text{cell}^{-1}$)	Mole C:N	Chl a ($\text{pg} \cdot \text{cell}^{-1}$)	Protein ($\text{pg} \cdot \text{cell}^{-1}$)	
		($\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$)							
<i>T. pseudonana</i>	24:0	25		16.3 ± 3.76	124 ± 25	8.90 ± 0.29	0.67 ± 0.025	7.74 ± 0.12	
		75		10.3 ± 0.04	69.4 ± 4.6	7.85 ± 0.48	0.28 ± 0.010	4.96 ± 0.25	
		150		10.0 ± 0.54	63.7 ± 2.1	7.40 ± 0.14	0.18 ± 0.015	4.19 ± 0.64	
	16:8	300		10.5 ± 1.19	65.2 ± 1.7	7.26 ± 0.62	0.13 ± 0.009	4.23 ± 0.53	
		75		10.7 ± 1.93	87.4 ± 21.3	9.51 ± 0.63	0.34 ± 0.071	5.04 ± 1.04	
		150		11.0 ± 3.11	72.7 ± 1.4	8.01 ± 2.12	0.21 ± 0.031	3.79 ± 0.28	
	8:16	300		7.53 ± 0.08	60.1 ± 1.6	9.28 ± 0.17	0.12 ± 0.001	4.03 ± 0.09	
		75		13.0 ± 0.77	85.4 ± 1.3	7.66 ± 0.31	0.46 ± 0.001	5.95 ± 0.01	
		150		14.0 ± 0.60	83.4 ± 4.6	6.93 ± 0.06	0.38 ± 0.004	5.63 ± 0.62	
	4:20	300		11.1 ± 0.03	65.9 ± 0.4	6.87 ± 0.07	0.23 ± 0.011	4.47 ± 0.04	
		75		16.3 ± 0.76	95.6 ± 9.2	6.81 ± 0.32	0.50 ± 0.012	7.50 ± 0.29	
		150		13.1 ± 0.80	83.5 ± 6.3	7.45 ± 1.02	0.37 ± 0.001	6.06 ± 0.03	
	<i>T. punctigera</i>	24:0	300		11.6 ± 0.49	74.1 ± 3.6	7.43 ± 0.41	0.33 ± 0.007	6.01 ± 0.19
			600		11.8 ± 0.54	90.1 ± 0.8	8.76 ± 0.52	0.34 ± 0.003	7.29 ± 0.96
			25		8.45 ± 0.34	42.5 ± 3.8	5.04 ± 0.65	0.53 ± 0.015	45.3 ± 8.63
16:8		75		10.4 ± 0.14	48.1 ± 0.1	4.66 ± 0.06	0.90 ± 0.097	44.3 ± 5.94	
		150		12.1 ± 0.23	58.6 ± 1.2	4.85 ± 0.02	1.21 ± 0.010	48.2 ± 0.03	
		300		10.2 ± 0.99	47.8 ± 4.7	0.69 ± 0.00	0.48 ± 0.044	35.3 ± 0.75	
8:16		75		7.76 ± 0.09	34.9 ± 0.3	4.50 ± 0.02	1.03 ± 0.022	28.4 ± 0.68	
		150		8.55 ± 0.66	39.1 ± 1.0	4.58 ± 0.22	1.28 ± 0.031	39.0 ± 1.77	
		300		10.3 ± 0.18	52.3 ± 0.6	5.10 ± 0.15	1.11 ± 0.012	50.0 ± 3.75	
4:20		75		9.44 ± 0.44	41.4 ± 0.2	4.38 ± 0.22	1.49 ± 0.092	43.5 ± 3.82	
		150		8.83 ± 0.33	39.7 ± 1.3	4.49 ± 0.02	1.10 ± 0.039	39.1 ± 4.17	
		300		8.80 ± 0.21	43.2 ± 0.7	4.89 ± 0.19	0.95 ± 0.033	32.3 ± 0.66	
		75		9.39 ± 0.25	39.1 ± 1.4	4.17 ± 0.03	0.50 ± 0.014	27.0 ± 4.78	
		150		10.0 ± 0.10	45.1 ± 1.3	4.50 ± 0.18	1.39 ± 0.164	45.3 ± 1.03	
		300		10.7 ± 0.01	50.0 ± 0.4	4.68 ± 0.04	1.24 ± 0.108	47.3 ± 1.37	
		600		10.4 ± 0.98	45.2 ± 0.9	4.36 ± 0.34	0.72 ± 0.036	38.5 ± 0.17	

T. punctigera (Fig. 4B) as photoperiod decreased to 4:20 L:D achieved RUBISCO performance increased. To our surprise, the achieved RUBISCO turnover rate for *T. punctigera* under 4:20 L:D cycle was 8 times higher than under 24:0 L:D cycle, with little effect of growth light intensity (Fig. 4B). The content of intact RbcL protein varied across growth conditions with photoperiod generally showing more effect than instantaneous growth light intensity, particularly for the smaller *T. pseudonana* (Fig. 4, C and D). *T. punctigera* also showed a significant accumulation of RbcL fragments and larger products consistent with reactive oxygen stress (Wilson et al. 1995, Gerhardt et al. 1999, Nakano et al. 2010; Fig. S3 in the Supporting Information), particularly under the 24 h photoperiod.

To integrate these complex patterns, we compared retained carbon assimilation with reductant generation. To do so, we needed a common basis of normalization. Our estimates of protein $\cdot \text{cell}^{-1}$ did not vary significantly through the diel cycle, and allowed comparison of patterns between the small *T. pseudonana* and the large *T. punctigera*. In this study and in related papers (Wu et al. 2014b, Li et al. 2015), protein content proved our most reliable and general basis for comparison of different data types across the samples and species.

Normalization to chlorophyll is common in the oceanographic literature, but chlorophyll is a major component of PSII and its associated pigment protein complexes, while chlorophyll is not present within RUBISCO. So normalization of carbon assimilation and electron generation to total protein avoided co-variance of chlorophyll and PSII content. We therefore plotted the retained carbon assimilation rate per total cellular protein versus electron transport rate per total protein (Fig. 5, A and B) to show the instantaneous coupling of reductant generation at PSII to carbon assimilation (Halsey et al. 2010, 2013, Lawrenz et al. 2013), across the range of growth light intensities and photoperiods. In Figures 5, A and B the fine dotted line shows the theoretical maximum of one carbon to four electrons ($1\text{C}: 4 e^-$), assuming no diversion of electrons to other metabolic pathways and a $4 e^-$ reduction of CO_2 to (CH_2O) , with no accumulation of more reduced (CH_2) compounds.

Thalassiosira pseudonana (Fig. 5A) showed a saturating response of carbon assimilation rate to photochemical electron generation rate, with an initial slope of ~ 0.1 at low electron generation rates. The carbon assimilation rate per cellular total protein for *T. pseudonana* reached a maximum of $3.7 \text{ pmol C} \cdot \mu\text{g protein}^{-1} \cdot \text{s}^{-1}$ when the electron transport rate saturated at $37 \text{ pmol } e^- \cdot \mu\text{g}$

FIG. 3. Photosystem II electron transport rate ($e^- \cdot \text{PSII}^{-1} \cdot \text{s}^{-1}$) under illumination (A, B) and $\text{PSII}_{\text{active}}$ content ($\text{fmol} \cdot \mu\text{g protein}^{-1}$) (C, D) versus growth light ($\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$) for *T. pseudonana* (A, C) and *T. punctigera* (B, D) at L:D cycles of 24:0 (filled circles), 16:8 (open circles), 8:16 (open triangles), and 4:20 (filled triangles). Points show averages from each of 2 or 3 independently grown cultures; error bars show range of 2 replicated points and standard deviation of 3 replicated points, often within symbols. Each replicate was in turn each based upon an average of 5–9 repeated determinations taken within a diel cycle from a culture.

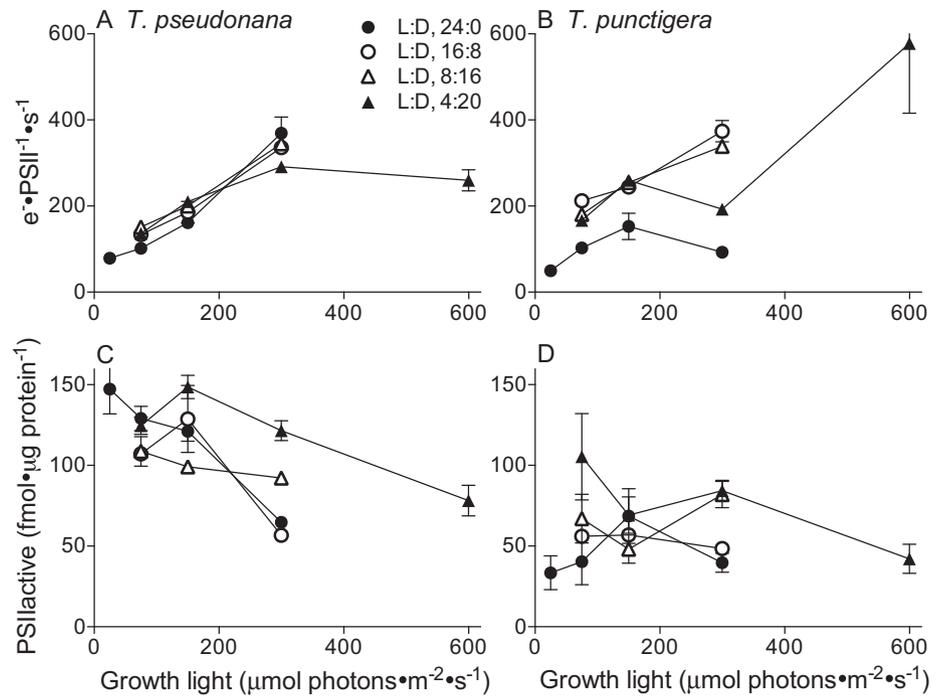
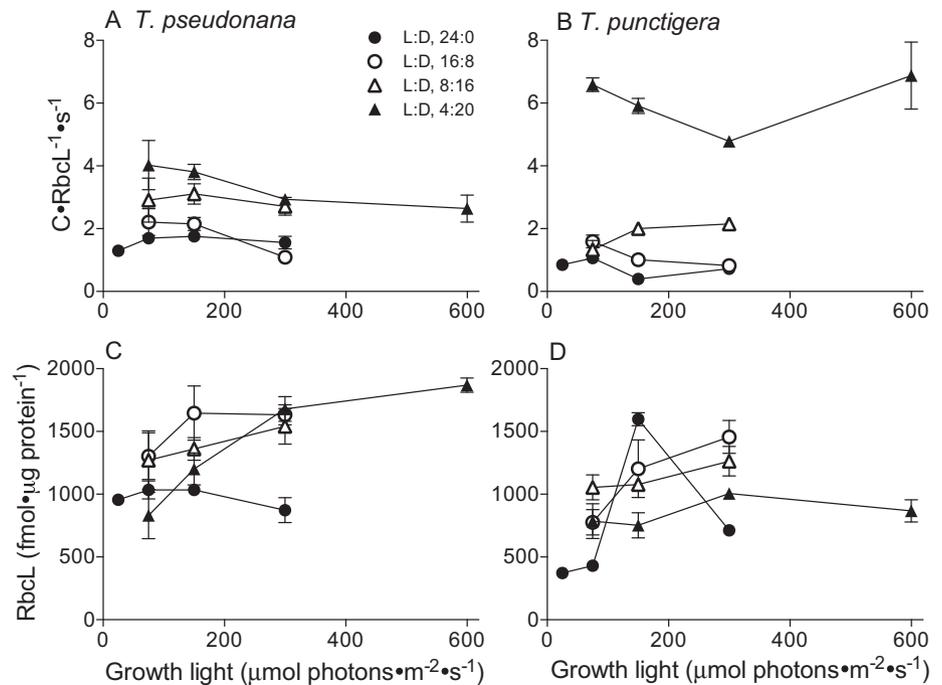


FIG. 4. Apparent carbon assimilation per RUBISCO active site ($\text{C} \cdot \text{RbcL}^{-1} \cdot \text{s}^{-1}$) under illumination (A, B) and RbcL content ($\text{fmol} \cdot \mu\text{g protein}^{-1}$) (C, D) versus growth light ($\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$) for *T. pseudonana* (A, C) and *T. punctigera* (B, D) at L:D cycles of 24:0 (filled circles), 16:8 (open circles), 8:16 (open triangles), and 4:20 (filled triangles). Points show averages from each of 2 or 3 independently grown cultures; error bars show range of 2 replicated points and standard deviation of 3 replicated points, often within symbols. Each replicate was in turn each based upon an average of 5–9 repeated determinations taken within a diel cycle from a culture.



$\text{protein}^{-1} \cdot \text{s}^{-1}$, derived from a fitted P-E curve (Peeters and Eilers 1978, Eilers and Peeters 1988, Macedo et al. 1998). *Thalassiosira pseudonana* cultures growing under 24:0 L:D and high instantaneous light intensity (closed circles, enclosed in oval) fell below the 95% confidence interval for the trend line fit to the rest of the data set and so were excluded from this curve fit. For 8:16,

16:8, and 24:0 L:D photoperiods *T. punctigera* showed a near linear response of carbon fixation to increasing photochemical electron generation ($r^2 = 0.73$; $F_{18} = 47.63$; slope significantly different from 0, $P < 0.0001$), with no indication of saturation (Fig. 5B). The slope of the regression was, however, only $0.07 \pm 0.01 \text{ C} \cdot \text{e}^{-1}$, far below the 0.25 slope of the 1 C : 4 e^{-} line. Therefore in

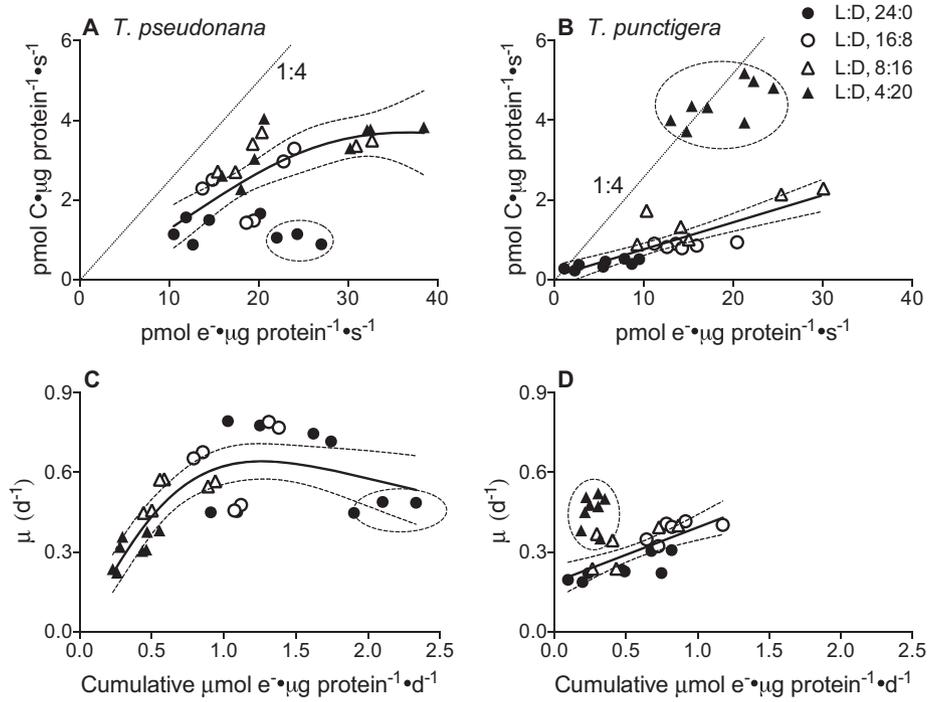


FIG. 5. Carbon fixation rate per cellular total protein ($\text{pmol C} \cdot \mu\text{g protein}^{-1} \cdot \text{s}^{-1}$) as a function of electron transport rate per total protein ($\text{pmol e}^- \cdot \mu\text{g protein}^{-1} \cdot \text{s}^{-1}$) for *T. pseudonana* (A) or *T. punctigera* (B) under L:D cycles of 24:0 (filled circles), 16:8 (open circles), 8:16 (open triangles), and 4:20 (filled triangles). The fine dotted line shows the theoretical maximum of 1C: 4 e⁻, assuming the 4 e⁻ reduction in CO₂ to (CH₂O) and no diversion of electrons to other metabolic pathways. Cell specific exponential growth rate (μ , d^{-1}) as a function of daily cumulative electron transport rate per total protein ($\mu\text{mol e}^- \cdot \mu\text{g protein}^{-1} \cdot \text{d}^{-1}$) for *T. pseudonana* (C) or *T. punctigera* (D). Solid lines: fitted curve with the growth-irradiance equation (A, C) or linear regression (B, D); thin dashed lines show 95% confidence intervals on the fitted curves. Oval in A outlines *T. pseudonana* measures from 24:0 L:D cycle at 300 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{d}^{-1}$ excluded from the curve fit. Measurements from the same samples in the oval in C were included in the curve fit. Ovals in B and D outline *T. punctigera* under 4:20 L:D cycles, excluded from the linear regression.

T. punctigera under most growth light intensities and photoperiods, the coupling of photochemical reductant generation to retained carbon assimilation was significantly lower than in *T. pseudonana* based upon the confidence intervals of the fitted data. *Thalassiosira punctigera* growing under 4:20 L:D was a notable exception. Under this short photoperiod, *T. punctigera* achieved a coupling of photochemical generation of reductant to retained carbon at or near the theoretical maximum of 1 C: 4 e⁻ (Fig. 5B), and well above the upper 95% CI for the fitted regression for the other photoperiods. This unexpectedly strong coupling of electron transport to retained carbon fixation then underpins the strong growth performance of *T. punctigera* under the shortest photoperiod.

To extend these measures of instantaneous photochemical performance, we plotted cell specific growth rate versus the cumulative diel photochemically generated reductant, normalized to total protein content. In *T. pseudonana*, μ shows a smoothly saturating response to increasing cumulative photochemical electrons per day across all combinations of growth light and photoperiod, with growth saturation of 0.64 d^{-1} determined from the curve fit,

achieved at $\sim 1.26 \mu\text{mol e}^- \cdot \mu\text{g protein}^{-1} \cdot \text{d}^{-1}$ (Fig. 5C). This plot of μ versus cumulative electron generation largely reconciles the complex interacting effects of growth light and photoperiod upon growth (Fig. 1).

In *T. punctigera* under 8:16, 16:8, and 24:0 L:D photoperiods μ responds linearly to increasing cumulative photochemical electron generation ($r^2 = 0.53$; $F_{1,8} = 19.8$; slope significantly different from 0, $P < 0.0003$; Fig. 5D), but with a significantly lower growth rate yield of reductant than in *T. pseudonana* (Fig. 5C) based upon the confidence intervals of the fitted data. The 4:20 L:D condition is again a notable exception (dotted oval; Fig. 5D), where *T. punctigera* achieves a growth yield per reductant higher than *T. pseudonana* at comparable daily reductant generation (Fig. 5C). We can thus largely reconcile the complex growth responses of *T. pseudonana* and *T. punctigera* to light intensity and photoperiod as driven by cumulative daily photochemical reductant generation, with higher performance in *T. punctigera* under short day/long night conditions because of high coupling of generated reductant to retained C under that combination of conditions.

We summarized these responses by plotting the retained carbon: electron ratio ($C:e^-$) versus cumulative daily photons (Fig. 6, A and B). *Thalassiosira pseudonana* shows the highest $C:e^-$ at lower daily photon doses, with some scatter depending upon the photoperiod over which the photons were delivered. *Thalassiosira punctigera* also shows maximal $C:e^-$ at the lowest daily photon dose (Fig. 6B), with $C:e^-$ values at the 4:20 L:D growth condition remarkably close to the theoretical maximum of 0.25 $C:e^-$.

DISCUSSION

We found systematic differences in the growth rate responses of two coastal marine centric diatoms differing by ~ 4 orders of magnitude in cell biovolume, cultured under a matrix of light intensities and photoperiods. We selected these related species as representatives from a wider panel of marine centric diatoms, which have shown consistent patterns of cell-size dependencies in their responses to light, photoinactivation, ocean acidification, and growth rates (Finkel 2001, Finkel et al. 2004, Key et al. 2010, Wu et al. 2014a,b).

The two diatoms showed complex and contrasting growth and photophysiological responses to light and photoperiod, which were not readily reconciled by separately examining their PSII activity (Fig. 3) nor their RUBISCO turnover (Fig. 4). Their overall growth responses were, however, largely reconciled by comparing their growth rate to cumulative photochemical reductant generation, which we were able to track using a new approach to quantitate the pool of active PSII contributing to reductant generation (Oxborough et al. 2012, Silsbe et al. 2015), in combination with an established approach to estimate the photochemical generation of electrons per PSII (Suggett et al. 2009, Huot and Babin 2010). In *T. pseudonana* growth showed a saturating response to increasing cumulative photochemical electrons, because the growth return per electron decreases as photoinhibition (Key et al. 2010) and dissimilatory electron flows (Waring et al. 2010)

accelerate under high light (Fig. 5C). In *T. punctigera*, growth showed a linear response to increase cumulative photochemical electrons, albeit at a lower growth yield per electron than in *T. pseudonana* (Fig. 5D). Growth under 4:20 L:D in *T. punctigera* proved the exception to the linear trend, with the strain achieving much higher growth yields per electron under this short photoperiod.

The smaller diatom *T. pseudonana* grew slowly under the shortest photoperiod, and fastest under 24:0 L:D (Fig. 1A), consistent with a pattern established for coastal strains (Brand and Guillard 1981). *Thalassiosira pseudonana* thus shows no need for a dark period to achieve maximum growth rate. In contrast, the larger coastal diatom *T. punctigera* achieved maximum growth rate under 4:20 L:D, while longer photoperiods slowed growth (Fig. 1B), consistent with a pattern common in offshore phytoplankton species (Brand and Guillard 1981). As noted by (Brand and Guillard 1981) for temperate strains, there is no direct selective pressure for exploitation of continuous 24:0 L:D, so the capacity to achieve maximum growth under 24:0 L:D suggests the absence of a requirement for a dark period, or absence of requirement for a regular photoperiod, rather than selection for exploitation of 24:0 L:D per se. Brand and Guillard (1981) argued that coastal strains are selected to exploit instantaneous light, of whatever duration, to cope with fluctuating light and nutrients in coastal environments (MacIntyre et al. 2000, Litchman et al. 2009), leading to a pleiotropic result of a capacity for rapid growth under continuous light. Indeed, *T. pseudonana* exploits rapidly fluctuating light regimes in part through an up-regulation of its light harvesting capacity (Grouneva et al. 2016) to exploit low light periods, which is feasible because of the rapid regulation of counter-balancing excitation dissipation in diatoms (Lavaud 2007, Lavaud and Lepetit 2013) after shifts to higher light.

In contrast to *T. pseudonana*, we noted that *T. punctigera* accumulated a ~ 118 kDa product that reacts with anti-RbcL antibody, consistent with an

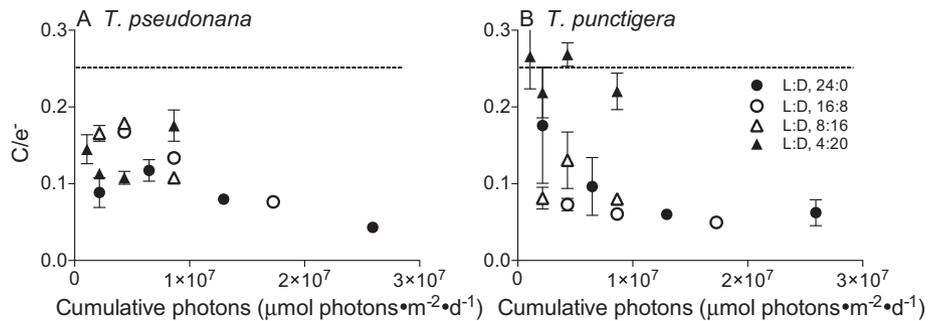


FIG. 6. Carbon fixation per PSII transported electron (C/e^-) versus daily cumulative photons ($\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{d}^{-1}$) for (A) *T. pseudonana* and (B) *T. punctigera* under L:D cycles of 24:0 (filled circles), 16:8 (open circles), 8:16 (open triangles), and 4:20 (filled triangles). Points show averages from 2 or 3 independently grown cultures; error bars show range of duplicated points and standard deviation of triplicated points, many bars fall within the size of the symbols. Horizontal dotted line shows 0.25 theoretical maximum 1C: 4 e^- .

RbcL-RbcL cross-link product, in parallel with accumulation of smaller fragments of 26 and 45 kDa that are immuno-reactive with the anti-RbcL antibody (Fig. S3, A and B). None of these products were apparent in the protein extracts from *T. pseudonana* from any growth condition. Proteins form di-tyrosine cross-links triggered by reactive oxygen species (ROS) (Berlett and Stadtman 1997, Larios et al. 2001, Yamamoto 2001) in a range of systems. The large subunit of RUBISCO, RbcL, indeed forms covalent cross-links with the smaller RbcS subunit under UV stress (Wilson et al. 1995, Gerhardt et al. 1999). Under ROS stress RbcL also fragments (Nakano et al. 2010) generating detectable products. In *T. punctigera*, the abundance of these RbcL products correlates with the 24 h continuous photoperiod growth condition, which strongly inhibited the growth rate of *T. punctigera*. We therefore hypothesize that *T. punctigera* has limited capacity to counter reactive oxygen toxicity during illumination and uses a daily dark period to allow detoxification or repair processes to catch up with products of light-driven reactive oxygen stress.

At least two mechanisms could cause cell size dependent changes in growth rate across different photoperiods. First, large cells with high capacity for variation in C:N ratio could be better adapted to long periods in the dark since they can draw upon a large energy reserve to fuel respiratory costs of nutrient assimilation and biosynthesis in the dark (Talmy et al. 2014). The diatom *Skeletonema costatum* growing under high light indeed showed large diurnal changes in C:N ratio (Anning et al. 2000, Ross and Geider 2009), consistent with respiratory consumption of organic C reserves over dark periods. Synchronized cultures of the green algae *Dunaliella bioculata* (Halsey et al. 2013, Halsey and Jones 2015) show diel fluctuations in allocations of photosynthetic production to proteins or polysaccharides, generating detectable changes in C:N through the day in synchrony with progression through the cell cycle. Over our range of photoperiods and low to moderately high growth light intensities, we observed only limited changes in C:N ratio over the diurnal cycle (Fig. S2), although our experimental design only tracked C:N through a single diel cycle, with limited temporal resolution so we lacked signal averaging to detect small or transient changes. Furthermore, we did not systematically establish synchronized cell cycles (Huysman et al. 2014), so cell-cycle dependent changes in C:N could be dampened in our cultures. The only combination of treatments where we detected a clear diel variation in C:N was the 4:20 L:D, 600 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ treatment of the smaller *T. pseudonana* (Fig. S2). Thus, for the diatoms and moderate growth light conditions considered here, diel variation in carbon reserves was not sufficient to cause large diel changes in C:N ratio, and so do not appear to explain the cell size related differences in growth rate responses.

Alternately, cell size dependent changes in growth rate under different photoperiods could result from differential coupling of photosynthetically generated reductant to retained carbon assimilation (Suggett et al. 2009, Halsey et al. 2010, 2011, 2013, Lawrenz et al. 2013). In *T. pseudonana* and *T. punctigera* assimilated C:e⁻ generally declined under increasing cumulative photon doses (Fig. 6). In contrast, short photoperiod allowed *T. punctigera* to convert reductant to retained carbon with remarkable efficiency (Fig. 6) across a range of daily cumulative photon doses. We did not track levels of excreted organic carbon in our experiments, which could vary across conditions or species. Therefore, the high ratio of retained C:e⁻ in *T. punctigera* under short photoperiods could result from a down-regulation of excretion of organic carbon under those particular conditions.

Tight coupling of photochemical reductant to retained carbon (Fig. 5B), along with the limited detected variation in C:N (Figs. S1 and S2) showed that the light-dependent metabolic activity needed for *T. punctigera* to maintain rapid growth can take place in as little as 4 h, and indeed that longer photoperiods (or shorter dark periods) were inhibitory to growth of *T. punctigera*. We used a maximum light intensity of 600 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$, but the full ability of large diatoms to utilize short photoperiods may only be observed at yet higher light intensities, e.g., the 1,180 $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ used by (Anning et al. 2000). This capacity for rapid burst exploitation of transient light periods likely relates to the tight metabolic coupling between diatom chloroplasts and mitochondria (Prihoda et al. 2012, Bailleul et al. 2015). On the other hand, the inhibition of growth of *T. punctigera* under longer photoperiods was surprising to us. Intrinsic susceptibility to photoinactivation of PSII decreases with increasing cell size across centric diatoms (Key et al. 2010), so we initially hypothesized that *T. punctigera* would show good resilience to prolonged photoperiods, since slower photoinactivation theoretically lowers the metabolic running costs needed to maintain the pool of active PSII (Wu et al. 2011, 2012, Campbell et al. 2013). We aim to pursue this finding in future work by analyzing the actual costs to sustain repair of PSII in smaller versus larger diatoms, and by confirming whether high generation or low detoxification of ROS leads to growth inhibition of *T. punctigera* under prolonged illumination.

To conclude, we found large differences in the growth and photosynthetic responses of two related diatoms of different cell sizes to a matrix of photoperiods and light intensities. The range of growth responses could be largely reconciled in terms of a saturable response of growth to photochemical reductant generation per diel cycle (Lawrenz et al. 2013). Short photoperiods specifically favor the larger diatom, which achieved a high coupling of photochemical reductant to retained C assimilation under short photoperiods. In addition to length of

photoperiod diatoms respond strongly to the pattern of light oscillation (Orefice et al. 2016) with sinusoidal oscillations generating responses distinct from the square-wave oscillations we imposed. We therefore sound a note of caution on meta-analytical extrapolations of photophysiological responses using data from studies applying different photoperiods or oscillation patterns to different taxa (Edwards et al. 2015), since different taxa show differential, even opposing, interactions between photoperiod and the light intensity responses of growth and photosynthesis. Pre-adapted capacities to exploit differing photoperiods (Brand and Guillard 1981) may prove important factors in the growth and biogeography of diatoms, even between two temperate coastal strains of the same genus.

Jessica Grant-Burt did cell count determinations for *T. punctigera*. Miranda Corkum and Dr. Ina Benner assisted with the CN analyses. Amy Woroch and Dr. Amanda Cockshutt provided the quantitations and analyses of RbcL products for Figure S3. This work was supported by a Canada Research Chair (DC) using equipment supported by the Canada Foundation for Innovation, New Brunswick Innovation Foundation and the Natural Sciences and Engineering Research Council of Canada. GL was supported by National Natural Science Foundation of China (41206132) and Natural Science Foundation of Guangdong Province, China (2015A030313826). DT was supported by NSF grant NSF/OCE/153795 and by the Gordon and Betty Moore Foundation through grant GBMF3778 to M.J. Follows. The authors thank Dr. H. MacIntyre for stimulating discussions on these topics, and an anonymous reviewer for extensive and constructive comments.

AUTHOR CONTRIBUTIONS

G.L. did the culturing and experimental work, assembled the underlying database, generated the draft figures, wrote drafts of the Materials & Methods and Results and contributed to the Introduction and Discussion. D.T. contributed to planning the experiments on the influences of photoperiod on diatom physiology, contributed to discussions of the data interpretation and plotting of the data, wrote a draft of the Introduction and contributed to the Discussion. D.A.C. planned the experiments with G.L., contributed to ongoing discussions of experimental progress and data analyses with G.L. & D.T., assisted with generation of figures, wrote the draft of the Discussion and assembled the submitted version of the manuscript.

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Supporting Information

Additional Supporting Information may be found in the online version of this article at the publisher's web site:

Figure S1. Cell volume (μm^3) (A, B) and carbon content per cell ($\text{pg} \cdot \text{cell}^{-1}$) (C, D) versus

growth light ($\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$) for *T. pseudonana* (A, C) and *T. punctigera* (B, D) at L:D cycles of 24:0 (filled cycles), 16:8 (open cycles), 8:16 (open triangles), and 4:20 (filled triangles). Points show averages of data from 2 or 3 independently grown cultures, averaged over five to nine repeated determinations from each culture within a diel cycle; error bars show range of two replicated points or standard deviation of three replicated points, often within symbols.

Figure S2. Diel changes of mole C:N ratios of *T. pseudonana* (A, C, E, G) and *T. punctigera* (B, D, F, H) at L:D cycles of 24:0 (A, B), 16:8 (C, D), 8:16 (E, F) and 4:20 (G, H), under growth light of 25 (reversed dark triangles), 75 (gray triangles), 150 (gray squares), 300 (open circles) or 600 (open diamonds) $\mu\text{mol photons} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$. Points show averages of 2 or 3 determinations on independently grown cultures; error bars show range of 2 replicated points or standard deviation of 3 replicated points, often falling within symbol size.

Figure S3. RbcL products as an index of reactive oxygen species stress correlate with growth inhibition in *Thalassiosira punctigera*. (A) Representative Immunoblot of RbcL (52 kDa) in total protein extracted from *Thalassiosira pseudonana*. (B) Representative Immunoblot of RbcL (52 kDa) and RbcL products (26, 45, and 118 kDa) in total protein extracted from *T. punctigera*. The immunoblots were set to equivalent exposures. The anti-RbcL antibody used for the immunoblots binds to a single conserved peptide region in the protein, so one RbcL protein offers only one potential antibody binding site. An RbcL product therefore derives from a single RbcL (in the case of 26 or 45 kDa fragments) or from one or more RbcL cross-linked together or to other proteins (in the case of 118 kDa product). (C) The sum of *T. punctigera* RbcL Products normalized to the RbcL band (52 kDa), averaged from samples from each photoperiod; 4, 8, 16, and 24 h. A one-way ANOVA with Tukey multiple comparisons of means shows that *T. punctigera* cultures growing under 24 h photoperiod (c) accumulate significantly more RbcL products than do cultures growing under 4, 8 ($P = 0.003$) (a) or 16 h ($P = 0.000$) (b) photoperiods. Error bars show one standard deviation of replicated points.