

Marine Phytoplankton Stoichiometry Mediates Nonlinear Interactions Between Nutrient Supply, Temperature, and Atmospheric CO₂

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Abstract

Marine phytoplankton stoichiometry links nutrient supply to marine carbon export. Deviations of phytoplankton stoichiometry from Redfield proportions (106C:1P) could therefore have a significant impact on carbon cycling, and understanding which environmental factors drive these deviations may reveal new mechanisms regulating the carbon cycle. To explore the links between environmental conditions, stoichiometry, and carbon cycling, we compared four different models of phytoplankton C:P: a fixed Redfield model, a model with C:P given as a function of surface phosphorus concentration ([P]), a model with C:P given as a function of temperature, and a new multi-environmental model that predicts C:P as a function of light, temperature, and [P]. These stoichiometric models were embedded into a five ocean circulation box model, which resolves the three major ocean biomes (high-latitude, subtropical gyres, and tropical upwelling regions). Contrary to the expectation of a monotonic relationship between surface nutrient drawdown and carbon export, we found that lateral nutrient transport from lower C:P tropical waters to high C:P subtropical waters could cause carbon export to decrease with increased tropical nutrient utilization. It has been hypothesized that a positive feedback between temperature and $p\text{CO}_{2,\text{atm}}$ will play an important role in anthropogenic climate change, with changes in the biological pump playing at most a secondary role. Here we show that environmentally driven shifts in stoichiometry make the biological pump more influential, and may reverse the expected positive relationship between temperature and $p\text{CO}_{2,\text{atm}}$. In the temperature-only model, changes in tropical temperature have more impact on the $\Delta p\text{CO}_{2,\text{atm}}$ (~ 41 ppm) compared to subtropical temperature (~ 4.5 ppm). Our multi-environmental model predicted a decline in $p\text{CO}_{2,\text{atm}}$ of ~ 46 ppm when temperature spanned a change of 10°C . Thus, we find that variation in marine phytoplankton stoichiometry and its environmental controlling factor can lead to non-linear controls on $p\text{CO}_{2,\text{atm}}$, suggesting the need for further studies of ocean C:P and the impact on ocean carbon cycling.

1 Introduction

50 The discovery of large-scale deviations of phytoplankton stoichiometry from the Redfield ratio in the past decade (Martiny et al., 2013a, 2013b; Weber and Deutsch, 2010) has significant consequences for our understanding of the biological carbon pump and global carbon cycling (Galbraith and Martiny, 2015; Moreno and Martiny, 2018). Traditionally, the biological pump is thought to be controlled by a combination of the vertical nutrient flux and
55 nutrient utilization efficiency (also known as elemental stoichiometry) (Sarmiento and Toggweiler, 1984). Evidence that elemental stoichiometry is variable thus adds a new dimension to the biological pump, and may lead to higher than currently expected carbon export ($C:P_{export}$) in subtropical regions (Emerson, 2014; Tanioka and Matsumoto, 2017; Teng et al., 2014). Global carbon export has been estimated to range between 5 and 12 Pg
60 C/year (Boyd and Trull, 2007; Henson et al., 2011), but these projections have yet to incorporate the environmental controls on $C:P_{export}$. Variation in $C:P_{export}$ from Redfield proportions can be linked to environmental conditions. There are two leading environmental parameters thought to control $C:P_{export}$; nutrients, predominantly phosphate concentrations, and temperature. Galbraith and Martiny used a simple three-box model to
65 show that variable stoichiometry driven by phosphate availability could enhance the efficiency of the biological pump in the low-latitude ocean (Galbraith and Martiny, 2015). In contrast, Yvon-Durocher and co-workers (2015) used a meta-analysis of global temperature and stoichiometric ratios to propose that C:P increased 2.6-fold from 0° C to 30° C. Thus, it is unclear if differences in nutrient supply, temperature, or some
70 combination of them, control the global variation in C:P of plankton and exported material.

There are two important ingredients missing from published studies that could alter the interactions among phytoplankton stoichiometry, carbon export, and atmospheric pCO_2 ($pCO_{2,atm}$). The first is the presence of two distinct low-latitude biomes, namely the equatorial upwelling regions and the macronutrient-depleted subtropical gyres. In direct
75 observations and inverse model analyses, these two biome types appear to have unique elemental compositions, which leads to relatively increased rates of export from oligotrophic gyres in comparison to equatorial upwelling regions (DeVries and Deutsch, 2014; Martiny et al., 2013a; Teng et al., 2014). Thus, in order to properly represent global variations in surface plankton C:P and carbon export, it is essential to separately model
80 both macronutrient-limited subtropical gyres and equatorial upwelling zones.

The second missing ingredient is that environmental factors beyond nutrient availability may impact the elemental composition of surface plankton and $C:P_{export}$. Temperature, irradiance, and nutrient concentrations are all important environmental factors, which influence the physiology and stoichiometry of phytoplankton. However,
85 surveys of phytoplankton C:P are insufficient to distinguish between the separate effects of each factor on C:P due to strong environmental covariance. Cellular trait-based models use detailed studies of phytoplankton physiology to determine how phytoplankton cells should allocate their resources as a function of environmental conditions, allowing us to model the interactive influence of temperature, nutrient concentrations, and irradiance on C:P ratios
90 (Clark et al., 2011; Daines et al., 2014; Shuter, 1979; Talmy et al., 2014; Toseland et al., 2013). Numerous physiological mechanisms have been proposed to explain variation in phytoplankton stoichiometry, including growth rate (Sterner and Elser, 2002), photoacclimation (Falkowski and LaRoche, 1991; Geider et al., 1996; Leonardos and Geider, 2004, 2005), nutrient-limitation responses (Garcia et al., 2016; Goldman et al.,

951979; Rhee, 1978), and temperature acclimation (Rhee and Gotham, 1981; Toseland et al., 2013; Yvon-Durocher et al., 2015). Through incorporation of such physiological responses, a trait-based model has revealed that differences in ribosomal content and cell size between warm-water, oligotrophic environments and cold-water, eutrophic environments are important mechanisms driving stoichiometric variation in the ocean (Daines et al., 1002014). Thus, linking biome-scale variations in environmental conditions with a detailed trait-based model of phytoplankton resource allocation and elemental composition may enable us to more fully explore interactions among multiple ocean environmental factors, the biological pump, and $p\text{CO}_{2,\text{atm}}$.

Here we create a five ocean circulation box model, incorporating the three major 105ocean biomes, to study the feedback effects of variable stoichiometry on carbon export and $p\text{CO}_{2,\text{atm}}$. We will explicitly address the following research questions: (1) How does environmental variability influence marine phytoplankton cellular allocation strategies and in turn the elemental stoichiometric ratio? (2) What are the effects of changing environmental conditions on stoichiometric ratios, carbon export, and $p\text{CO}_{2,\text{atm}}$?, and (3) 110What is the influence of the environmental conditions among the three major surface biomes on carbon export and $p\text{CO}_{2,\text{atm}}$?

2 Methods

2.1 Stoichiometric Models

115To quantify and understand the feedbacks between carbon export and $p\text{CO}_{2,\text{atm}}$, we embedded four stoichiometric models into our five ocean circulation box model. Each model differs according to its complexity and how much environmental information they utilize. These are a static Redfield model that assumes that $\text{C:P}_{\text{export}}$ is constant across environmental conditions, a nutrient-only model that uses surface $[\text{P}]$ to predict $\text{C:P}_{\text{export}}$ 120(from Galbraith and Martiny, 2015), a temperature-only model that uses T to predict $\text{C:P}_{\text{export}}$ (modified from Yvon-Durocher et al., 2015), and a multi-environmental model that uses light (I), T , and $[\text{P}]$ to predict $\text{C:P}_{\text{export}}$.

2.1.1 Static Redfield Model

125Our control model uses a static Redfield stoichiometry. The Redfield ratio is based on an average value of organic carbon to phosphorus of 106:1.

2.1.2 Nutrient-Only Model

The nutrient-only stoichiometric model expresses phytoplankton C:P as a function of the 130ambient phosphate concentration:

$$C:P = \frac{1}{\kappa[\text{P}] + [\text{P}]_0} \quad (1)$$

where the parameters $\kappa = 6.9 \times 10^{-3} \mu\text{M}^{-1}$ and $[\text{P}]_0 = 6.0 \times 10^{-3}$ were obtained by regressing the reciprocal of C:P onto $[\text{P}]$ (Galbraith and Martiny, 2015).

2.1.3 Temperature-Only Model

135The temperature-only stoichiometric model expresses phytoplankton C:P as a function of temperature:

$$\ln(C:P) = \Pi(T - 15^{\circ}\text{C}) + b, \quad (2)$$

where the parameters $\Pi = 0.037/^{\circ}\text{C}$ and $b = 5.5938$ (Yvon-Durocher et al., 2015). The temperature-only model was created to determine the temperature responses of log-transformed C:P ratios centered at 15°C .

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2.1.4 Multi-Environmental Model

We created a multi-environmental model which predicts how cell size, biomass allocations to biosynthesis and photosynthesis, and C:P ratios vary with light levels, temperature, and phosphorus concentrations. The multi-environmental factor model was derived from a non-dynamic physiological trait-based model. We used a theoretical cellular-allocation trait model based on phytoplankton physiological properties that divides the ‘cell’ into several functional pools which represent cellular investments in biosynthesis, photosynthesis, and structure, and a storage pool, which represents variations in the level of P-rich molecules such as polyphosphates. The functional pools are composed of biological macromolecules such as ribosomes, proteins, carbohydrates and lipids. The model predicts the size of each pool as a function of light, temperature, and [P]. The size of each functional pool is modeled by using subcellular resource compartments, which connect the fitness of a hypothetical phytoplankton cell in a given environment to its cellular radius and the relative allocation of cellular material to photosynthetic proteins, ribosomes, and biosynthetic proteins. We assume that real phytoplankton populations have physiological behaviors that cluster around the strategy that produces the fastest growth rate in each environment (Norberg et al., 2001), and use the stoichiometry of this optimal strategy to model the elemental composition of cellular material (Figure 1).

Phytoplankton can accumulate large reserves of nutrients that are not immediately incorporated into the functional components of the cell (Diaz et al., 2016; Mino et al., 1998; Van Mooy and Devol, 2008; Mouginot et al., 2015). This storage capability varies among phytoplankton species, and depends on the particular nutrient under consideration: the cost for storing physiologically relevant quantities of nutrients is low for nutrients with low quotas such as phosphorus, in comparison to nitrogen and carbon. Thus, the phosphorus storage is assumed highly plastic in comparison to carbon storage (Moore et al., 2013).

Further, we assume that each cell dedicates a fixed fraction of its biomass to carbon reserves, and focus our modeling efforts on the variability of the stored phosphorus pool. To predict the size of the storage pool, we assume a linear relationship between stored phosphorus and ambient environmental phosphorus levels and used statistical modeling of an oceanic C:P dataset (Martiny et al., 2014) to calculate the constant of proportionality. The result is a relatively simple model that both qualitatively and quantitatively predicts the variation of C:P in plankton throughout the oceans.

Phytoplankton physiology is modeled through allocations of cell dry mass to three distinct pools: structure ($S(r)$), biosynthesis (E), and photosynthesis (L). Allocations satisfy:

$$1 = S(r) + E + L, \quad (3)$$

where the variables S , E , and L represent the *specific* allocations of cellular biomass.

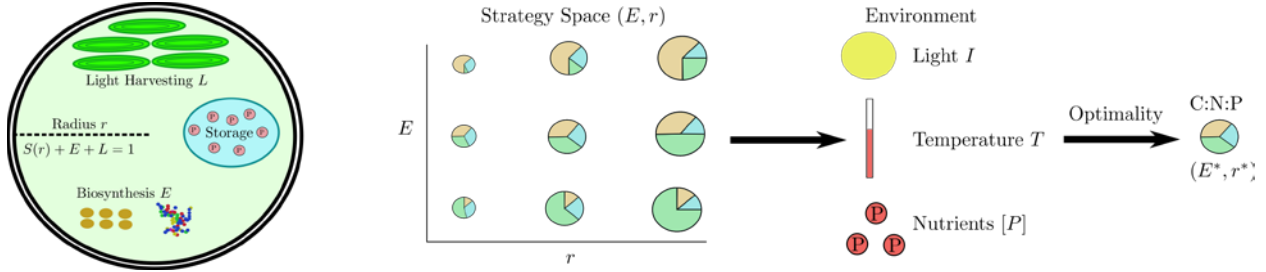


Figure 1: Diagram of physiological model. Phytoplankton strategies are represented in a two-dimensional strategy space (E, r) . Each strategy is assigned a fitness in each environment using physiological principles, and the strategy with the highest fitness is selected to represent the local population. The stoichiometry of cellular components is used to calculate the stoichiometry of the functional pools in the cell.

The specific allocation of biomass to the cell membrane is inversely proportional to the cell radius $\left(\frac{\alpha}{r}\right)$ (Clark et al., 2011), which accounts for the changing relative volume of the cell-membrane with radius. The structure pool includes the cell membrane plus wall and other components (γ), which are not related to photosynthesis or biosynthesis and is given by:

$$S(r) = \frac{\alpha}{r} + \gamma. \quad (4)$$

In an environment specified by I , T , and $[P]$, the growth rate of a cell using a given strategy is the minimum of the following growth rates:

$$\mu = \min(\mu_E, \mu_L, \mu_P). \quad (5)$$

Here μ_E is determined by the specific rate of protein synthesis, μ_L is determined by the specific rate of carbon fixation, and μ_P is determined by the specific rate of phosphorus uptake, or:

$$\mu_E = k_E(T)E, \mu_L = \frac{f_P(L, I) - \Phi_M(T)}{1 + \Phi_S}, \mu_P = \frac{1}{Q_P(r, E)} \frac{V_m(r)[P]}{K_P(r) + [P]}. \quad (6)$$

We assume that part of the energy captured by a cell via photosynthesis is used for maintenance (Φ_M), whereas the rest is used to drive the synthesis of new macromolecules (Φ_S), so that a cell growing at rate μ_L is in energy balance. The efficiency of biosynthesis k_E and the carbon cost of maintenance Φ_M are functions of T , whose dependence is modeled using $Q_{10} = 2.0$ (Van Bogelen and Neidhardt, 1990; Broeze et al., 1978; Shuter, 1979). Uptake is regulated by a Monod function with kinetic parameters depending on the radius through the allometric scaling relationships derived from measurements of phytoplankton populations (Edwards et al., 2012):

$$V_m(r) = a_P r^{b_P}, K_P(r) = a_K r^{b_K}. \quad (7)$$

This use of allometric scaling relationships departs from the conventions adopted by Shuter (1979) (Shuter, 1979) or Daines et al. (2014) (Daines et al., 2014), who assume that uptake rates are diffusion-limited.

The phosphorus quota for functional elements of the cell (thus not including any storage) is determined by the allocation to biosynthesis E and the percentage p_{DNA} of cellular dry mass allocated to DNA:

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$$Q_{p,biosynthesis}(E, r) = \frac{4}{3} \pi r^3 \rho_{cell} p_{dry} \frac{(\alpha_E E P_{rib} + p_{DNA} P_{DNA})}{31}. \quad (8)$$

Here we assume that there is no contribution to the functional-apparatus P quota from phospholipids, which instead are merged with storage molecules. This differs from Daines et al. (2014), who assume that phospholipids occupy 10% of the cell by mass.

Phytoplankton can substitute sulfoquinovosdiacylglycerol (SQDG) for phospholipids in their cell membranes under low P conditions (Van Mooy et al., 2009). Similarly, P storage molecules are also regulated by P availability. Thus, we here assume that phospholipids and P-storage exhibit the same behavior and thus model-wise treated as one pool (Van Mooy et al., 2009).

The function f_p is the cellular response to light levels, and is chosen to capture the effects of both electron transport and carbon fixation on photosynthesis, and is closely related to a previous model (Talmy et al., 2013)(Talmy et al., 2014). This prior model included four compartments: electron transport, carbon fixation, photoprotection, and biosynthesis. The outcome was that photoprotection allocation was not a large or greatly changing component of their allocations. We therefore do not include this within our model due to its high complexity with little qualitative results. Our biosynthesis was also separately parametrized.

The decomposition of photosynthesis into light harvesting and carbon fixation components is critical, and makes our model predictions agree much better with experiments studying the variations of C:P or N:P ratios with irradiance. Models that do not have this decomposition predict too large a decrease in cellular allocations to photosynthesis at high-light levels. In a two-compartment model, increases in allocations to carbon fixation cause the overall allocation to light harvesting to have a more mild decrease. The two-compartment treatment also seems more physiologically realistic than a 1-compartment treatment, which only models photosynthetic pigments. Thus, we used the functional forms and parameters that were derived (experimentally) previously for carbon-fixation and light harvesting (Talmy et al., 2013).

Our model interprets the light harvesting allocation, L , as being composed of proteins dedicated to carbon fixation (F_1), such as RuBisCO, and proteins dedicated to light harvesting (F_2), such as photosynthetic pigments. The rate of photosynthetic carbon fixation is a function of the allocations to each of these, which satisfy $F_1 + F_2 = L$. The relative allocations together determine the overall photosynthetic rate:

$$P_{max} = \min(k_1 F_1, k_2 F_2), f_p = P_{max} \left(1 - \exp \left(\frac{-\alpha_{ph} \phi_M F_2 I}{P_{max}} \right) \right). \quad (9)$$

For a given I and L , there is a pair of values $(F_{1,opt}, F_{2,opt})$ that maximize the photosynthetic rate f_p . We estimate the photosynthetic rate $f_p(L, I)$ under the assumption that cells assume the optimal allocations to carbon fixation and electron transport. This model

departs from the models developed by Shuter (1979) and Daines et al. (2014), which assume that energy acquisition is a linear function of light levels, with functional response linearly proportional to the cellular investment in light harvesting proteins.

We model photosynthesis as having a $Q_{10}=1$, which is consistent with physiological studies going back to Shuter (1979) that suggest that photosynthetic efficiency does not depend on temperature over physiologically relevant ranges. The discrepancy between photosynthetic and biosynthetic temperature dependence has traditionally been explained by referring to the differences in the chemistry and physics of the two processes. The electron transport chain relies on quantum mechanical processes, which are unaffected by variations in temperature in a physiologically relevant range (Devault, 1980). Required maintenance respiration rates are also modeled as having a $Q_{10}=2.0$. We model the phytoplankton community residing in a given environment by assuming it consists solely of the phytoplankton type using the highest growth rate strategy in that environment. This strategy is found by solving for the values of r and E that make

$$\mu = \mu_L = \mu_P = \mu_E \quad (10)$$

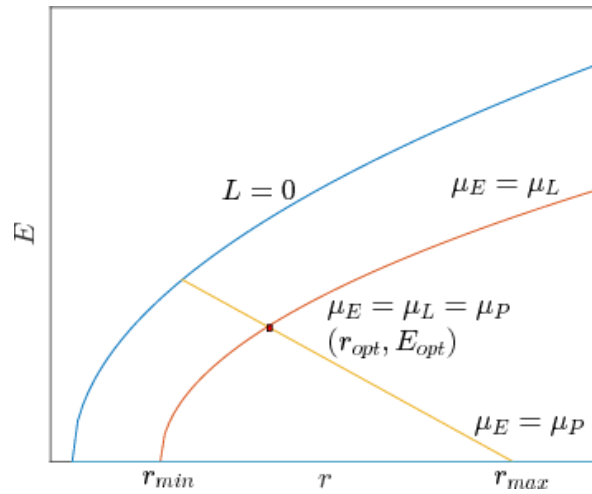


Figure 2: Diagram of strategy space. The (r, E) plane is divided into a region in the first quadrant where $L > 0$, corresponding to the set of allowable strategies. The optimal strategy occurs at the point (r_{opt}, E_{opt}) , denoted by the red rectangle, where $\mu = \mu_L = \mu_P = \mu_E$.

We will now show that under two assumptions that will be true in nearly any realistic situation, a strategy maximizing μ always exists, is unique, and satisfies $\mu = \mu_L = \mu_P = \mu_E$ (Figure 2). The function μ_L is a function of the chosen strategy (r, E) , and it is an increasing function of r and decreasing function of E . The first assumption is that light levels are sufficiently high that there exists some r_{min} such that $\mu_L(r_{min}, 0) > 0$, which means that light is sufficient for some phytoplankton to be able to overcome maintenance costs. The function μ_P is a monotonically decreasing function of both r and E . Because there is a non-zero amount of P contained in the structure pool, and because uptake rates decline to zero with r , there will be some r_{max} at which $\mu_P(r_{max}, 0) > 0$. The second assumption is that $r_{min} < r_{max}$, which will be true for most realistic values of the light level. We note that for fixed r , μ_E is a monotonically decreasing function of E . Since none of μ_E , μ_L or μ_P have

critical points, the function μ can only have a maximum at places where two or more of μ_L , μ_P , and μ_E are equal, or at the boundaries of the strategy space. On the boundaries of strategy space, $E = 0$ or $L = 0$ so that $\mu \leq 0$. We can exclude the boundary and focus on places where two or more of μ_L , μ_P , and μ_E are equal. We define two curves, one on which $\mu_L = \mu_E$, and the other on which $\mu_P = \mu_E$. The curve for which $\mu_L = \mu_E$ begins at the point $r = r_{min}$ and can be described by a monotonically increasing function $E = g(r)$ on the interval $[r_{min}, \infty]$. This curve exists because $\mu_E = 0$ when $E = 0$, $\mu_L > 0$ when $E = 0$ and $r_{min} < r$, and $\mu_L < 0$ when $L = 1 - S(r) - E = 0$, so that there is always a solution to $\mu_L = \mu_E$ for fixed $r > r_{min}$. To see that the curve is an increasing function of r , consider the function $V(E, r) = \mu_L - \mu_E$ and apply the chain rule to the equation $V(g(r), r) = 0$ to find that along the curve $E = g(r)$:

$$\frac{dE}{dr} = g'(r) = \frac{-\frac{\partial V}{\partial r}}{\frac{\partial V}{\partial E}} \quad (11)$$

We consider the terms in equation 11 carefully. The function V is an increasing function of r because μ_E is independent of r and because μ_L is an increasing function of r (for a fixed investment in biosynthesis, a larger radius leads to a greater investment in photosynthesis and greater photosynthetic growth rate). Thus, the numerator of equation 11 is negative. The function V is a decreasing function of E because μ_L is a decreasing function of E (greater investments in biosynthesis at fixed radius lead to smaller investments in photosynthesis) and μ_E is an increasing function of E . Thus the denominator of equation 11 is negative, and the quotient on the right hand side is positive, so $g'(r)$ is positive and describes an increasing curve.

By similar logic, we can define a curve $h(r)$ that solves the equation $\mu_P(h, r) = \mu_E(h, r)$. This curve exists on the finite interval $[r_l, r_{max}]$, where r_l solves the equation $\mu_P(1 - S(r_l), r_l) = \mu_E(1 - S(r_l), r_l)$. Thus $h(r)$ represents a decreasing curve from the point $(1 - S(r_l), r_l)$ to $(0, r_{max})$. We can see that $h(r)$ is always decreasing by using the chain rule on $\mu_P(h, r) - \mu_E(h, r) = 0$, just as in the previous argument.

The growth maximizing strategy must occur somewhere on the curves described by $(g(r), r)$ and $(h(r), r)$. The functions $\mu_1(r) = \mu(g(r), r)$ and $\mu_2(r) = \mu(h(r), r)$ are continuously differentiable functions of r except where $g(r) = h(r)$ (which must exist by the intermediate value theorem). Therefore the only place where μ can have a maximum is at the place where $g(r)$ and $h(r)$ intersect, which is the strategy that leads to equality of all the growth rates. We refer to this strategy, as a function of environmental conditions, as $(r_m(P, I, T), E_m(P, I, T), L_m(P, I, T))$. Using this strategy we can predict the stoichiometry of the functional components of the phytoplankton population in a given environment.

We assume that real phytoplankton populations cluster near the optimal strategy in the local environment (Norberg et al., 2001):

$$(E_m, r_m) = \operatorname{argmax}_{(E, r)} \mu. \quad (12)$$

For all values of environmental parameters used in this study, the unique maximum of the growth rate occurs for the set of parameter values that lead to co-limitation by nutrients,

photosynthesis, and biosynthesis, analogously to the predictions of Klausmeier and co-workers (2004). The optimal strategy determines the model prediction of the C:P of functional components in a given environment by taking the quotient of the carbon and phosphorus quotas.

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Table 1. *Physiological Model Constants.*

PARAMETER	DESCRIPTION	VALUE	UNITS	SOURCE
α	Proportionality coefficient for radius	0.12	-	(Toseland et al., 2013)
γ	Percent dry mass devoted to structure other than membrane	0.2	-	(Toseland et al., 2013)
k_{E0}	Synthesis rate of biosynthesis apparatus at $T_0=25$	0.168	hr^{-1}	(Shuter, 1979)
$Q_{10,E}$	Q_{10} of biosynthetic apparatus	2.0		(Shuter, 1979)
Φ_{M0}	Specific carbon cost of maintenance at $T_0=25$	10^{-3}	hr^{-1}	(Shuter, 1979)
$Q_{10,M}$	Q_{10} of maintenance	2.0	-	(Shuter, 1979)
$Q_{10,P}$	Q_{10} of photosynthesis	1.0		(Shuter, 1979)
Φ_S	Carbon cost of synthesis	0.67	-	(Shuter, 1979)
a_P	Allometric scaling constant for VMP	1.04×10^{-16}	$(\text{mol P})(\text{hr})^{-1}$	(Edwards et al., 2012)
b_P	Allometric scaling exponent for VMP	3.0	-	(Edwards et al., 2012)
a_K	Allometric scaling constant for KP	6.4×10^{-8}	$(\text{mol P})(\text{L})^{-1}$	(Edwards et al., 2012)
b_K	Allometric scaling exponent for KP	1.23	-	(Edwards et al., 2012)

ρ_{cell}	Cell Density	10^6	g/m^3	(Shuter, 1979)
ρ_{dry}	Fraction of dry mass in cell	0.47	-	(Toseland et al., 2013)
α_E	Fraction of dry mass in biosynthetic apparatus devoted to ribosomes	0.55	-	(Toseland et al., 2013)
ρ_{rib}	Fraction of ribosomal mass in phosphorus	0.047	-	(Sternier and Elser, 2002)
ρ_{DNA}	Fraction of cell dry mass in DNA	0.01	-	(Toseland et al., 2013)
ρ_{DNA}	Fraction of DNA mass in phosphorus	0.095	-	(Sternier and Elser, 2002)
k_1	Specific Efficiency of Carbon Fixation Apparatus	0.373	hr^{-1}	(Talmy et al., 2013)
k_2	Specific Efficiency of Electron Transport Apparatus	0.857	hr^{-1}	(Talmy et al., 2013)
α_{Ph}	Light Absorption	1.97	m^2/gC	(Morel and Bricaud, 1981)
ϕ_M	Maximum Quantum Efficiency	10^{-6}	$gC/\mu mol \text{ photons}$	(Falkowski and Raven, 1997)
m_{lip}	Fraction of cell membrane composed of lipids	0.3	-	(Toseland et al., 2013)
m_{prot}	Fraction of cell membrane composed of protein	0.7	-	(Toseland et al., 2013)
p_{lip}	Fraction of cell dry mass in storage lipids	0.1	-	(Sternier and Elser, 2002)
p_{carb}	Fraction of cell dry mass in storage	0.04	-	(Sternier and Elser, 2002)

	carbohydrates			
C_{DNA}	Fraction of DNA mass in Carbon	0.36	-	(Sternner and Elser, 2002)
C_{rib}	Fraction of ribosomal mass in Carbon	0.42	-	(Sternner and Elser, 2002)
C_{prot}	Fraction of protein mass in Carbon	0.53	-	(Sternner and Elser, 2002)
C_{lip}	Fraction of lipid mass in Carbon	0.76	-	(Sternner and Elser, 2002)
C_{carb}	Fraction of carbohydrate mass in Carbon	0.4	-	(Sternner and Elser, 2002)

The carbon quota is calculated as:

$$Q_C = \frac{\left(\frac{m_{lip}\alpha}{r} p_{lip} C_{lip} + p_{carb} C_{carb} + \alpha_E E C_{rib} + \left((1 - \alpha_E) E + L + \frac{m_{prot}\alpha}{r} \right) C_{prot} + p_{DNA} C_{DNA} \right)}{\frac{4}{3} \pi r^3 \rho_{cell} p_{dry}} \quad (13)$$

Here we see the contributions of carbon contained in both functional and storage pools, the latter of which are assumed to occupy a fixed fraction of the cell independent of the environment (but linked to cell size).

Measurements of cellular P partitioning indicate that the ribosomal RNA can sometimes contribute only 33% of the total P quota (Garcia et al., 2016). The additional phosphorus includes membrane phospholipids and storage compounds, each of which can be up- or down-regulated in response to phosphorus availability in the environment. To model this phenomenon, we assume the existence of an additional stored P pool, whose size is a linear function of environmental P, or:

$$(P:C)_{storage} = \epsilon[P], \quad (14)$$

where ϵ is determined by the best fit to the Martiny et al. (2014) data. Our model then predicts C:P as:

$$C:P = \frac{1}{(P:C)_{(E_m, r_m)} + \epsilon[P]}. \quad (15)$$

The model parameter ϵ is calculated by minimizing the residuals of the P:C ratio predicted by Eq.13 in comparison to the global data-set on particulate organic matter stoichiometry compiled by Martiny and others (2014). To maintain consistency with the linear regression model of Galbraith and Martiny (2015), we restrict the dataset to observations from the upper 30 meters of the water column containing particulate organic phosphorus and

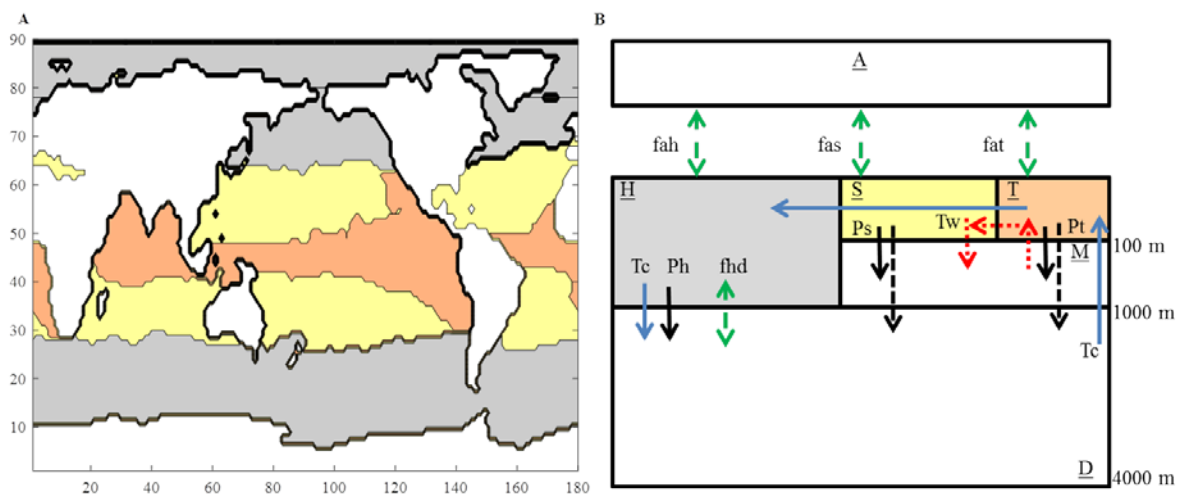
340 carbon concentrations of greater than 5 nM. Observations from the same station and the same day, but at different depths in the water column are averaged together. The P:C ratio of the functional apparatus is calculated using irradiance, temperature, and [P] data from the World Ocean Atlas (Garcia et al., 2014; Locarnini et al., 2013; oceancolor.gsfc.nasa.gov/data/10.5067/AQUA/MODIS/L3B/PAR/2014/), which are used
345 to estimate environmental conditions at the location and date of particulate organic matter measurements. Light levels are computed by averaging irradiance over the top 50 meters of the water column, assuming an e-folding depth of 20 meters. Linear regression determines $\epsilon = 2500 \text{ M}^{-1}$ which fits the data with an $R^2 = 0.28$. All parameters for the model are listed in Table 1.

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2.2 Box Model Design

To quantify the feedbacks between phytoplankton stoichiometry, carbon export, and $\text{pCO}_{2,\text{atm}}$, we formulated a five-box ocean circulation model of the phosphorus and carbon cycles in the ocean coupled to an atmospheric box. The foundation of our model is based on
355 the models introduced in Ito and Follows (2003) and DeVries and Primeau (2009).

Phosphorus is used to represent the role of nutrient availability in controlling stoichiometry and C export. We chose this over N to avoid having to include a parameter rich N cycle. Furthermore, P rather than N is commonly regarded as the ultimate limiting nutrient (Tyrrell, 1999) and thus P availability represents the long-term steady-state
360 biogeochemical equilibrium. The model includes three surface boxes, each corresponding to one of the major biomes: the tropical equatorial upwelling regions (labeled T), the subtropical gyres (labeled S), and the high-latitude regions (labeled H) (Figure 3). We define the oligotrophic subtropical gyre regions where the mean annual phosphate concentration is less than $0.3 \mu\text{M}$ (Teng et al., 2014), with the remainder of the surface
365 ocean assigned either to box T or box H based on latitude. We use these assignments to calculate the baseline physical properties of each region, including mean annual averaged irradiance and temperature. The subsurface ocean is divided into two regions: the thermocline waters that underlies the subtropical gyres and the equatorial upwelling regions (labeled M), and deep waters (labeled D) (DeVries and Primeau, 2009).



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Figure 3: Box Model Design. A) Sea surface breakdown by region. All peach-colored regions represent the tropical surface ocean box, the cream-colored regions represent the subtropical surface ocean box, and grey

regions represent the high-latitude surface ocean box. B) The model includes tropical (*T*), subtropical (*S*), and high-latitude (*H*) surface ocean boxes, a mixed thermocline (*M*) box, and a deep water (*D*) box. The thermohaline circulation *T_c* is set to 20 Sv, while the wind driven shallow overturning circulation is set to 5 Sv. The high-latitude mixing flux *f_{hd}* is set to 45.6 Sv. The thickness of Box *H* is 1000 m, and Box *M* is 900 m. Box *T* has a temperature of 26°C, box *S* has a temperature of 24°C, and box *H* has a temperature of 7°C. Box *S* covers 39% and Box *T* covers 25% of the ocean surface area.

To simulate the global transport of water between boxes, our model includes a thermohaline circulation (labeled *T_c*) that upwells water from the deep ocean into the tropics, laterally transports water into the subtropics and high-latitudes, and downwells water from the high-latitude region to the deep ocean. Surface winds produce a shallow overturning circulation (labeled *T_w*), that transports water from the thermocline to the tropics and then laterally into the subtropics. These circulations create teleconnections of nutrient supply in the surface ocean boxes. A bidirectional mixing term that ventilates the deep box directly through the high-latitude surface box (labeled *f_{hd}*) represents deep water formation in the Northern Atlantic region and around Antarctica (Sarmiento and Toggweiler, 1984). The parameters *T_c*, *T_w* and *f_{hd}* are considered adjustable parameters, which we calibrate using phosphate data from WOA13 (Garcia et al., 2014). In order to simulate the movement of particles, we included export fluxes (*P_t*, *P_s*, and *P_h*) of organic phosphorus out of each surface box.

Our box model simulates [*P*], alkalinity and various forms of *C*; total carbon in the surface boxes is partitioned into carbonate, bicarbonate, and *pCO₂*. The global mean [*P*] is prescribed according to the observed mean ocean value (Garcia et al., 2014). The export of carbon is linked to phosphorus export using the *C:P_{export}* ratio. To quantify the breakdown of carbon into these components, we model the solubility pump, using temperature and salinity to determine the partitioning of inorganic carbon among total carbon within a box. The global mean alkalinity is prescribed according to the observed mean ocean values but is also subject to transport (Sarmiento and Toggweiler, 1984). Our box model simulates alkalinity and total inorganic carbon, which are conserved tracers from which the speciation of inorganic carbon in sea-water can be calculated. Biome specific salinity and temperature are used to prescribe the solubility constants of *CO₂* in seawater and the bromine concentration, which is taken to be proportional to salinity. *CO₂* cycles through the atmosphere via the air-sea gas exchange fluxes (*f_{ah}*, *f_{as}*, *f_{at}*). We used a uniform piston velocity of $5.5 \times 10^{-5} \text{ m s}^{-1}$ to drive air-sea gas exchange (DeVries and Primeau, 2009; Follows et al., 2002).

We calibrated our model parameters (*T_c*, *T_w*, *f_{hd}*) so that the macronutrients were at similar average values compared to World Ocean Atlas 2013 dataset for each location. We tested the sensitivity of modeled *pCO_{2,atm}* to the fluxes *T_c*, *T_w*, and *f_{hd}* and found that with *T_c* = 20 Sv and *T_w* = 5 Sv (values that allowed the model to match [*P*] and alkalinity), the *pCO_{2,atm}* was sensitive to the value of *f_{hd}* (Sarmiento and Toggweiler, 1984). Guided by values previously used in the literature we set *f_{hd}* to 45.6 Sv (Table 2) but we also present results for the nutrient-only stoichiometry model at two extreme values of *f_{hd}*, 18 and 108 Sv (Figure 4). The functional dependence of *pCO_{2,atm}* with changing subtropical and tropical [*P*] for each extreme value of *f_{hd}* was quite similar, though the value of *pCO_{2,atm}* for the high *f_{hd}* simulation was approximately twice that of the low *f_{hd}* simulation (Figure 4). We found that our value of 45.6 Sv provides a modern *pCO_{2,atm}* value. Although the focus of this study is to determine the impact of low latitude biogeochemistry on *pCO_{2,atm}*, we point out that at

Redfield stoichiometry, $p\text{CO}_{2,\text{atm}}$ increases by 100 ppm when fhd is increased to 108 Sv from its default value 45.6 Sv. For certain values of the parameters, the model produced excessive nutrient trapping in the thermocline. In order to dampen the nutrient trapping, we tuned the remineralization depth. Assuming that 25% of the total export is respired in the thermocline with the remaining 75% exported into the deep ocean, produced a better match between the modeled and observed $[P]$ in the thermocline box. Total export is made from both the stoichiometry of sinking particulate and of primary producers, based on Teng et al. (2014) this is a reasonable first order assumption.

Table 2: High-latitude deep water exchange range

RANGE OF FHD [SV]	SOURCE
38.1	(Sarmiento and Toggweiler, 1984)
3-300	(Toggweiler, 1999)
60	(DeVries and Primeau, 2009)
30-130	(Galbraith and Martiny, 2015)
18-108 (default value 45.6)	This Study

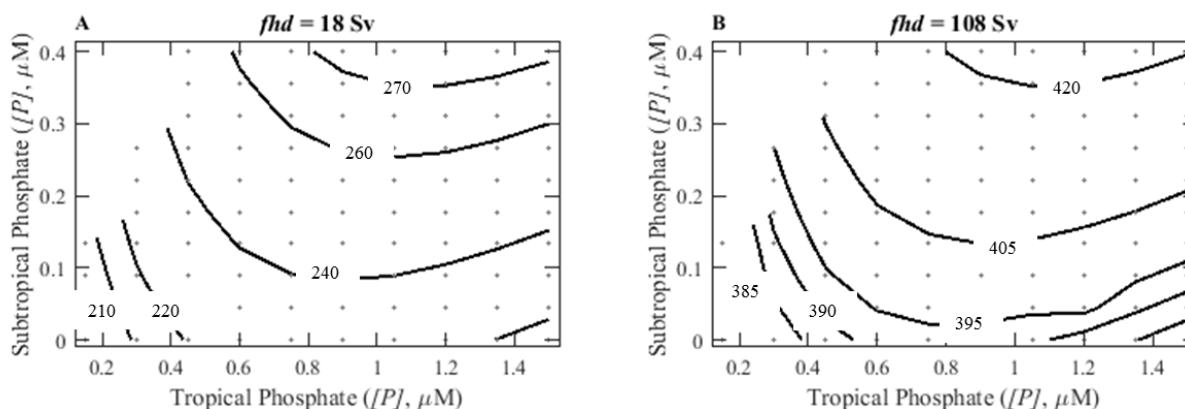


Figure 4: $p\text{CO}_{2,\text{atm}}$ (ppm) sensitivity to extreme fhd values under changing surface phosphate concentrations. A.) Range of $p\text{CO}_{2,\text{atm}}$ (ppm) using an fhd value of 18 Sv. B.) Range of $p\text{CO}_{2,\text{atm}}$ (ppm) using an fhd value of 108 Sv.

2.2.1 Experimental Design

To address how changing environmental conditions affected stoichiometric ratios, carbon export, and $p\text{CO}_{2,\text{atm}}$ we performed two tests; a change in nutrients and a change in sea surface temperature. These tests allowed us to observe how the relationships between environmental conditions, carbon export and $p\text{CO}_{2,\text{atm}}$, depend on the mechanisms responsible for stoichiometric variation in the ocean. In order to account for the effects of

particulate inorganic carbon (PIC) export, we multiply model predicted $C:P_{export}$ by 1.2, consistent with previous studies (Broecker, 1982; Sarmiento and Toggweiler, 1984).

The first set of numerical experiments examined the sensitivity of $pCO_{2,atm}$ to nutrient availability in the tropical and subtropical boxes for each of the three stoichiometric models. We varied tropical $[P]$ from 0.15 to 1.5 μM and subtropical $[P]$ from 1×10^{-3} to 0.5 μM by adjusting the implied biological export and determined the equilibrium $pCO_{2,atm}$ values.

The second set of experimental tests was done to quantify how temperature modifies carbon export and $pCO_{2,atm}$ for each stoichiometric model. Temperature influences carbon cycling in two ways within our model: through the solubility of inorganic carbon in seawater and through changes in phytoplankton stoichiometry within the temperature-only and multi-environmental models. Due to the well-known effects of temperature on CO_2 solubility, it is generally predicted that there is to be a positive feedback between $pCO_{2,atm}$ and temperature mediated by declining CO_2 solubility at high temperatures. To study the relative strengths of the temperature solubility feedback and the temperature regulation of C:P feedback, we performed a numerical experiment in which we varied the sea surface temperature by five degrees in either direction of modern sea surface temperature. This represents a plausible range of variation under both ice-age and anthropogenic climate change scenarios. We varied tropical temperature from 21° to 31°C and subtropical temperature from 19° to 29°C, determining equilibrium $pCO_{2,atm}$ values for combinations of temperature conditions.

3 Results

To quantify the linkages between phytoplankton physiology, elemental stoichiometry, and ocean carbon cycling, we divide our results into two parts. The first is a direct study of the stoichiometric models, comparing their predictions about the relationship between stoichiometry and environmental conditions, and in the case of the trait-based model, illustrating how cellular physiology is predicted to vary across these conditions. In the second part, we show how variable stoichiometry influences carbon export and $pCO_{2,atm}$, under changing phosphorus concentrations and temperature. Within these results, we distinguish the influence or lack thereof of the three distinct biomes; in particular the equatorial upwelling regions and the macronutrient depleted subtropical gyres.

3.1 Multi-environmental and physiological controls on plankton stoichiometry

Our multi-environmental model captured several major mechanisms hypothesized to be environmental drivers of C:P ratios including a temperature dependence of many cellular processes, a link between growth rate and ribosome abundance, and storage drawdown during nutrient limitation. The predicted relationship between environmental conditions and C:P can be understood through the environmental regulation of three factors: (i) the balance between photosynthetic proteins and ribosomes, (ii) the cell radius and associated allocation to structural material, and (iii) the degree of phosphorus storage. Our model predicted that for an optimal strategy, specific protein synthesis rates will match specific rates of carbon fixation. Thus, the ratio of photosynthetic machinery to biosynthetic machinery is primarily controlled by irradiance and temperature. Increases in light levels lead to higher photosynthetic efficiency, higher ribosome content, smaller cells (due to a

lower requirement for photosynthetic machinery), and lower C:P ratios (Figure 5). The response of C:P to light levels predicted by our model was muted in comparison to other subcellular compartment models because we separately modeled electron transport and carbon fixation (Talmy et al., 2013), and our predictions were consistent with the weak relationship between irradiance and C:P (Thrane et al., 2016) (Figure 5A).

Increases in temperature increase the efficiency of biosynthesis, but not photosynthesis ($Q_{10} = 1$). Therefore elevated temperature lead to a reduced ribosome content relative to photosynthetic proteins and higher C:P ratios (Figure 6A). There leads to a non-monotonic, concave relationship between temperature and cell size, which is due to a subtle interaction between biosynthesis efficiency (which varies greatly with temperature), maintenance costs, and size dependent uptake rates.

Nutrient concentrations do not affect the ratio of biosynthetic to photosynthetic machinery but positively relate to both P storage and cell radius. Cell radius directly influences the specific rate of nutrient uptake, and indirectly biosynthesis and photosynthesis as the cell membrane and wall affects the space available for other investments. This effect is pronounced in oligotrophic conditions ($[P] < 100\text{nM}$), cell radius declines below $1\mu\text{m}$, decreasing the allocations to both photosynthesis and biosynthesis and driving up C:P ratios. Much larger values of the cell radius are observed at high nutrient concentrations.

P concentrations also influenced C:P through their direct control of P storage. We plotted the relative contribution of the storage compartment and the functional compartment to the P quota, as a function of environmental conditions. The impact of the residual pool on the overall size of the P pool is heavily dependent on environmental conditions, varying from a minimum of close to 0% to a maximum of just under 50%, for the combinations of parameter values used in all of our numerical experiments. In the vast majority of the parameter range considered here, the contribution of the residual pool is much more modest, 10-20%. High values occur when phosphorus is available and the temperature is high. In these conditions, ribosomal contributions are decreased, but the residual contribution is high. In cold water, high P ecosystems, the residual contribution is approximately 25%, and in oligotrophic ecosystems it is close to 0. Thus, C:P was predicted to be a decreasing function of $[P]$ with two distinct regimes: a moderate sensitivity regime for $[P]$ above 100nM , and a high sensitivity regime for $[P]$ below 100nM .

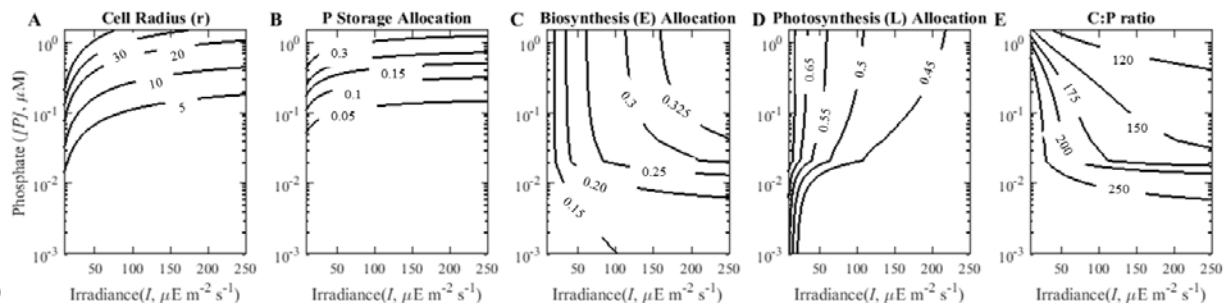


Figure 5: Influence of phosphate concentration and irradiance on cellular stoichiometry and cellular traits, at a constant $T = 25\text{ °C}$. A) Cell radius (r). B) P storage allocation. C) Biosynthesis allocation. D) Photosynthesis (L) allocation. E) The C:P ratio. As irradiance increases, there is a tendency towards greater allocation to biosynthesis and lesser allocation to photosynthesis, which leads to lower C:P ratios. When phosphorus is very low, there is a large decrease in both biosynthesis and photosynthesis allocations due to the

large relative allocation to the cell membrane. C:P ratios are inversely proportional to phosphorus concentration, driven by an increase in luxury storage and ribosomal content as P increases.

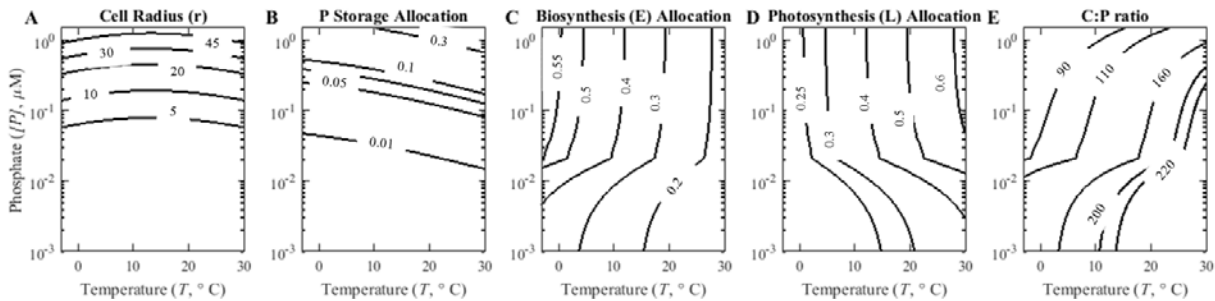


Figure 6: Influence of phosphate concentration and temperature on cellular stoichiometry and cellular traits, at a constant irradiance $I = 50 \mu E m^{-2} s^{-1}$. A) Cell radius (r). B) P storage allocation. C) Biosynthesis allocation. D) Photosynthesis (L) allocation. E) The C:P ratio. Consistent with the translation compensation hypothesis, increases in T led to a reduction in the allocation to biosynthesis and an increase in C:P.

We next used the outcome of the trait model as a multi-environmental model to predict C:P ratios in the modern ocean based on annual mean light, temperature, and [P]. Our predictions reproduced the global pattern (Martiny et al., 2014) with C:P ratios above the Redfield ratio in subtropical gyres and C:P ratios below the Redfield ratio in equatorial and coastal upwelling regions and subpolar gyres (Figure 7A). Additionally, our model also reproduced basin-scale stoichiometric gradients among similar biomes in each ocean, predicting the highest C:P ratios in the western Mediterranean Sea and the western North Atlantic Subtropical Gyre, and somewhat elevated C:P ratios in the South Atlantic Subtropical Gyre as well as the North and South Pacific Subtropical Gyres.

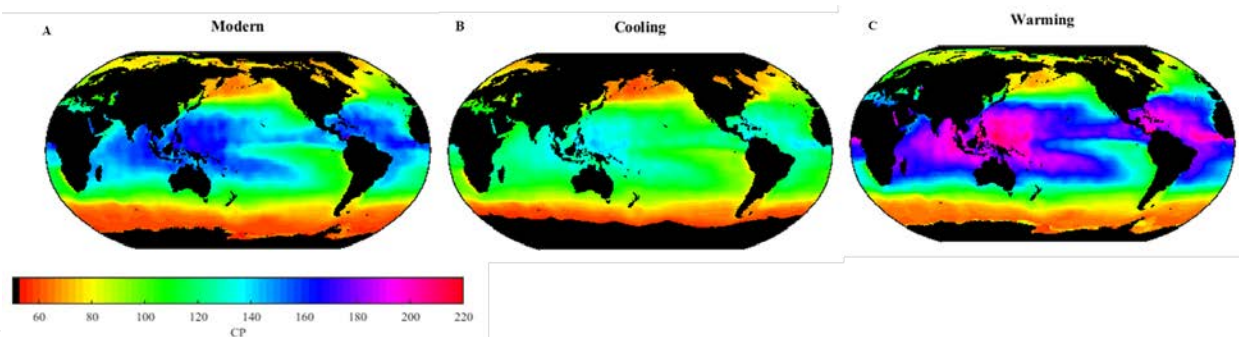


Figure 7: Predicted C:P ratios in the global ocean in differing climatic regimes. A) C:P ratio under modern ocean conditions. Large differences in C:P are predicted between distinct types of ocean biome, with low C:P in equatorial upwelling regions and subpolar gyres, and high C:P in subtropical gyres. Regional differences between biomes of similar type are observed as well, with the low phosphorus Atlantic having a higher C:P than the Pacific. B.) C:P ratio under cooling temperature conditions ($-5^{\circ}C$ from the modern ocean). C) C:P ratio under warming temperature conditions ($+5^{\circ}C$ from the modern ocean). Each 5 degree change leads to a shift of 15% in the mean C:P ratio of organic matter.

To study the potential impact of sea surface temperature on phytoplankton resource allocation and stoichiometry, we used our multi-environmental model to predict C:P in ocean conditions both five degrees colder (Cooling environments) and warmer (Warming environments) than the modern ocean. According to our model, a five-degree increase (or

decrease) in sea surface temperature would cause a 15% rise (or fall) in C:P ratios (Figure 7). This sensitivity suggested that the relative effect of *temperature* on biochemical processes could have large implications for biogeochemical cycles, making it important to determine the relative importance of physiological mechanisms regulating C:P ratios.

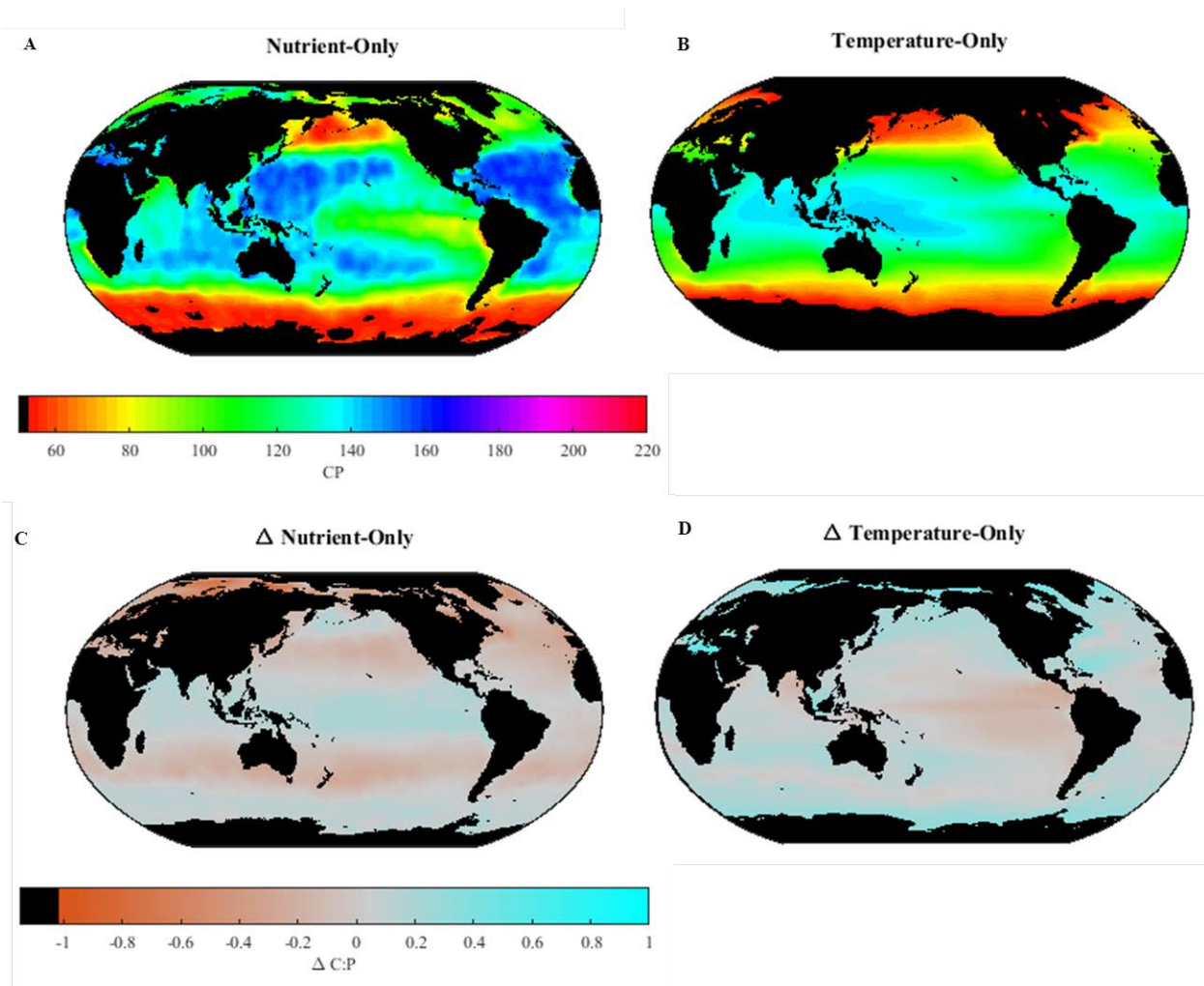


Figure 8: Comparison of C:P between the multi-environmental model and the nutrient-only model and temperature-only model. The upper panels show predicted C:P for the global ocean under the nutrient-only (A) and temperature-only (B) models, and the lower panels show the normalized difference, i. e. $\frac{C:P_{subcell} - C:P_{other}}{C:P_{subcell}}$, between the C:P in the subcellular model (C, D).

We compared the multi-environmental model to the predictions made by two other models: the nutrient-only model used by the Galbraith and Martiny model (2015), and our temperature-only model modified from Yvon-Durocher and co-workers (2015). These two models also successfully predicted the qualitative pattern of stoichiometric variation in the ocean, but were unable to replicate the full range of variation observed in the data (Figure 8). In particular, they misrepresented the North Atlantic Subtropical Gyre and the Southern Ocean, where the C:P ratio is at the extreme (Figure 8A, B). The nutrient-only model had a tendency to predict lower C:P ratios than the multi-environmental model in warm tropical

and subtropical waters, and predict higher C:P ratios in cold waters (Figure 8A). This difference is driven by the *temperature* sensitivity of biosynthesis in the multi-environmental model, leading to increasing C:P in all warm water regions and decreasing C:P in cold water regions (Figure 8C). The multi-environmental model predicted a wider range of C:P in the ocean. The temperature-only model overall had higher C:P ratios globally compared with the multi-environmental model (Figure 8B) but suggested lower C:P in the gyres and higher C:P in high latitudes, especially in the Southern Ocean (Figure 8D).

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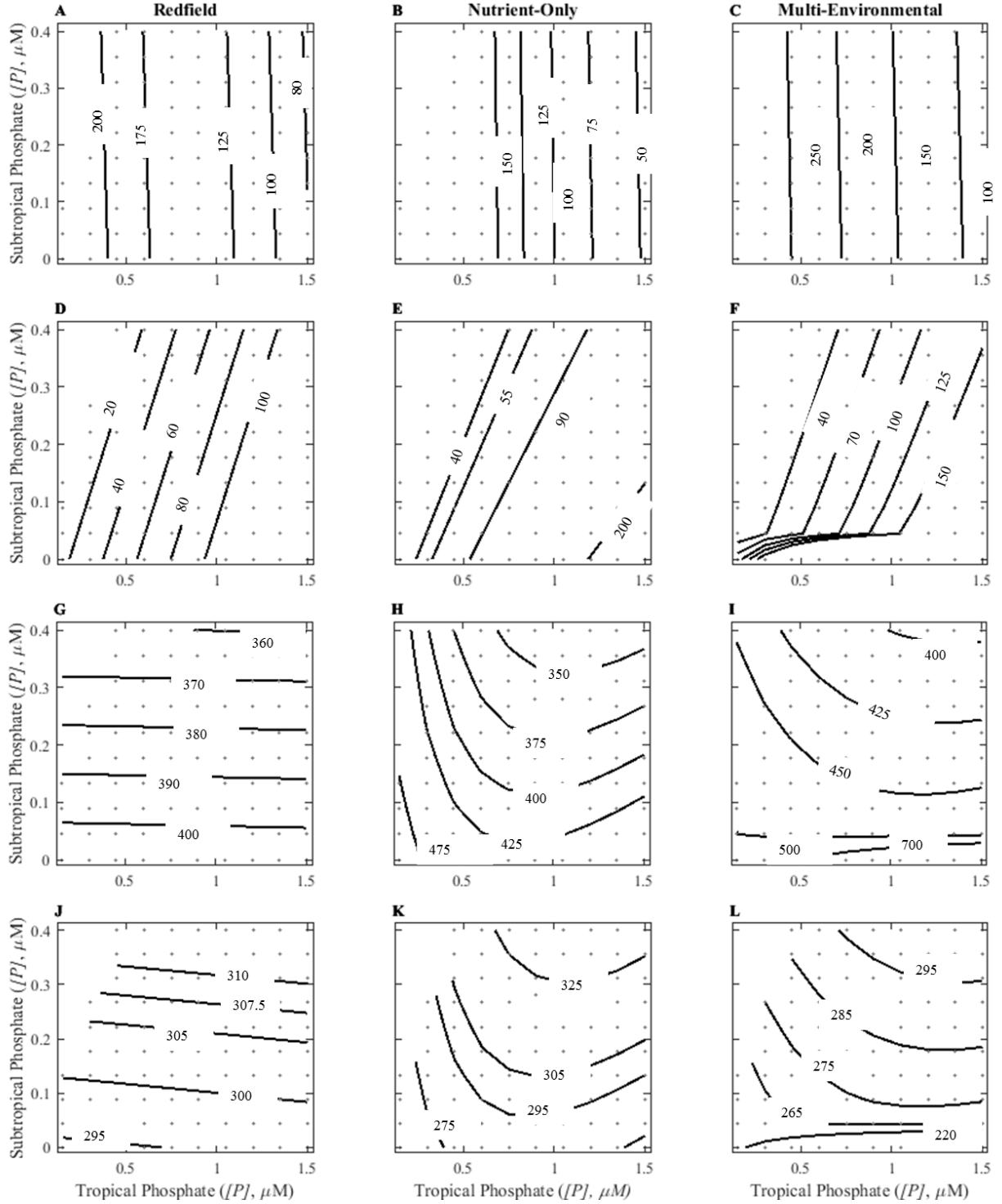
3.2 Impact of nutrient availability on carbon export and atmospheric pCO₂

We next quantified the impact of nutrient availability in the tropics and subtropics on stoichiometry, carbon export and pCO_{2,atm} (Figure 9A-L). Using a constant Redfield model (or the temperature-only model), we replicated the previously observed approximately linear relationship between surface [P] and pCO_{2,atm} (equivalent to how pre-formed [P] will influence pCO_{2,atm}) (Ito and Follows, 2003; Sigman and Boyle, 2000). We found that [P] drawdown in the subtropical box had a greater impact on carbon export, since export from the high-latitude box was not enhanced by the [P] supply from the subtropical box (Figure 9A, D, G). In the Redfield model, pCO_{2,atm} appeared to be much more sensitive to subtropical [P] than tropical [P], which was partially due to enhanced carbon export in the subtropical box and partially due to the larger surface area of the subtropical box (implying a greater potential for CO₂ exchange) (Figure 9J).

In contrast to the predictions made using Redfield stoichiometry, when we used the nutrient-only model for phytoplankton stoichiometry, we observed a non-linear relationship between surface [P] and pCO_{2,atm} (Figure 9B, E, H, K). At fixed tropical [P], there was a strong relationship between subtropical [P], export, and pCO_{2,atm} in accordance with the findings of Galbraith and Martiny (2015) (Figure 9B, E, H). The total decline in pCO_{2,atm} as subtropical [P] declined from 0.4 μM to 1x10⁻³ μM could be more than 60 ppm, which was more than twice the decline that occurred in the fixed stoichiometry experiment (Figure 9K). We found a non-linear monotonic relationship between tropical [P] and pCO_{2,atm}: when tropical [P] was high, declines in tropical [P] led to lower carbon export and increased pCO_{2,atm}. However, this trend reversed when tropical [P] was lower (Figure 9K). The counter intuitive decline in pCO_{2,atm} with higher export from tropics was driven by a teleconnection in nutrient delivery between the subtropical and tropical boxes. Increases in export in the tropical box due to increased [P] drawdown decreased the supply of [P] to the subtropics, which led to a decrease in the more efficient (higher C:P) subtropical export. Thus, the nutrient-only model predicted a greater decrease in subtropical export than the counter increase in tropical export.

The multi-environmental model also predicted a non-linear relationship between surface P, carbon export, and pCO_{2,atm}. However, the pattern was somewhat distinct from that of the nutrient-only model results (Figure 9C, F, I, L). First, subtropical [P] drawdown had a nonlinear relationship with pCO_{2,atm}: when subtropical [P] was high, declines in subtropical [P] led to slight declines in pCO_{2,atm}, and when subtropical [P] is low, small declines in tropical [P] lead to large declines in pCO_{2,atm}. This intensification of the relationship between subtropical [P] and pCO_{2,atm} was due to the nonlinear relationship between [P] and C:P predicted by the trait-based model (Figure 9I). The multi-environmental model predicted extremely high export, but only when [P] was lower than

0.05 μM (Figure 9C, F, I). Second, the effect of tropical [P] levels on $\text{pCO}_{2,\text{atm}}$ was strongly modulated by subtropical [P], reversing from a negative to a positive relationship as subtropical [P] declines (Figure 9I, L). The difference between the nutrient-only model and the multi-environmental model arose because the multi-environmental model incorporated a temperature impact on resource allocation and elemental ratios. Although we were not varying temperature in these experiments, we did represent regional temperatures differences between the different boxes. The result is that a large stoichiometric contrast between the tropical and sub-tropical regions only arose when there was a large difference in nutrient levels between the two regions (Fig. 9L). However, both the nutrient-only model and the multi-environmental model predicted that carbon export and $\text{pCO}_{2,\text{atm}}$ were sensitive to the interaction between regional nutrient availability and $\text{C:P}_{\text{export}}$.



635 **Figure 9: Carbon export (Tmol C yr^{-1}) and $\text{pCO}_{2,\text{atm}}$ (ppm) in changing surface phosphate concentrations.**
 Columns correspond to type of stoichiometry; Redfield (Left), nutrient-only (Middle), and multi-environmental
 model (Right). Rows correspond to either tropical carbon export (A through C), subtropical carbon export (D
 through F), total carbon export (G through I) or atmospheric pCO_2 (J through L). The grey points represent
 640 where $\text{pCO}_{2,\text{atm}}$ was calculated, between spaces are interpolated.

3.3 Interactive effect of temperature on stoichiometry, carbon export and atmospheric $p\text{CO}_2$

We next quantified the impact of temperature in the tropics and subtropics on $\text{C:P}_{\text{export}}$, carbon export, and $p\text{CO}_{2,\text{atm}}$ (Figure 10A-D). The Redfield model predicts that increases in temperature lead to a decline in the solubility of CO_2 in seawater and consequently an increase in $p\text{CO}_{2,\text{atm}}$ from 288 to 300 ppm ($\Delta p\text{CO}_{2,\text{atm}} = 12$) (Figure 10A). This feedback was present with the same strength in the nutrient-only model (with no temperature dependence on C:P), in which $p\text{CO}_{2,\text{atm}}$ ranged from 268 to 280 ppm ($\Delta p\text{CO}_{2,\text{atm}} = 12$) (Figure 10B).

In contrast to the Redfield and nutrient-only models, the temperature-only model predicted a negative linear relationship between $p\text{CO}_{2,\text{atm}}$ and tropical temperature and a positive linear relationship between $p\text{CO}_{2,\text{atm}}$ and subtropical temperature (Figure 10C). The decline in $p\text{CO}_{2,\text{atm}}$ with tropical temperature was driven by an enhancement of export due to increased C:P at higher temperatures (Figure 11). At 5°C below modern ocean temperature, the model predicted C:P in the tropics was 131 and subtropical was 121, resulting in a $p\text{CO}_{2,\text{atm}}$ of 305 ppm. At 5°C above modern ocean temperature, the model predicts a C:P ratio in the tropics of 189 and C:P ratio of 175 in the subtropics, resulting in a $p\text{CO}_{2,\text{atm}}$ of 263 ppm. Tropical temperature had more impact with $\Delta p\text{CO}_{2,\text{atm}} = 41$ ppm compared to subtropical temperature with a $\Delta p\text{CO}_{2,\text{atm}}$ ranging from 4 to 5 ppm (Figure 11).

Similar to the temperature-only model, the multi-environmental model predicted a negative linear relationship between $p\text{CO}_{2,\text{atm}}$ and tropical temperature and a positive linear relationship between $p\text{CO}_{2,\text{atm}}$ and subtropical temperature (Figure 10D). The decline in $p\text{CO}_{2,\text{atm}}$ with tropical temperature was driven by an enhancement of export due to increased C:P at higher temperatures (Figure 11). In the subtropical region, the expected increase in export was mitigated by a decline in solubility. At 5°C below modern ocean temperature, the trait-based model predicted that C:P in the tropics was 147 and that C:P in the subtropics was 155, resulting in an increase of $p\text{CO}_{2,\text{atm}}$ to 279 ppm (Figure 11). Variation in tropical SST over a 10°C span led to a significant decline in $p\text{CO}_{2,\text{atm}}$, with a $\Delta p\text{CO}_{2,\text{atm}}$ of approximately 46, and tropical C:P ranging from 147 to 210 (Figure 11). Because the subtropical box has a large surface area, the decrease in surface CO_2 solubility at high temperatures is sufficient to overcome the increase in export due to higher C:P leading to a positive relationship between $p\text{CO}_{2,\text{atm}}$ and subtropical temperatures.

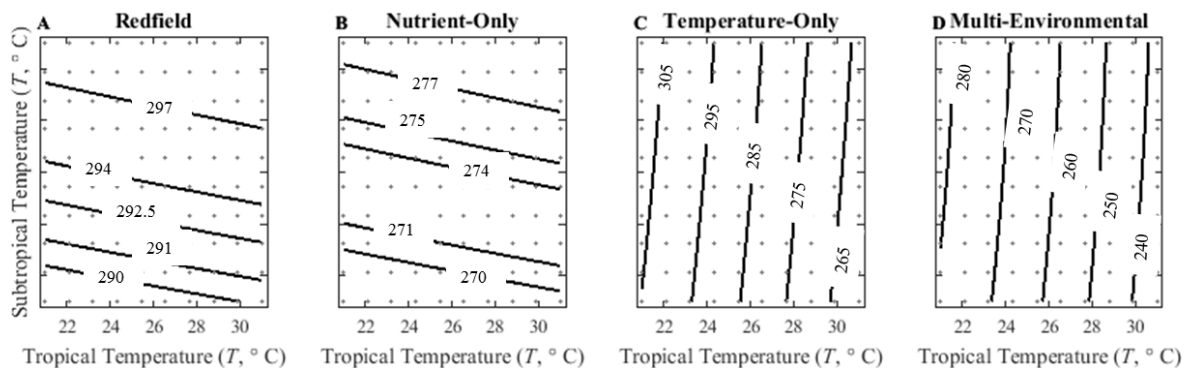
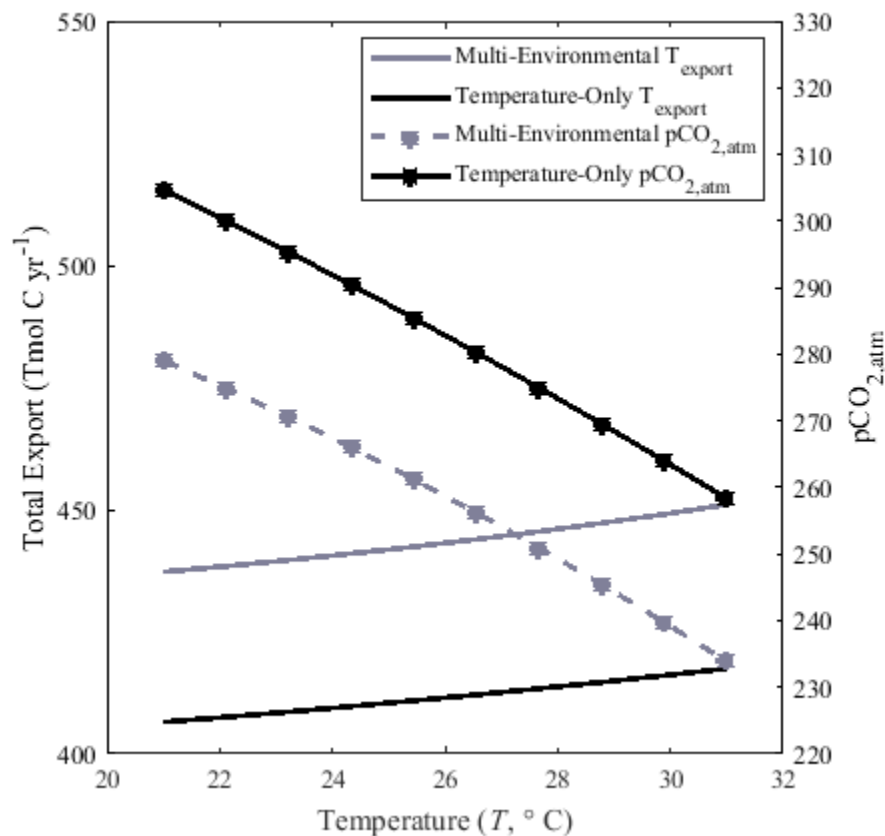


Figure 10: $p\text{CO}_{2,\text{atm}}$ (ppm) as a function of changing surface temperature concentrations. Based on A) Redfield (fixed) stoichiometry model, B) nutrient-only stoichiometry model, C) temperature-only stoichiometry model, and D) multi-environmental stoichiometry model.



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Figure 11: The effect of changing sea surface temperature (°C) on $p\text{CO}_{2,\text{atm}}$ and total carbon export (Tmol C yr⁻¹) in the temperature-only and multi-environmental model. Phosphate concentrations are 0.3 μM in the tropical and 0.05 μM in the subtropical box. Multi-environmental model total carbon export is the solid gray line, and $p\text{CO}_{2,\text{atm}}$ is the dashed gray line. Temperature-only model total carbon export is the solid black line, and $p\text{CO}_{2,\text{atm}}$ is the dashed black line.

4 Discussion

Here, we found that variable stoichiometry of exported organic material moderates the interaction between low-latitude nutrient fluxes and ocean carbon cycling. A full connecting circulation allows for complete movement of nutrients between ocean regions resulting in strong linkages between nutrient supply ratios and cellular stoichiometric ratios (Deutsch and Weber, 2012). It has been shown that the inclusion of an oceanic circulation connecting high and low-latitude regions results in a feedback effect between high-latitude nutrient export and relative nutrient supply in low-latitudes (Sarmiento et al., 2004; Weber and Deutsch, 2010). Together, the inclusion of lateral transport between ocean regions and of deviations from Redfield stoichiometry within our model led us to predict the existence of strong teleconnections between the tropics and the macronutrient limited subtropics. The degree of nutrient drawdown in the tropics had a strongly non-monotonic relationship with $p\text{CO}_{2,\text{atm}}$ because this drawdown influenced both nutrient

supply to the subtropics and tropical C:P. The idea of biogeochemical teleconnections has been proposed before, but we found that variations in stoichiometry greatly enhance the importance and strength of such linkages (Sarmiento and Toggweiler, 1984). Thus biome-scale variations in phytoplankton elemental stoichiometry may change the sensitivity of the carbon pump to other phenomena that regulate patterns of nutrient drawdown. We also see that the degree of nutrient drawdown had a strong impact on predicted (and observed) C:P leading to highly non-linear controls on $p\text{CO}_{2,\text{atm}}$ whereby increased export in the tropics counter intuitively leads to increasing $p\text{CO}_{2,\text{atm}}$. Large-scale gradients in stoichiometry can alter the regional efficiency of the biological pump: [P] supplied to high C:P regions leads to a larger export of carbon than [P] supplied to low C:P regions, giving an important role to the details of the ocean circulation and other processes that alter nutrient supply and phytoplankton physiological responses in different surface ocean regions. Therefore, biome-scale variations in phytoplankton elemental stoichiometry can lead to a fundamental change in the partitioning of carbon between the atmosphere and the ocean.

We have created a box model to simulate the impact of the low latitude stoichiometric ratios, its environmental controlling factors and its relationships on $p\text{CO}_{2,\text{atm}}$. Low latitude phosphorus concentrations can be set in one of two fashions; through iron limitation and through nutrient availability. Here we will briefly discuss how iron limitation could play a significant role on phosphorus concentrations. The biogeochemical functioning of tropical regions are commonly influenced by iron availability in such a way that macronutrient levels cannot be fully drawn down by phytoplankton (Coale et al., 1996; Moore, 2004; Raven et al., 1999). The degree of nutrient drawdown has a strong impact on predicted (and observed) C:P. This environmental control on C:P could lead to highly non-linear controls on $p\text{CO}_{2,\text{atm}}$ whereby increased export in the tropics leads to increasing $p\text{CO}_{2,\text{atm}}$. This relationship would differ in the subtropics, where iron is thought to stimulate nitrogen levels through nitrogen fixation, an iron exhaustive metabolic process (Wu et al., 2000). Iron's potential control on nitrogen fixation could promote higher carbon fixation and further exported stoichiometric ratios in the subtropical regions leading to increasing $p\text{CO}_{2,\text{atm}}$ (Wu et al., 2000). Thus, iron availability may play a complex role depending on whether there is an increased delivery in upwelling zones (leading to a potential declining global C export) or in the subtropical gyres (leading to a potential increase in global C export).

Past studies using box models have found $p\text{CO}_{2,\text{atm}}$ to be insensitive to low-latitude nutrients (Follows et al., 2002; Ito and Follows, 2003; Sarmiento and Toggweiler, 1984; Toggweiler, 1999). This phenomena was explored by DeVries and Primeau (2009), who showed that the strength of the thermohaline circulation is the strongest control on $p\text{CO}_{2,\text{atm}}$, and that changes in low-latitude export are relatively unimportant. Unlike our study, such earlier work relied on a uniform Redfield stoichiometry. However, we find that when stoichiometric variation is included, carbon export and $p\text{CO}_{2,\text{atm}}$ become dependent on details of low-latitude processes.

It is important to recognize that a five-box model is an incomplete description of ocean circulation, and is meant only to identify the most important mechanisms, not to make precise quantitative predictions. In order for our model to adequately reflect important features of the carbon and phosphorus nutrient distributions, we had to carefully select the values of the thermohaline and wind-driven upper ocean circulations that lead to reasonable nutrient fluxes and standing stocks. The value of thermocline

circulation, T_c , has been calibrated in different box models to range from 12 to 30 Sv (DeVries and Primeau, 2009; Galbraith and Martiny, 2015; Sarmiento and Toggweiler, 1984; Toggweiler, 1999). Representation of the wind driven overturning, T_w , in a simple box model has received less attention. Variations in the thermohaline circulation influence the abundance of nutrients in different boxes. Depending on the strength of this circulation, our model accumulated nutrients in the thermocline box and we tuned this parameter to most accurately mimic nutrient variation across ocean regions. Another caveat relates to our choice of the two-way flux values. Similar to the circulation values, a wide range of two-way flux values have been used in the literature. We therefore performed sensitivity experiments to find the best value for our full model set-up but the qualitative trends observed are insensitive to the choice of such fluxes.

Nutrient availability and temperature have been alternatively proposed as drivers of variation in stoichiometric ratios in the global ocean, and the strong statistical correlation between temperature and nutrients throughout the ocean has prevented identification of the relative importance of each factor (Martiny et al., 2013b; Moreno and Martiny, 2018). We see that although temperature regulation of $C:P_{\text{export}}$ can influence $pCO_{2,\text{atm}}$, this regulation is strongly dependent on the detailed control mechanism and also generally diverge from expectations based on the solubility pump. The decrease in surface CO_2 solubility at elevated temperature is sufficient to overcome the increase in export due to higher C:P leading to a positive relationship between $pCO_{2,\text{atm}}$ and subtropical temperatures. It is important to point out that the relative importance of the two competing effect depends critically on the physical circulation of the ocean. Predicted increases in stratification are often invoked as a mechanism that would decrease the vertical supply of nutrients, which one might think would further compensate for the effect of higher C:P. However, the strength of the biological pump in the subtropics is also influenced by lateral transport of nutrients (Letscher et al., 2015) so we argue that it is unclear if you should expect increasing, unchanged or decreasing C export in low latitude regions with ocean warming and stratification. Similarly, it is unclear how increases in stratification might affect the strength of the solubility pump. The sensitivity of $pCO_{2,\text{atm}}$ to changes in subtropical temperatures depends critically on the volume of the ocean ventilated from the subtropics, i.e. on the volume of the thermocline box in our model. How this volume might change in response to a warming world is a complicated dynamical problem that is beyond the scope of the present work.

Our results do not identify whether temperature or nutrient concentrations is the most important driver of phytoplankton C:P, but do suggest that the physiological effect of temperature could be important for ocean carbon cycling. Both the temperature-only and multi-environmental models predict that temperature increases enhance tropical export, causing substantial decreases in $pCO_{2,\text{atm}}$ with temperature. This relationship is the reverse of that predicted by the nutrient-only and Redfield models, and represents a sizable potential negative feedback on carbon cycling. The multi-environmental model also predicted that C:P responds in a nonlinear fashion to $[P]$, with significantly increased sensitivity in highly oligotrophic conditions. Thus, a deeper understanding of the physiological mechanisms regulating phytoplankton C:P ratios are key to understanding the carbon cycle.

Our derivation of the multi-environmental model relies on several important assumptions. The growth rate in the multi-environmental model is determined by a set of

environmental conditions and quantified by the specific rate of protein synthesis, carbon fixation and phosphorus uptake. The effect of growth rate on stoichiometry will likely be dependent on whether light, a specific nutrient, or temperature controls growth. The value of specific species of Q_{10} leads to uncertainty in our multi-environmental model because of the range of possible values is highly dependent on the cell or organism being tested. In a study examining Q_{10} of various processes within the cell, it was found that the Q_{10} of photochemical processes ranged from 1.0 to 2.08, and for carboxylase activity of RuBisCO to be 2.66 (Raven and Geider, 1988). In addition to the high uncertainty between Q_{10} values, there is high ambiguity associated with cellular inorganic P stores (e.g., polyphosphates and phospholipids) (Kornberg et al., 1999). P storage, such as polyphosphates, can serve as both energy and nutrient storage that may be regulated by unique environmental factors. Finally, we assume that our choice of the value of Q_{10} for each metabolic process is a potential source of error within our model, because measured values are highly dependent on the cell or organism being tested, and it is difficult to extend these single-organism observations across species. Thus, we recognize multiple caveats within the trait-based model but expect that it improves our ability to link environmental and phytoplankton stoichiometry variation.

5 Conclusions

We find that processes that affect nutrient supply in oligotrophic gyres, such as the strength of the thermohaline circulation, are particularly important in setting $pCO_{2,atm}$ but via a complex link with $C:P_{export}$. By explicitly modeling the shallow overturning circulation, we showed that increased export in the tropics, which might be influenced by increased atmospheric iron dust deposition, may lead to increases, rather than decreases, in $pCO_{2,atm}$. Increased [P] drawdown in the tropics shifts export away from the subtropical gyres, and changes the mean export C:P in the low-latitude ocean. We would expect that nutrient drawdown leads to high export and declines in $pCO_{2,atm}$, but instead we find that variation in cellular allocation and adaptation can lead to counterintuitive controls on $pCO_{2,atm}$. Additionally, we find that it is even more difficult to separate nutrient supply and temperature controls on marine phytoplankton stoichiometry, carbon export, and $pCO_{2,atm}$ and we need better physiological experiments and field data to fully understand the relative impact of the two factors. Nevertheless, it is likely that both play a key role in regulating phytoplankton stoichiometry, $C:P_{export}$, and ultimately ocean carbon cycling.

Author Contribution: ARM - creation and analysis of the box model and primary writer of manuscript. GIH - creation and development of the trait-based model, and writing. FWP - assistance on the box model and editing of manuscript. SAL - assistance on the trait-based model and editing of manuscript. ACM - assistance on both models and writing of manuscript.

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