





Is There A Universal Relationship Between Phytoplankton Specific Growth Rate And Temperature?

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Metabolic Theory of Ecology (MTE)

predicts: thermal sensitivity of a biological rate

- = "activation energy" (E_a)
- = slope

(Gillooly et al 2001)



Boltzmann constant = k = 8.62 x 10⁻⁵ eV. K⁻¹

Metabolic Theory of Ecology (MTE)

predicts: $E_a \sim 0.32 \text{ eV}$

for photosynthesis

(Allen et al. 2005)







Metabolic Theory of Ecology (MTE)

<u>assumes</u> E_a:

is the same for

- acute
- acclimatory
 - &
- evolutionary

responses to temperature







In Rate

Metabolic Theory of Ecology (MTE)

predicts *E_a*:

- is the same for
- photosynthesis

&

specific growth rate, μ
(= intrinsic rate of natural increase, r)

(Savage et al. 2004)





Tests of Prediction and Assumption

 E_a for microalgal specific growth rate is:

- i) 0.32 eV
- ii) The <u>same</u> for <u>acclimation</u> (within-species) and <u>evolutionary adaptation</u> (across-species)



Data

- 30 published single-sp growth responses to acclimation temperature
- Species from 8 divisions (27 spp, 3 strains)
- > 5 orders of magnitude of cell volume
- Temperature range ~30 °C





Methods: Sensitivity of *μ* to Temperature: Within-spp *versus* Across-spp

Stressfully high, supraoptimal temperatures, not included



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Stressfully high, supraoptimal temperatures, not included In Rate Cell size and light climate corrections followed López-Urrutia et al. (2006) 37 40 42 38 39 43 41 Ln (Rate. (Cell vol) -b). 1/(k.T) (eV⁻¹) ((PFD. Kpfd)/PFD) Temperature

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Reduced other confounding effects (e.g. nutrient limitation differences) : slope of <u>top</u> of data envelope by quantile regression (large triangles)

Results: *E_a* Across-species

- A. Microalgae (N = 138; 96th quantile): $E_a = 0.48$
- C. Diatoms (N = 52; 90th quantile): $E_a = 0.41$



Results: Withinspecies *E_a*

- Frequency distributions of activation energies
- Dotted horizontal lines: weighted means ± 2 SE
- A. Microalgae (N = 30 responses).

Mean $E_a = 0.46$

B. Diatoms (N = 11). Mean $E_a = 0.53$



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Solid line: Mean E_a (± 2 SE) from across-species quantile regression



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No: Mean $E_a = 0.41 - 0.53 \text{ eV}$

p<0.001, ex. diatoms within-spp = NS

→ ~25% higher µ over 10 °C compared with MTE

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Yes

 iii) The <u>same</u> for <u>acclimation</u> (within-species) and <u>evolutionary adaptation</u> (across-species)

Explaining These Patterns of E_a

Hypothesis: Thermal sensitivity is reduced where maximum growth falls below top of data envelope

(e.g. under nutrient limitation, nutrient diffusion (which has low E_a) has larger influence on specific growth rate, thus lowering E_a of μ)



Results: *E_a* Across-species

A. Microalgae (N = 138; 96th quantile): $E_a = 0.48$ 50th quantile (median): $E_a = 0.28$

Not significantly diff from 0.32 eV

B. Diatoms (N = 52; 90th quantile): $E_a = 0.41$



Conclusions

- Microalgal specific growth rate (μ) is 25% higher per 10 °C than expected from MTE, both within species and for μ_{max} across-species
- E_a within species is similar to that <u>across</u> species at maximum μ
- We hypothesize: E_a is reduced by nutrient limitation of μ (a nutrient x temperature interactive effect on μ). This could explain lower E_a for average community μ (mean or median) compared with maximum μ
- Research needed to better understand nutrient x temperature interactions

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Results: E_a Acrossspecies

C. Diatoms (
$$\mu_{max}$$
 only; N = 11)
E_a = 0.47

[Microalgae (μ_{max} only; N = 16): NS regression, not shown]

