Improving biogeochemical models through single-cell and taxon-specific analyses of marine plankton physiological plasticity, genetic diversity and evolutionary processes

Michael Lomas

(workshop steering committee: Susanne Neuer, Steph Dutkiewicz, Steve Giovanonni, Adam Martiny, Ben Twining, Ramunas Stepanauskas, Adrian Marchetti, Alison Taylor)

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Emergent Biogeography of Microbial Communities in a Model Ocean

Michael J. Follows,¹* Stephanie Dutkiewicz,¹ Scott Grant,^{1,2} Sallie W. Chisholm³

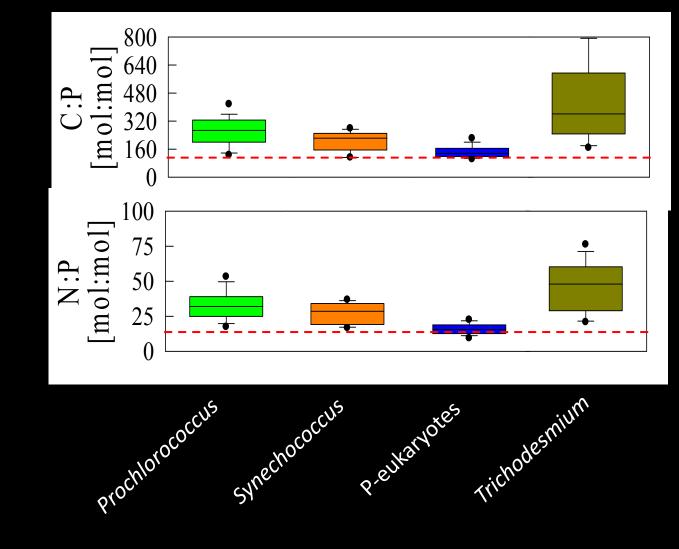
Variations in *Synechococcus* cell quotas of phosphorus, sulfur, manganese, iron, nickel, and zinc within mesoscale eddies in the Sargasso Sea

Benjamin S. Twining,^{a,1,*} Daliangelis Nuñez-Milland,^a Stefan Vogt,^b Rodney S. Johnson,^c and Peter N. Sedwick^d

Single-Cell Genomics Reveals Hundreds of Coexisting Subpopulations in Wild *Prochlorococcus*

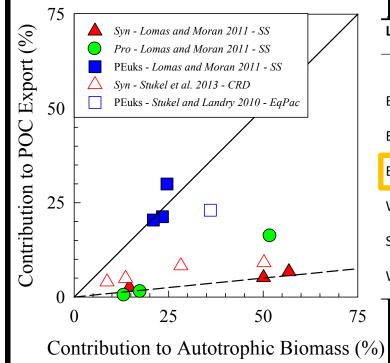
Nadav Kashtan,¹* Sara E. Roggensack,¹ Sébastien Rodrigue,^{1,2} Jessie W. Thompson,¹ Steven J. Biller,¹ Allison Coe,¹ Huiming Ding,^{1,3} Pekka Marttinen,⁴ Rex R. Malmstrom,⁵ Roman Stocker,¹ Michael J. Follows,⁶ Ramunas Stepanauskas,⁷ Sallie W. Chisholm^{1,3}*







Martiny et al. 2013 Lomas et al. unpubl. data



Location	Export Flux (mol C m ⁻² yr ⁻¹)		
	Ref	C:P=110	VCP
E. Subarctic N. Pac. (OSP) 50°N, 145°W	2.3±0.6	3.7±0.3	2.8±0.2
E. Subtropical N. Pac. (HOT) 23°N, 158°W	2.5±0.7	1.3±0.6	2.4±0.9
E. Equatorial Pac. 170°-95°W	3.3±1.8	3.0±0.5	2.7±0.4
W. Equatorial Pac. 150°E-140°W	1.5	2.5±0.7	2.0±0.5
Subarctic N. Atl. 40°N-65°N, 10°W-60°W	2.8±2.7	5.9±1.1	3.5±1.1
W. Subtropical N. Atl. (BATS) 32°N, 64°W	3.8±1.2	1.2±0.5	3.8±1.5

'Ref' from Emerson 2014 Teng et al. 2014, ASLO abstract

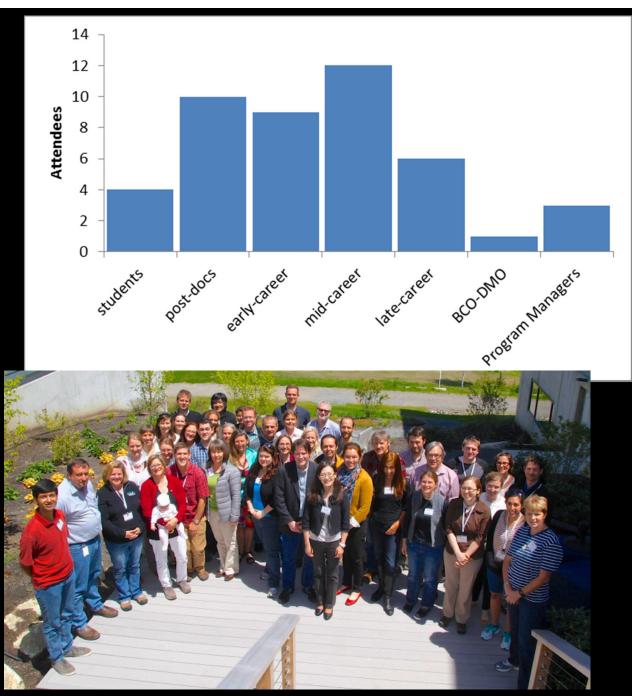


To understand cell- and taxon-specific physiological responses of marine microbes and phytoplankton in the current and a changing environment;

To understand the interactions and feedbacks between plankton physiological plasticity, genetic potential and evolution and the impacts on global ocean biogeochemical cycles



Attendees:





Questions we tried to address:

What are the single cell-specific and population-level traits required for parameterizing microbial "plasticity" in marine biogeochemical models of the current ocean?

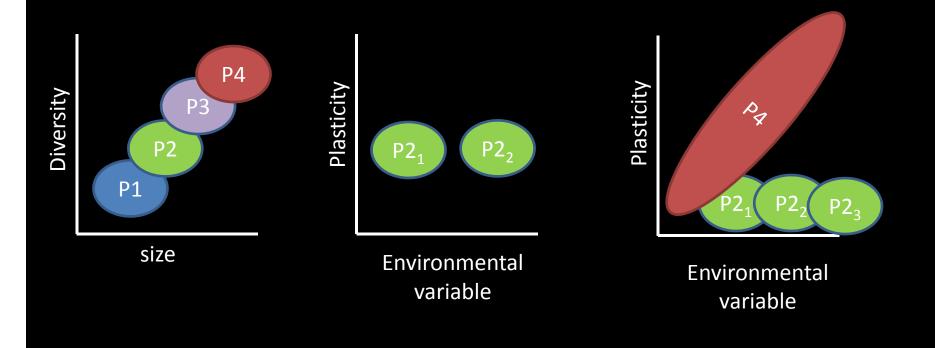
Do genetic diversity, evolution and physiological plasticity have similar or different impacts on ocean biogeochemistry, particularly the production and export of particulate organic matter from the surface ocean?

What roles do taxonomic diversity and physiological plasticity play in controlling the response of microbial communities to current and future environmental stressors, for example, oxygen minimum zones, ocean acidification, ocean warming, stratification and changing nutrient concentrations or supply rates?



Preliminary summary of outcomes:

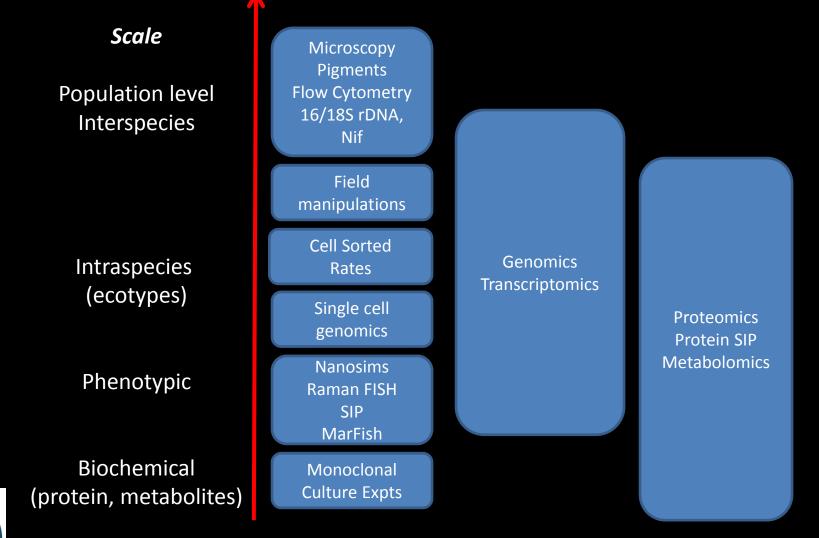
Types and scales of biodiversity





Preliminary summary of outcomes:

Linking biodiversity and measurements





Preliminary summary of outcomes:

Scale Population level Interspecies

Intraspecies (ecotype to SNPs)

Phenotypic

Biochemical (protein, metabolites)

Production e.g., through trait selection of highest u? Grazing, e.g., through cell size changes, encounter probability EXPORT e.g., through stoichiometric variability of exported cells, DNA fingerprints

Need methods improvements to connect plasticity with flux !



Potential products:

- 1) A formal workshop report to the OCB Office in the form of a white paper that communicates the vision and forward-looking priorities for this research.
- **2)** An article for EOS highlighting the workshop conclusions and vision at a high level.
- **3)** A detailed review publication in *L&O: Review, Frontiers in Aquatic Microbiology,* or similar journal.
- 4) Identification of key topics and possible speakers for a special session at the 2015 Aquatic Sciences meeting, potentially to include a town hall meeting to discuss the outcomes of the meeting and future research avenues.



For more information:

http://www.whoi.edu/website/taxon-specific-biogeochemistry/

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