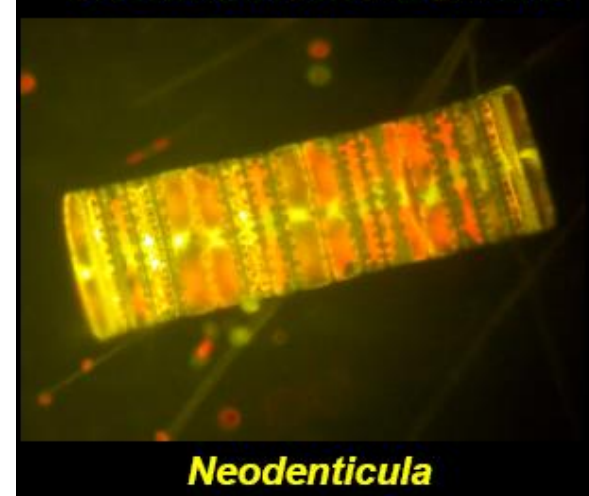
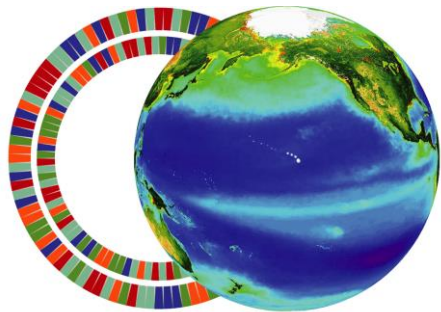


Planktonic community structure and the biological pump



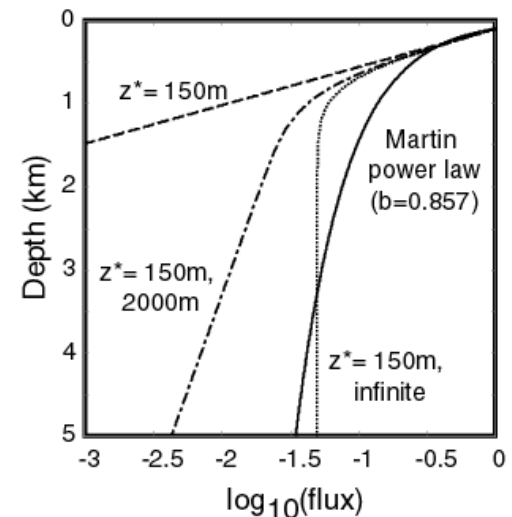
Philip Boyd

Institute for Marine & Antarctic Studies



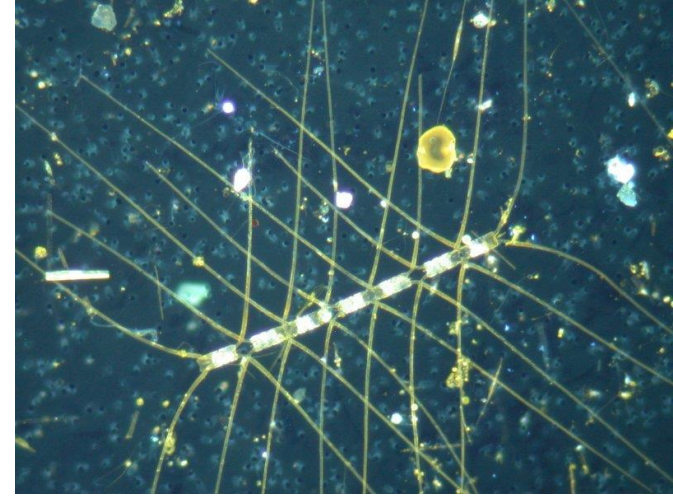
cmare

2014



OUTLINE

Multiple roles of the biological pump
What goes up must come down?



Particle transformations and foodwebs

Surface ocean

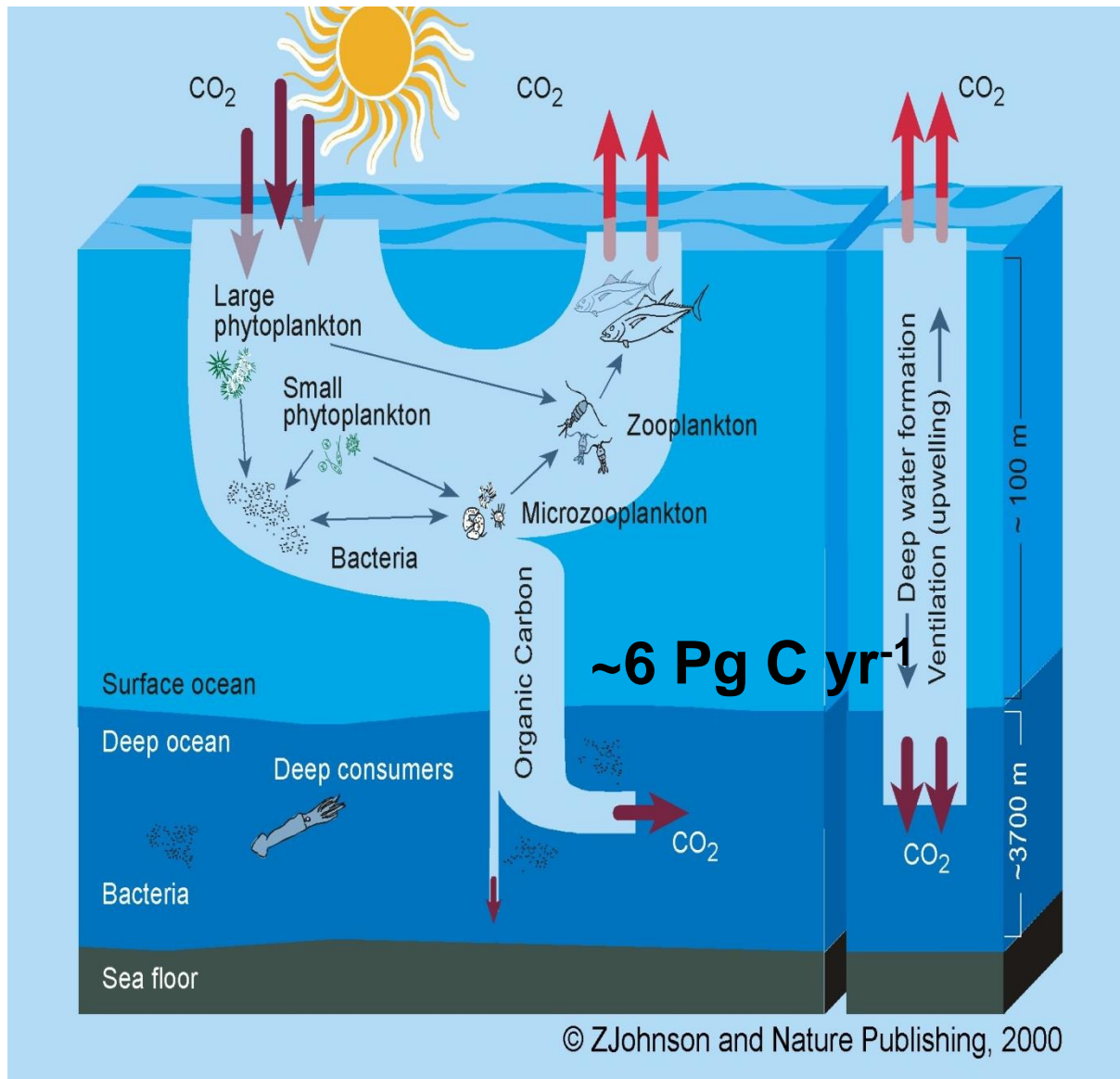
Subsurface ocean – zooplankton
- microbes

Particle attenuation – surface vs. subsurface waters

Biological pump – at a cellular level

Conclusions

Multiple roles of the biological pump



Carbon
export

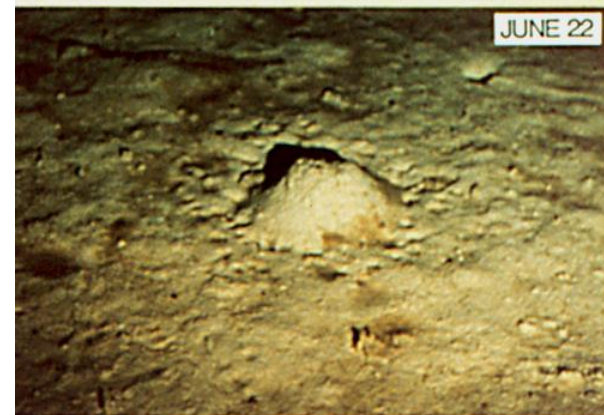
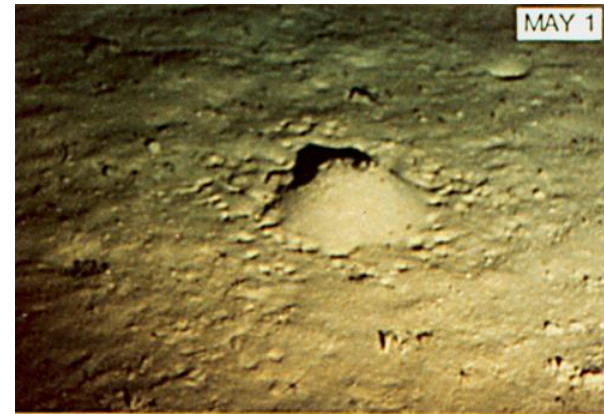
3.4 to 4.7×10^9 tons
C yr⁻¹ (Eppley & Peterson)
New production

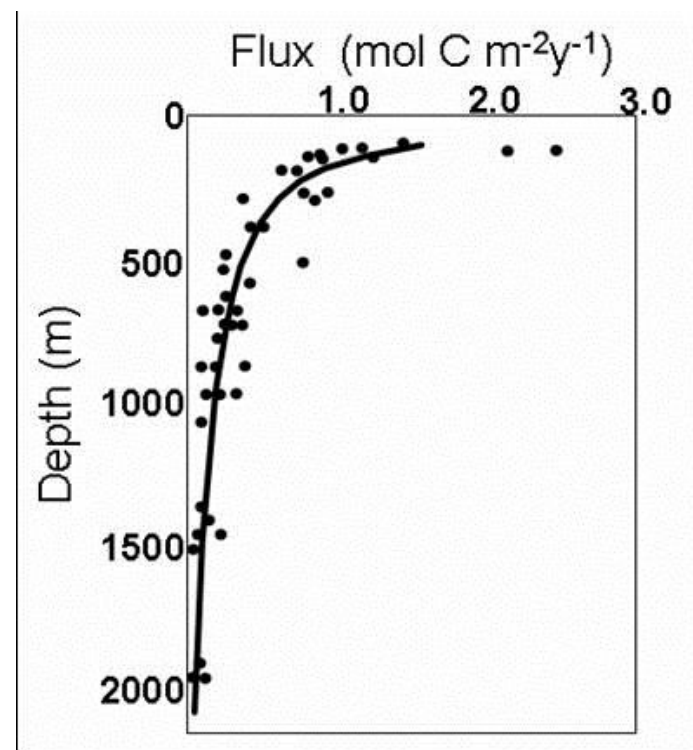
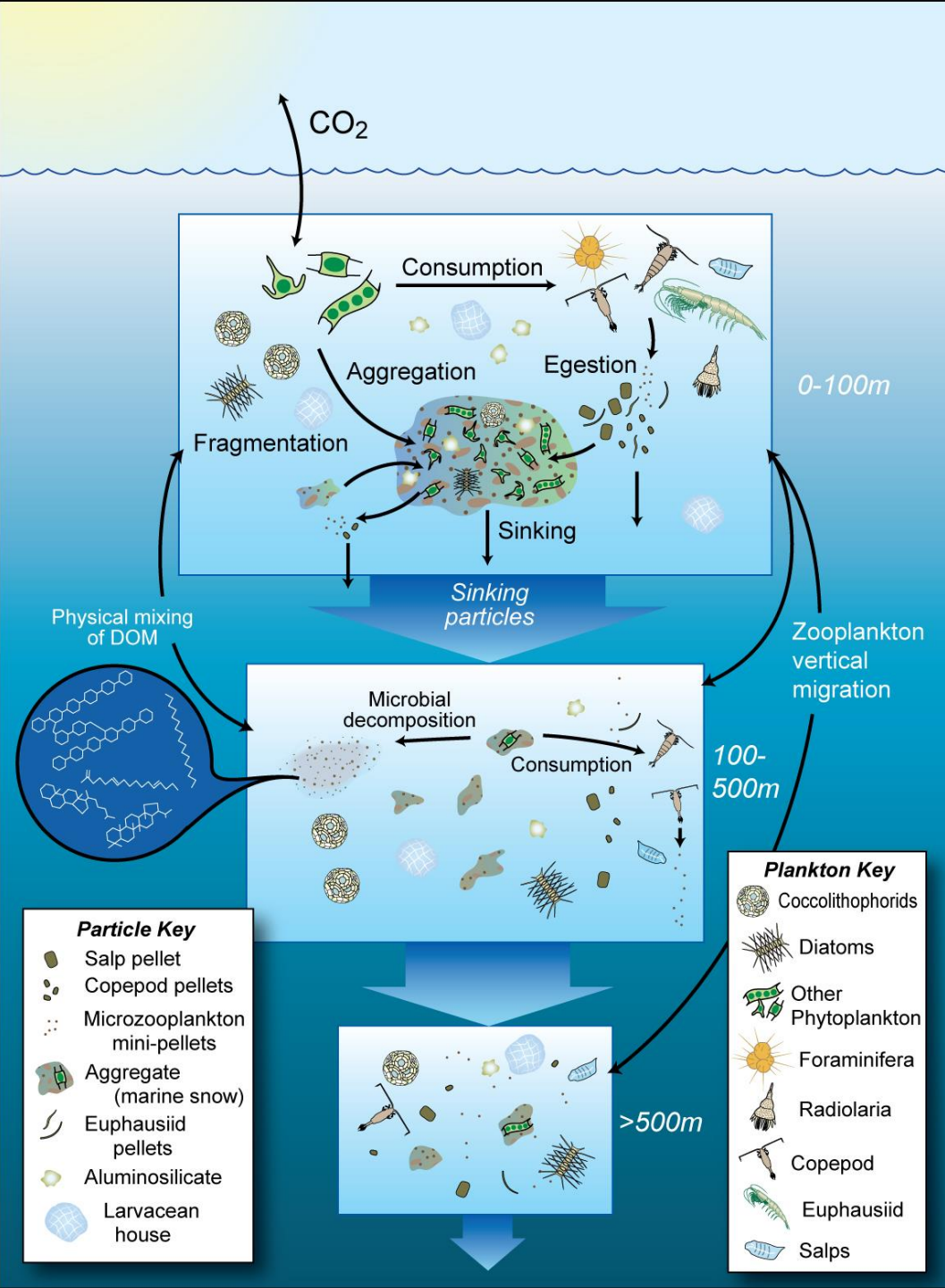
Multiple roles of the biological pump

Nutrition of the deep ocean & benthos

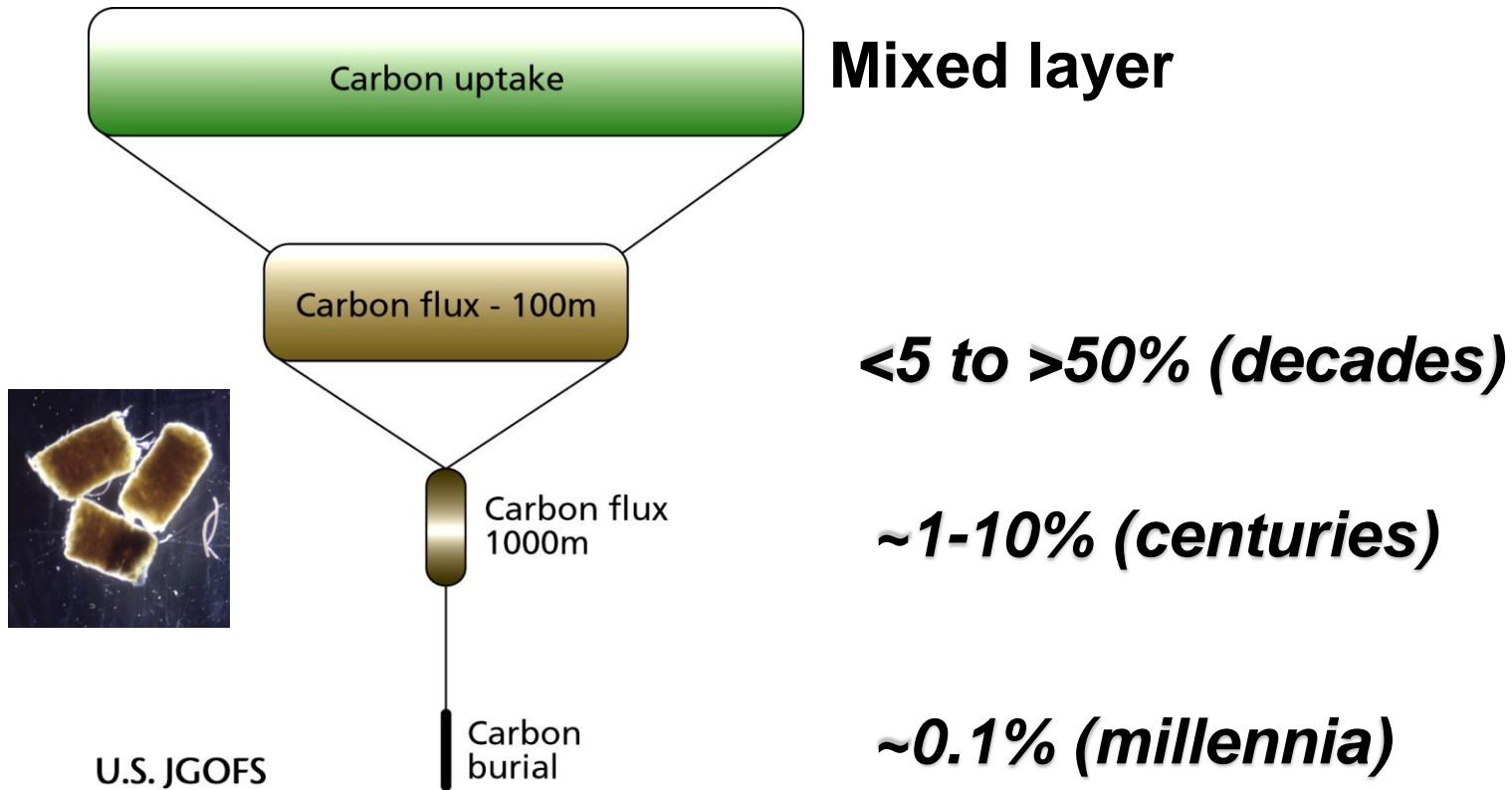
"...deep-sea organisms are nourished by a "rain" of organic detritus from overlying surface waters."

Alexander Agassiz (1888)





Sinking flux - vertical attenuation of particles



The rain of sinking particles scavenges dissolved materials

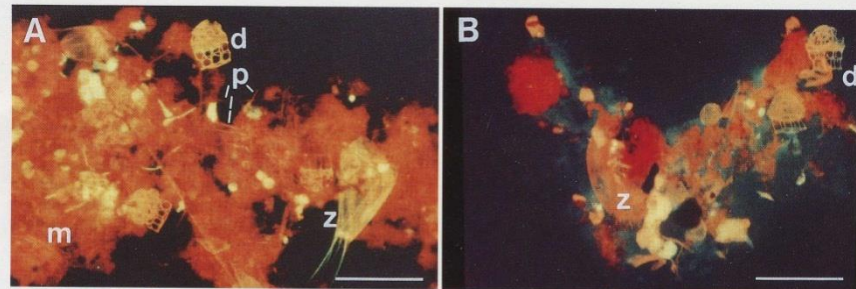


Fe

Fe

Fe

Nutrient regeneration



N



C

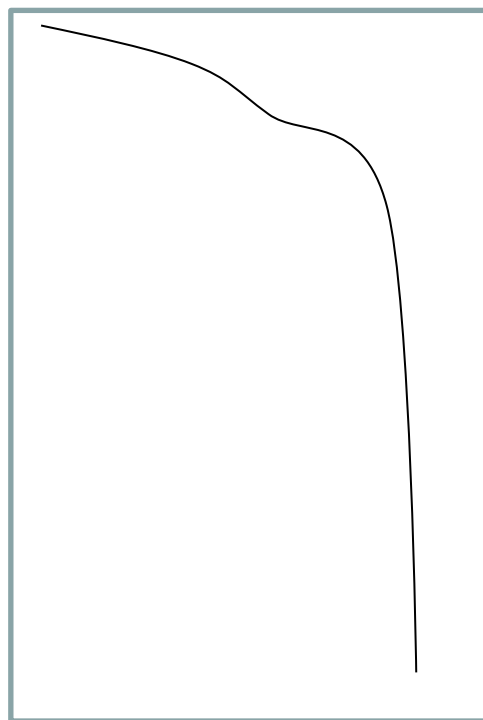


Si

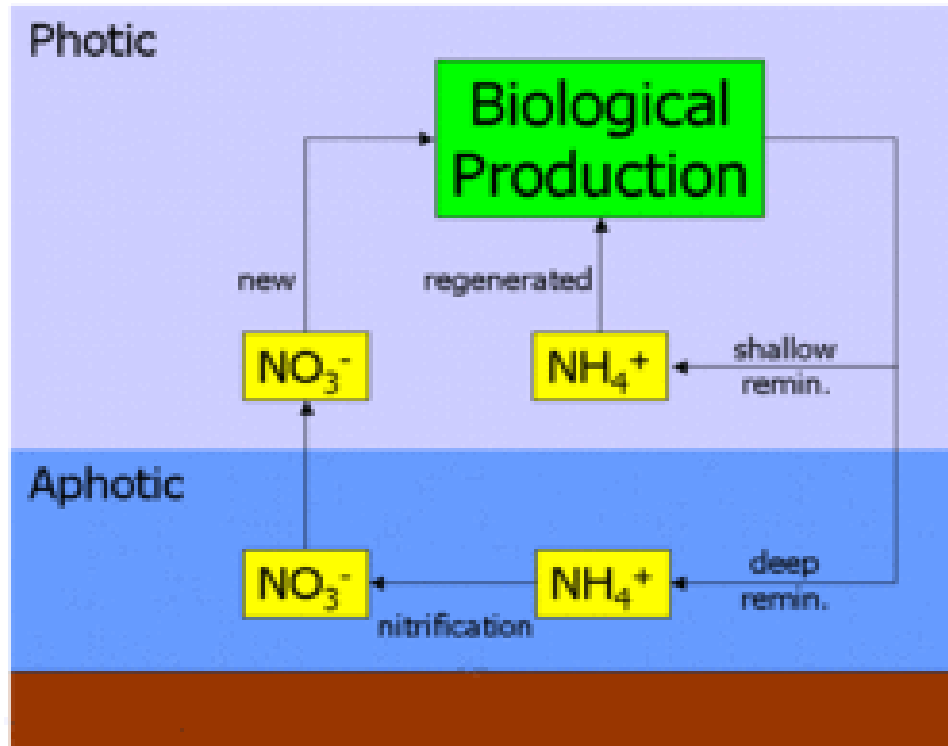


Fe

Remineralization length scales Of major & minor elements

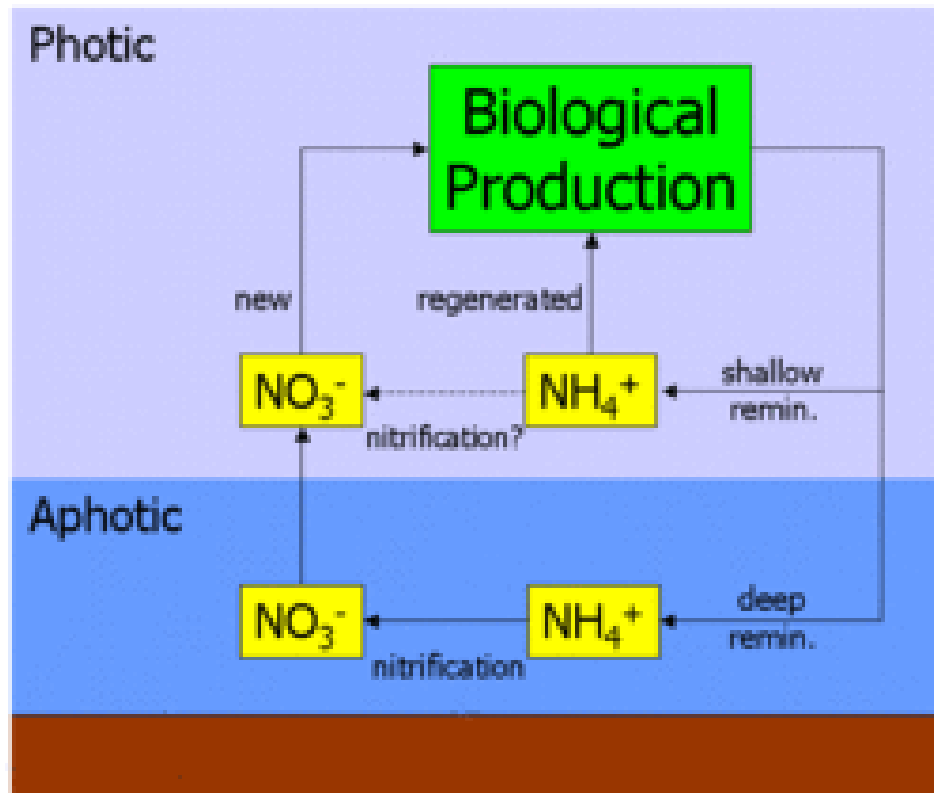


What goes up must come down?

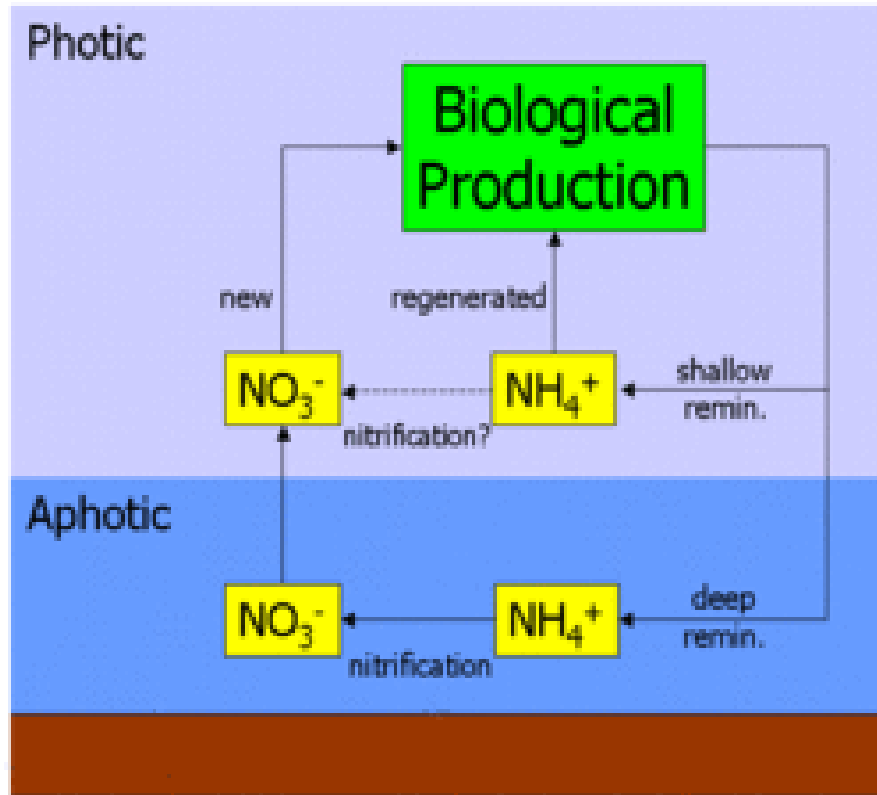


Particulate organic matter flux and planktonic new production in the deep ocean
Richard W. Eppley & Bruce J. Peterson

“Global new production is of the order of $3.4\text{--}4.7 \times 10^9$ tons of carbon per year and approximates the sinking flux of particulate organic matter to the deep ocean.”

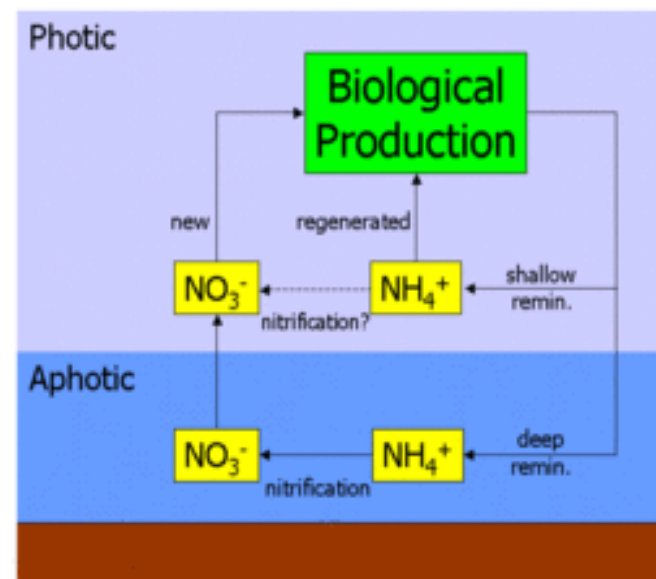


"To relate new production to export requires that nitrification in the euphotic zone be negligible".



Yool et al. predict that up to 50% of surface nitrate is supplied by surface nitrification rather than vertically.

Yool et al. (2007).
"The significance of nitrification for oceanic new production".



So what metric(s) should we use to compare surface mixed layer productivity to downward particulate export?

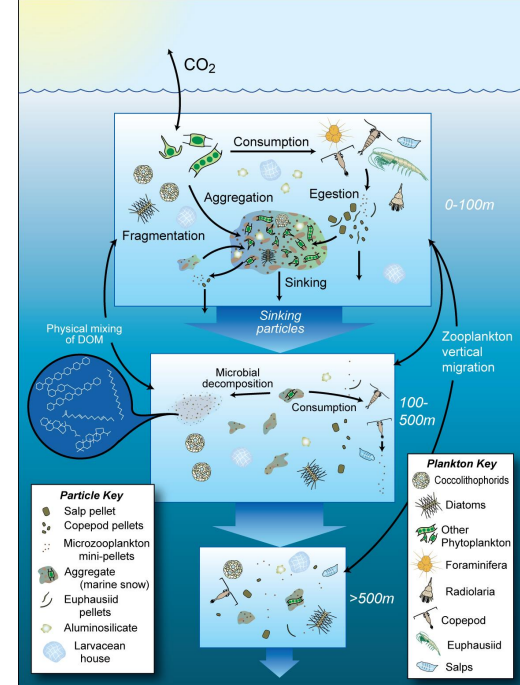
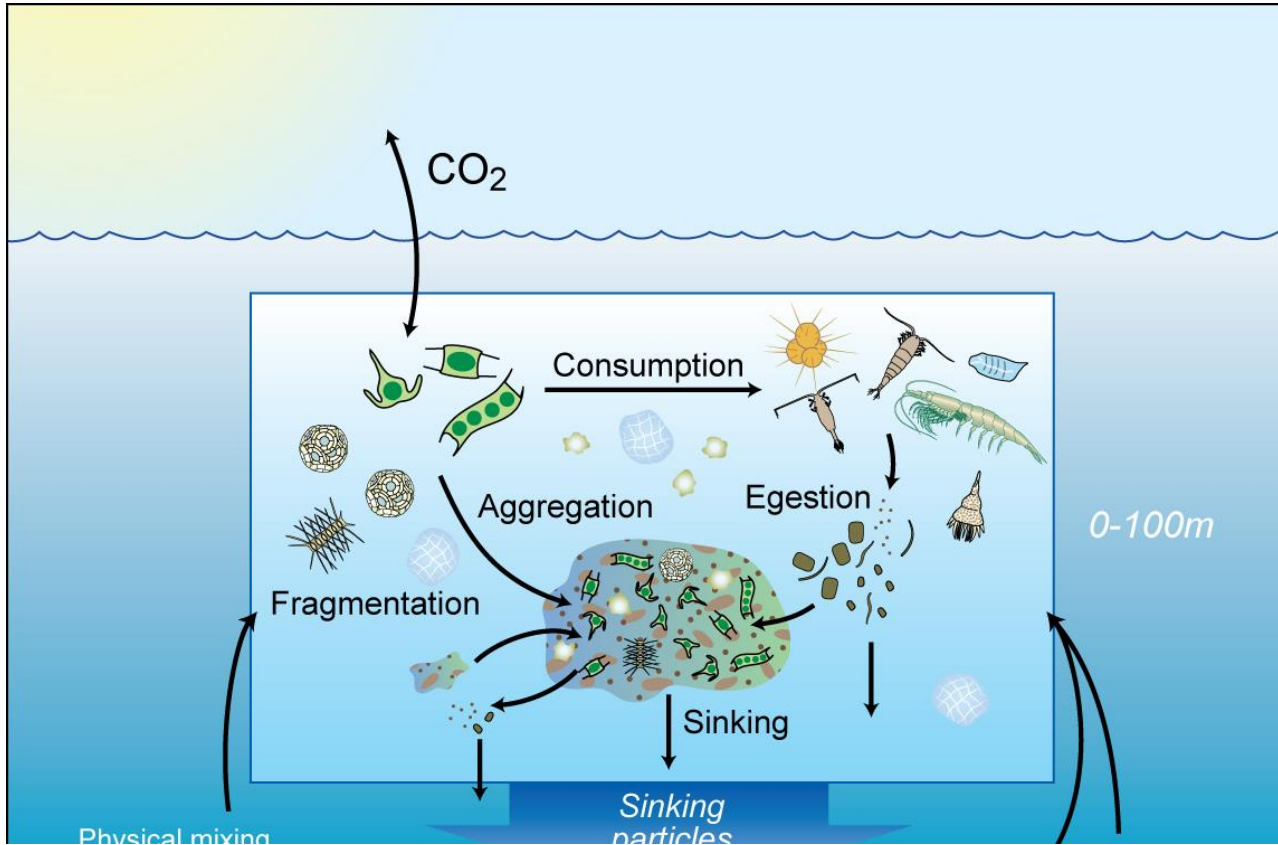
Back to NPP – the pe ratio of Dunne et al. (2005) GBC?

Clark DR et al. 2008. Ammonium regeneration and nitrification rates in the oligotrophic Atlantic Ocean: Implications for new production estimates. *Limnology and Oceanography*, 53(1), 52-62.

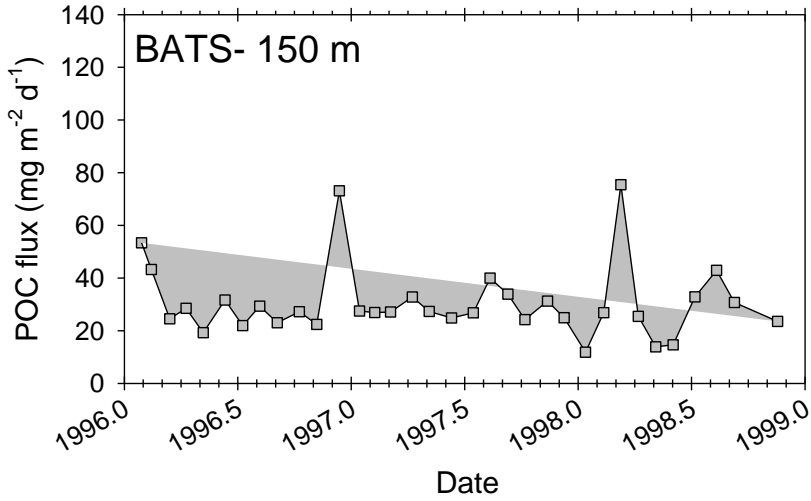
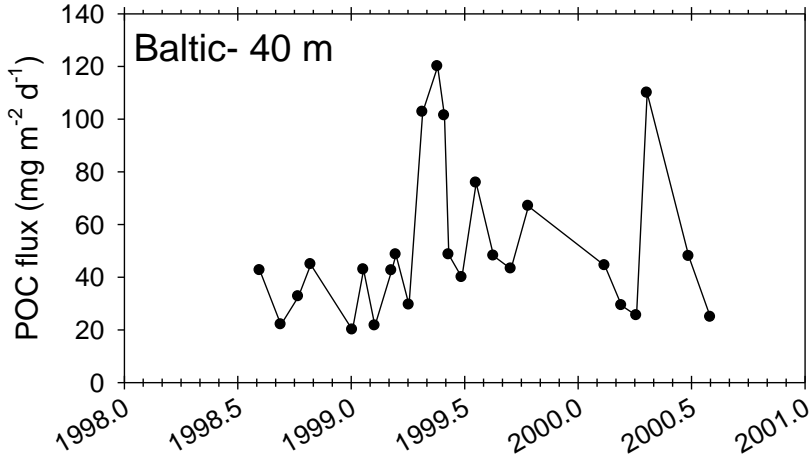
Yool, Andrew (2011) [Modeling the Role of Nitrification in Open Ocean Productivity and the Nitrogen Cycle.](#)

Particle transformations and foodwebs

Surface ocean

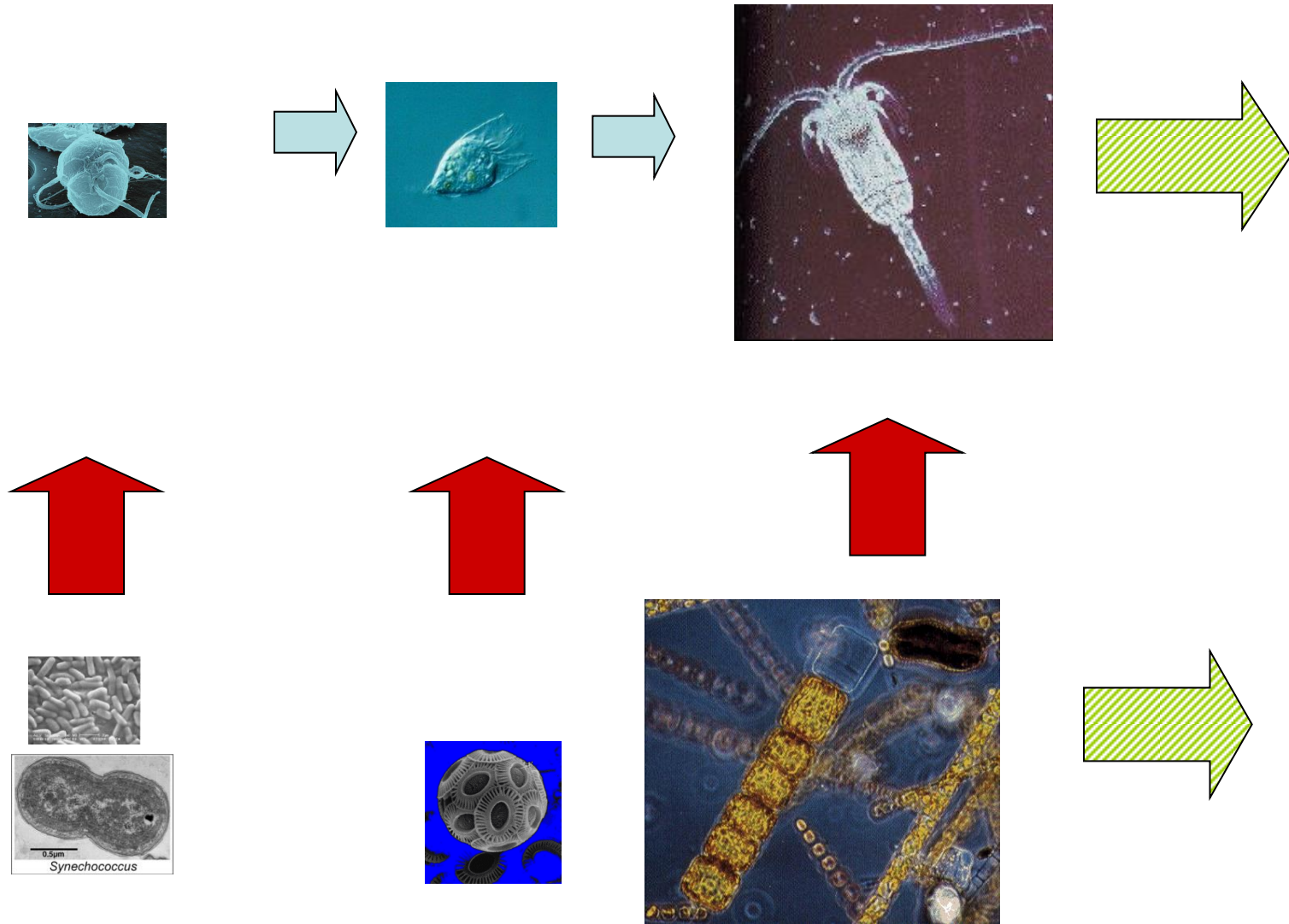


What is the imprint of the biological pump Just below the surface mixed layer?

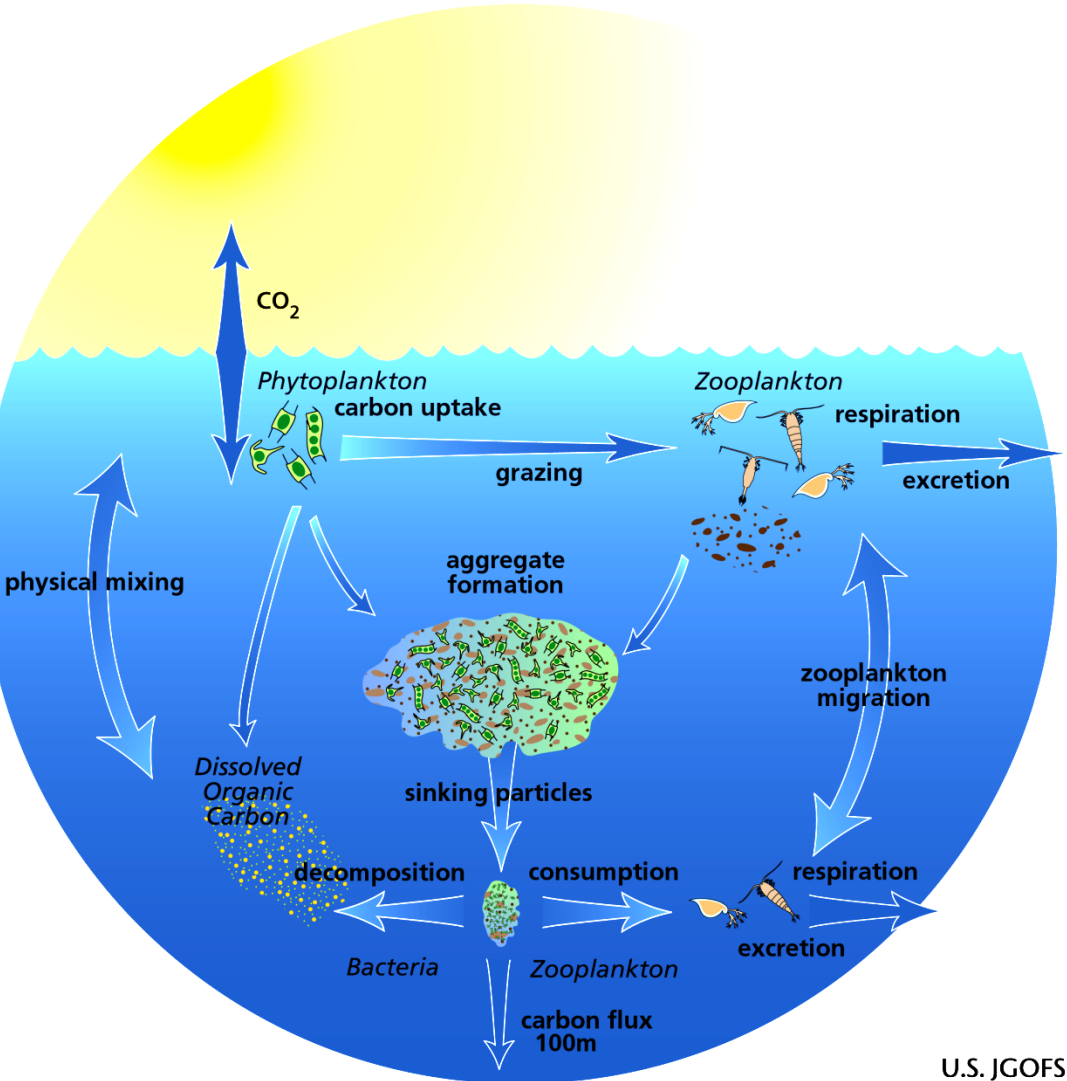
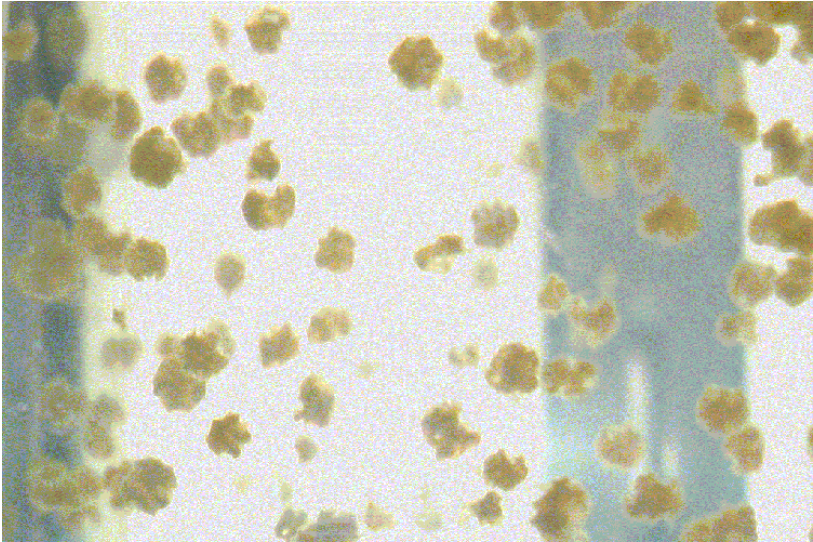
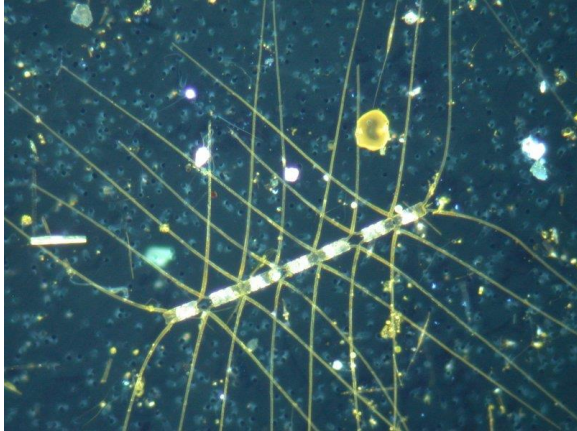


From Asper 1997

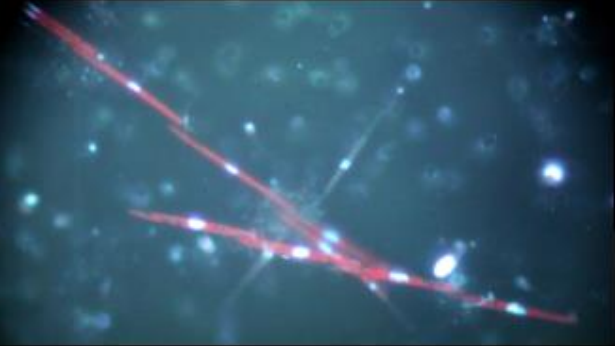
The foodweb – controlling particle transformations



DIRECT ALGAL SINKING



>20 μm Phytoplankton - K2



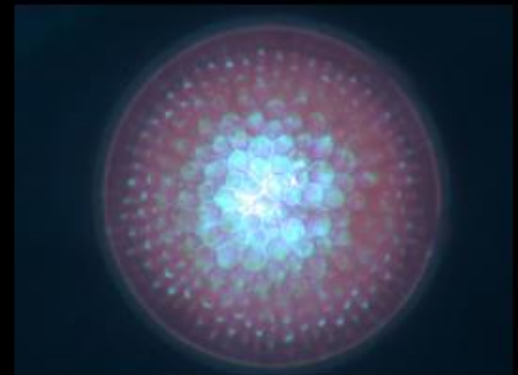
Pseudo-nitzschia



Dictyocha



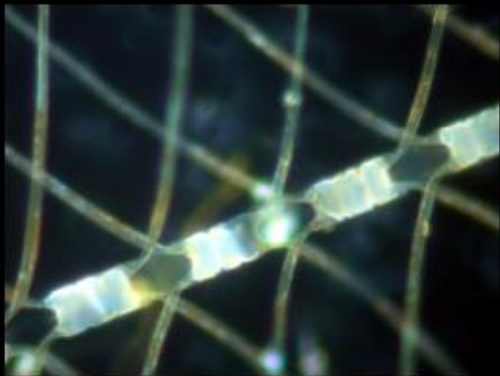
Rhizosolenia



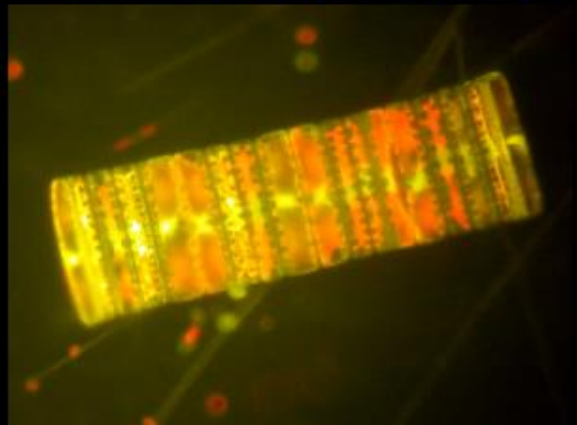
Coscinodiscus marginatus



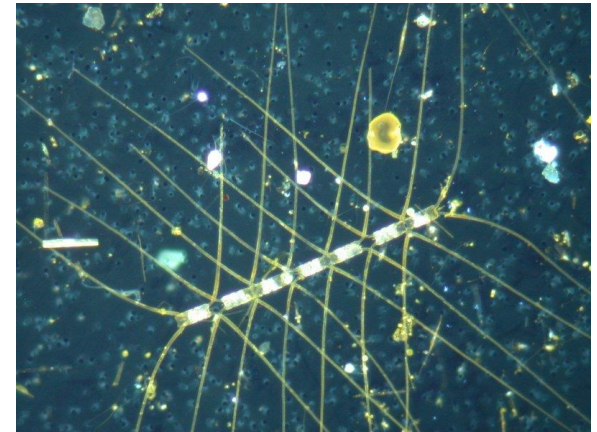
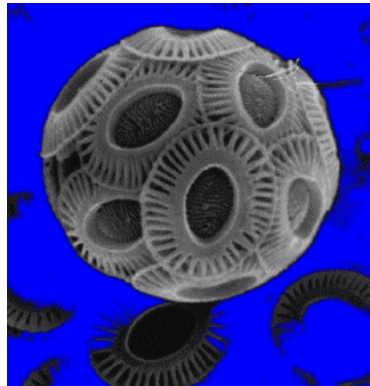
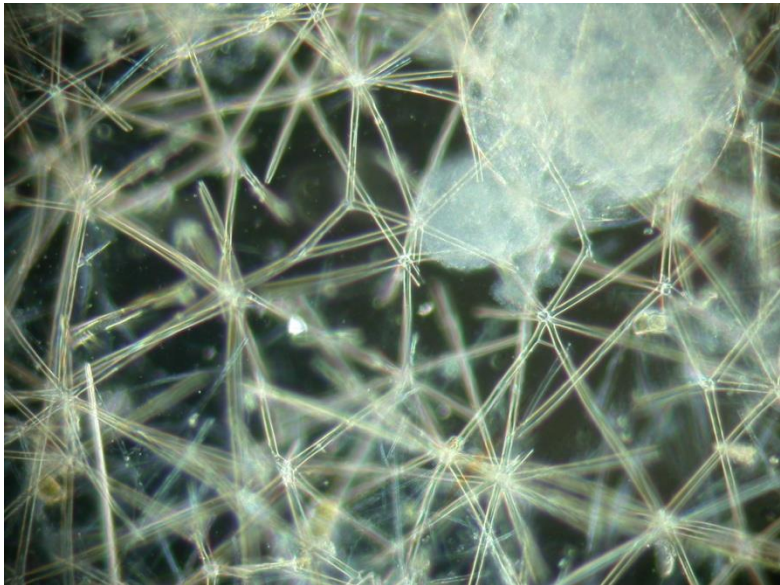
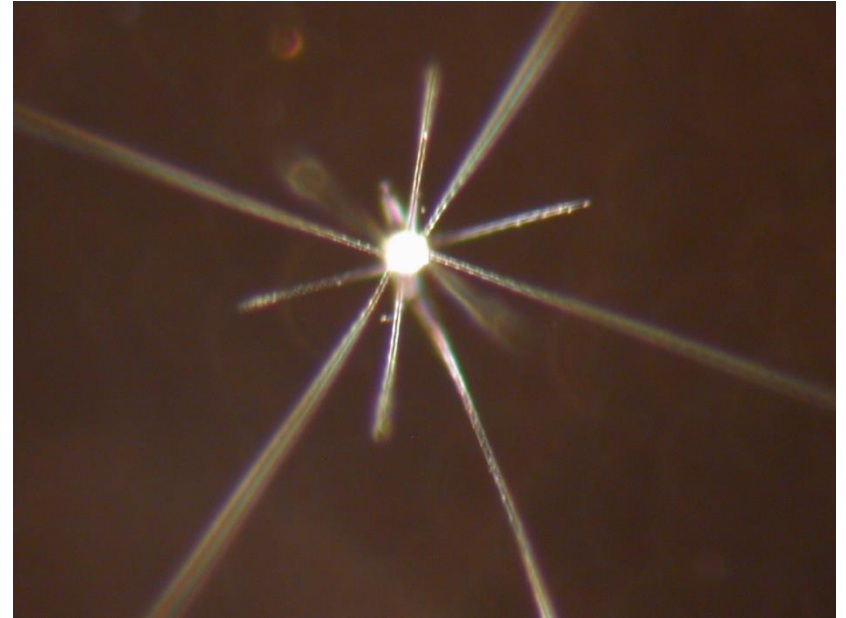
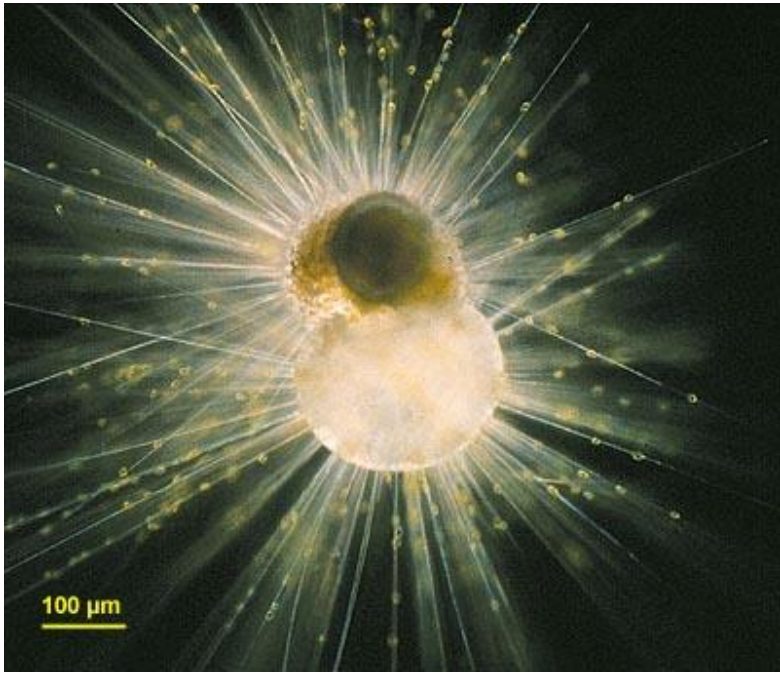
Corethron



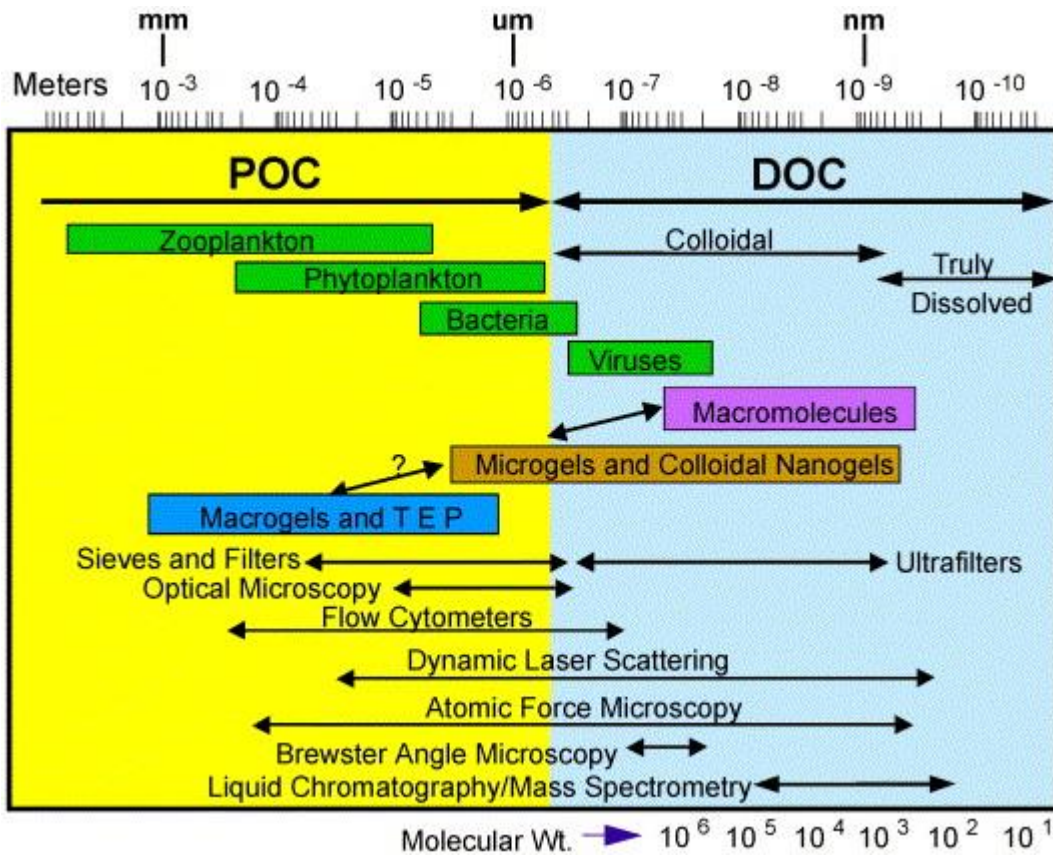
Chaetoceros



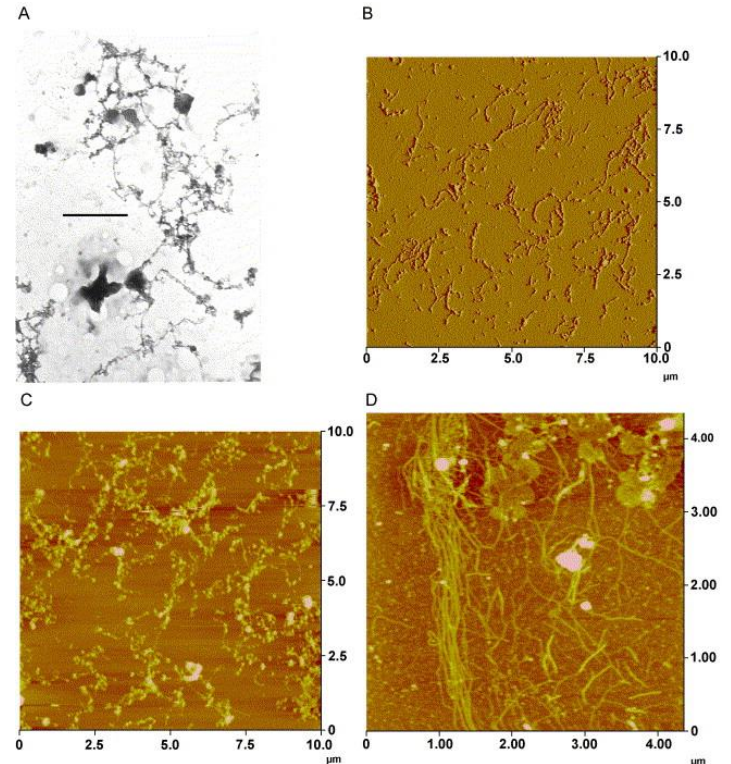
Neodenticula



Agents of mineral ballast – opal and calcite



Gels and Biological glues

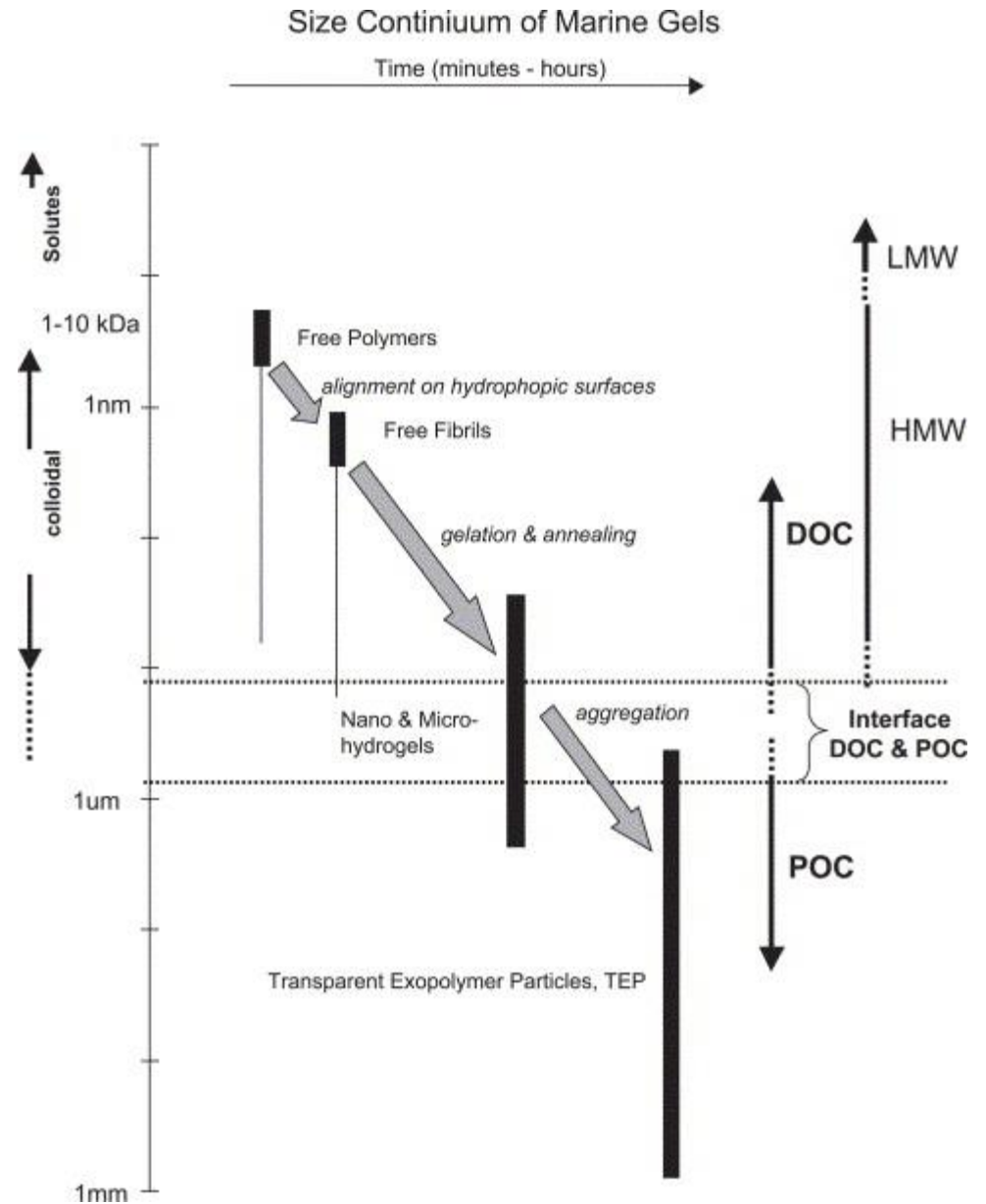


VERDUGO et al, (2004)

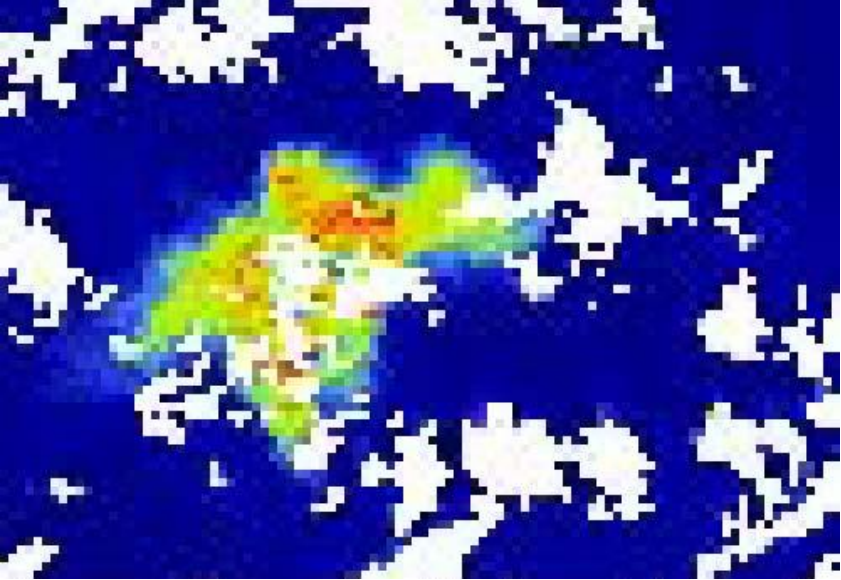
Exudation



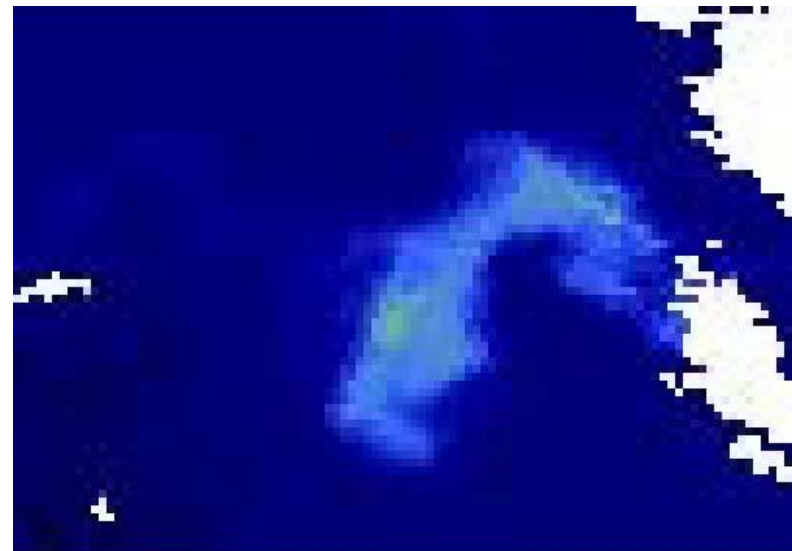
TEP – Passow and Alldredge



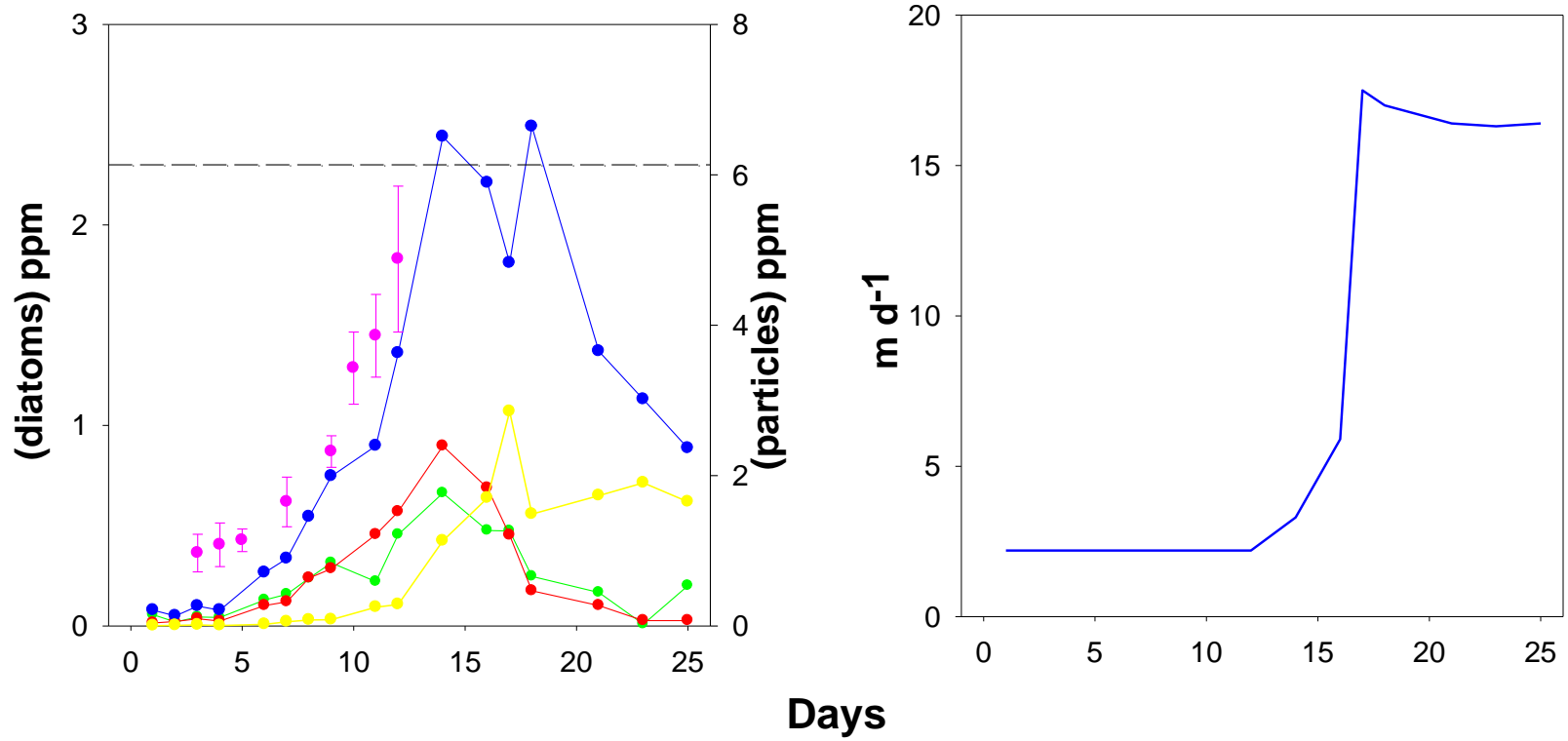
VERDUGO et al, (2004)



**A case study – the SERIES mesoscale in situ Fe enrichment
What halts bloom development and initiates bloom decline?**



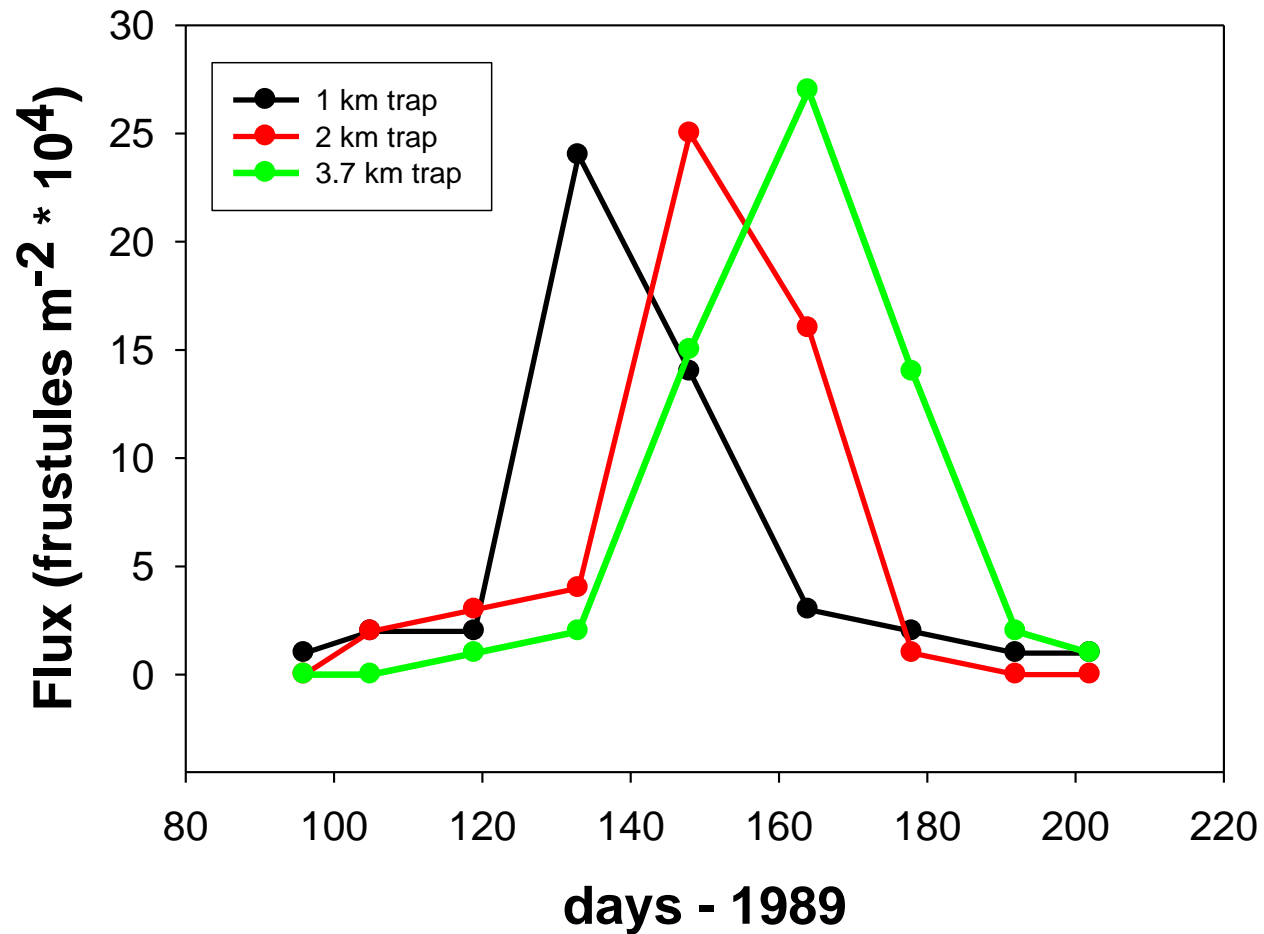
Boyd et al. (2005)



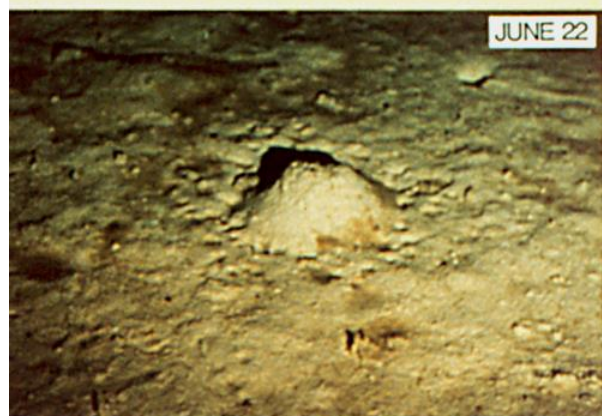
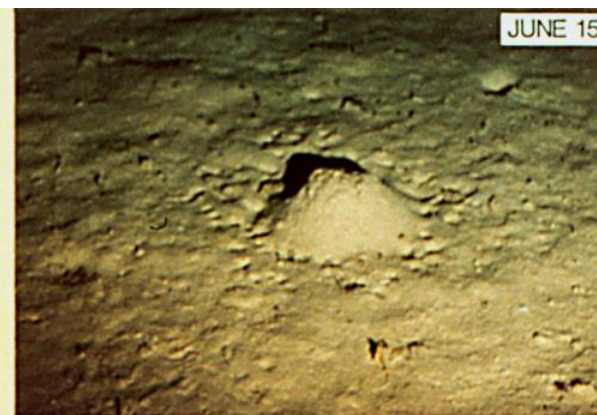
Boyd et al. (2005)

TRACKING THE ALGAL AGGREGATION SIGNATURE TO DEPTH
Honjo and Manganini (1993) NABE site

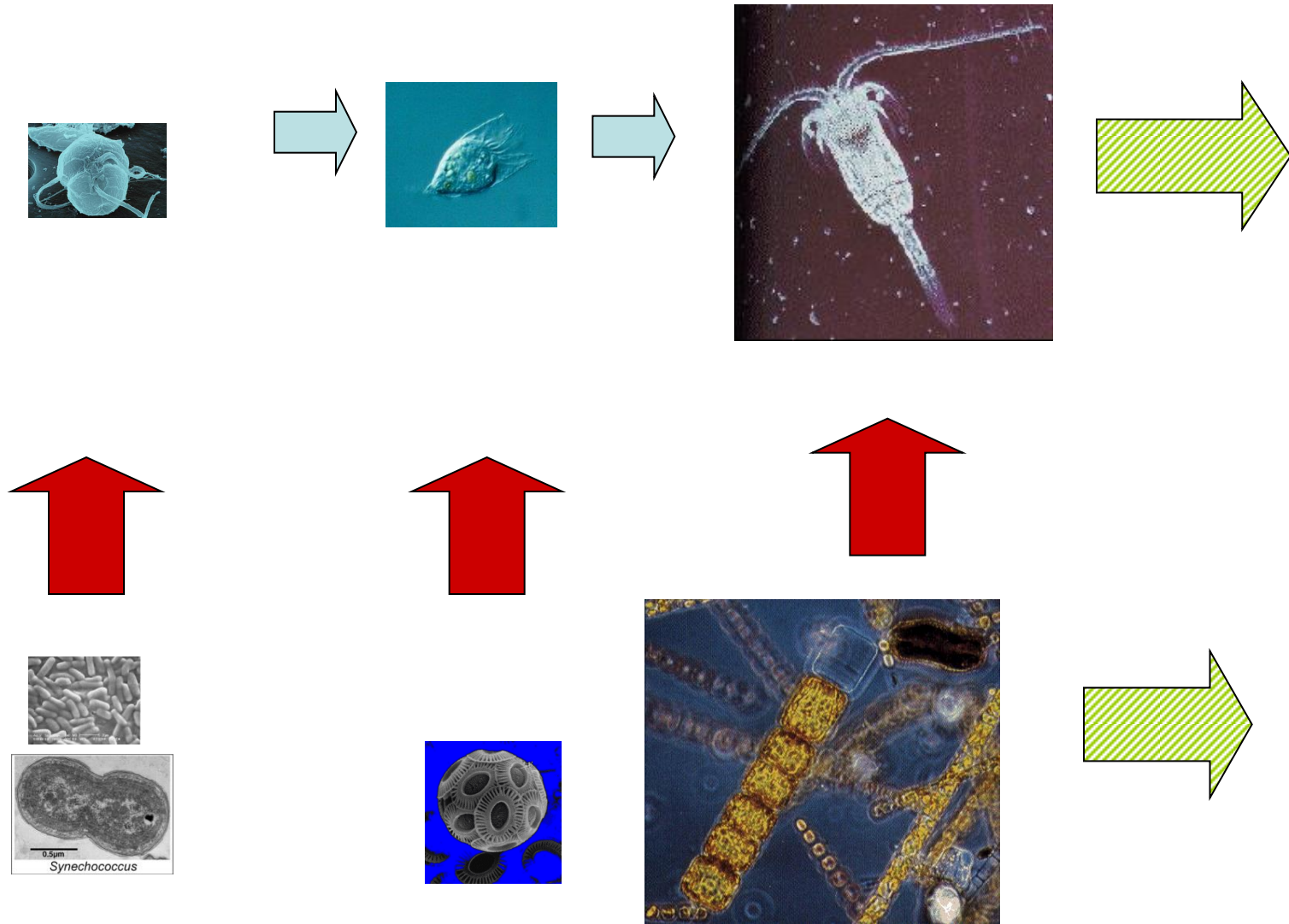
Chaetoceros messanensis



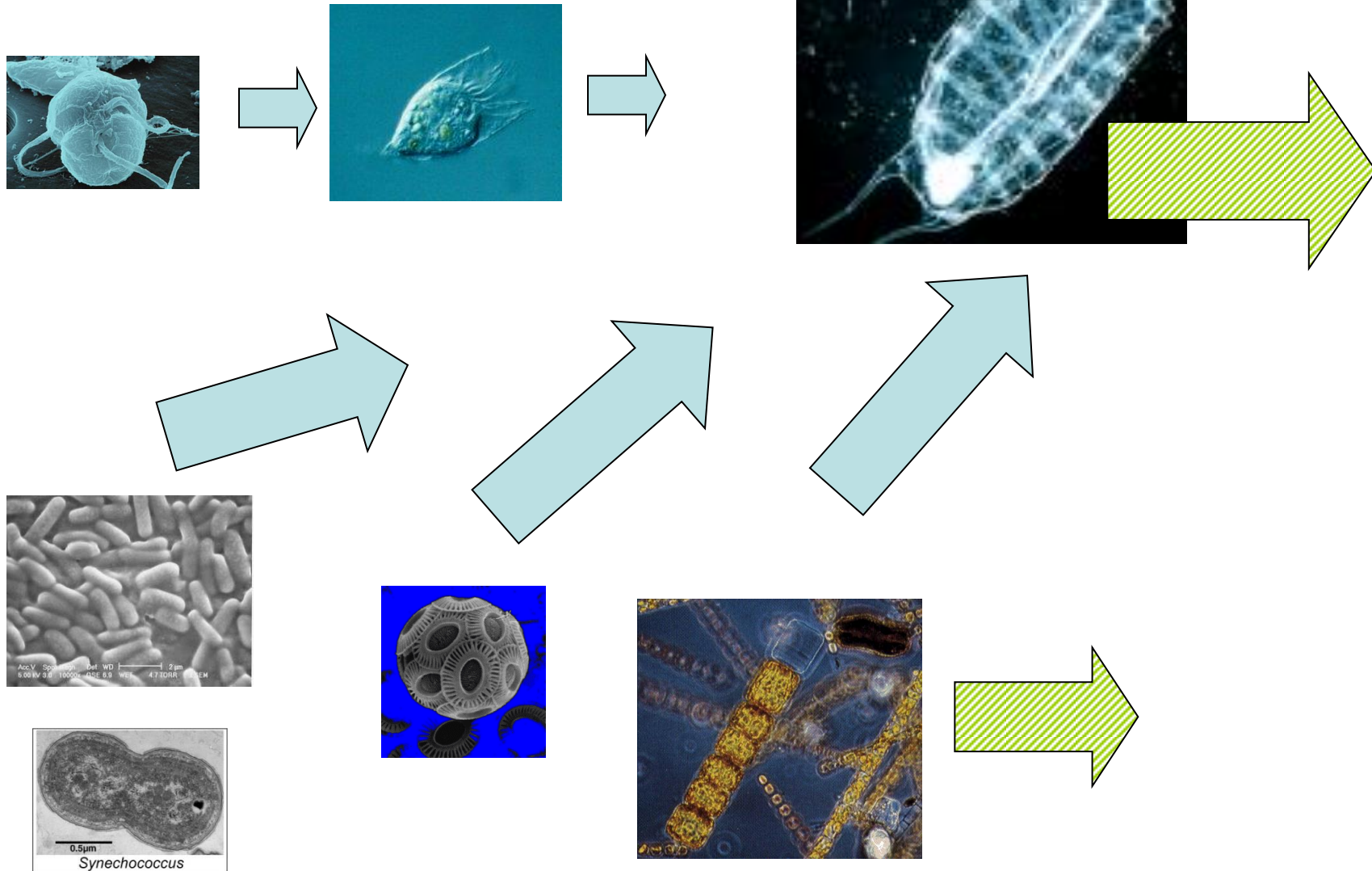
Impact of direct algal sinking



The foodweb – controlling particle transformations



Different foodweb structures result in A range of export efficiencies (pe ratio)



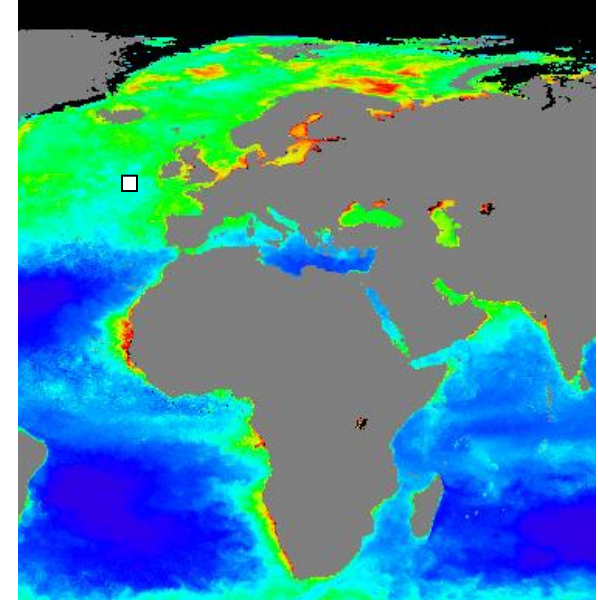
pe ratio = particle export/NPP



SALPS CRASH BLOOM

***Biomass of Salpa fusiformis
of up to 360 mg C m⁻³***

***Herbivory during the diatom bloom ended
the bloom before nutrients had run out.***

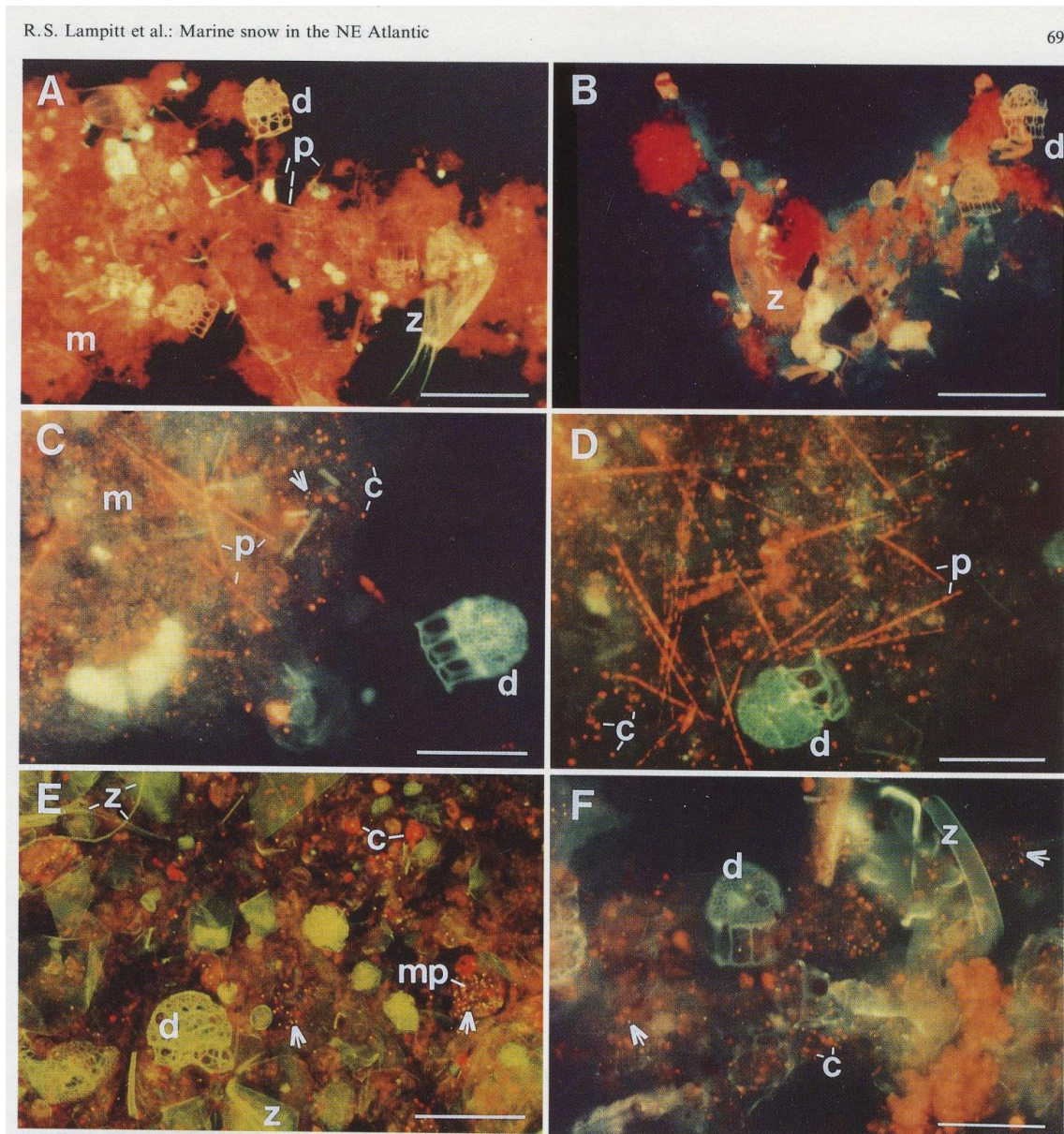


**Sedimentation of diatom-rich salp fecal
pellets
> 1 mm long, 350 μm wide, 10 μg C per
pellet**

BATHMANN (1988)

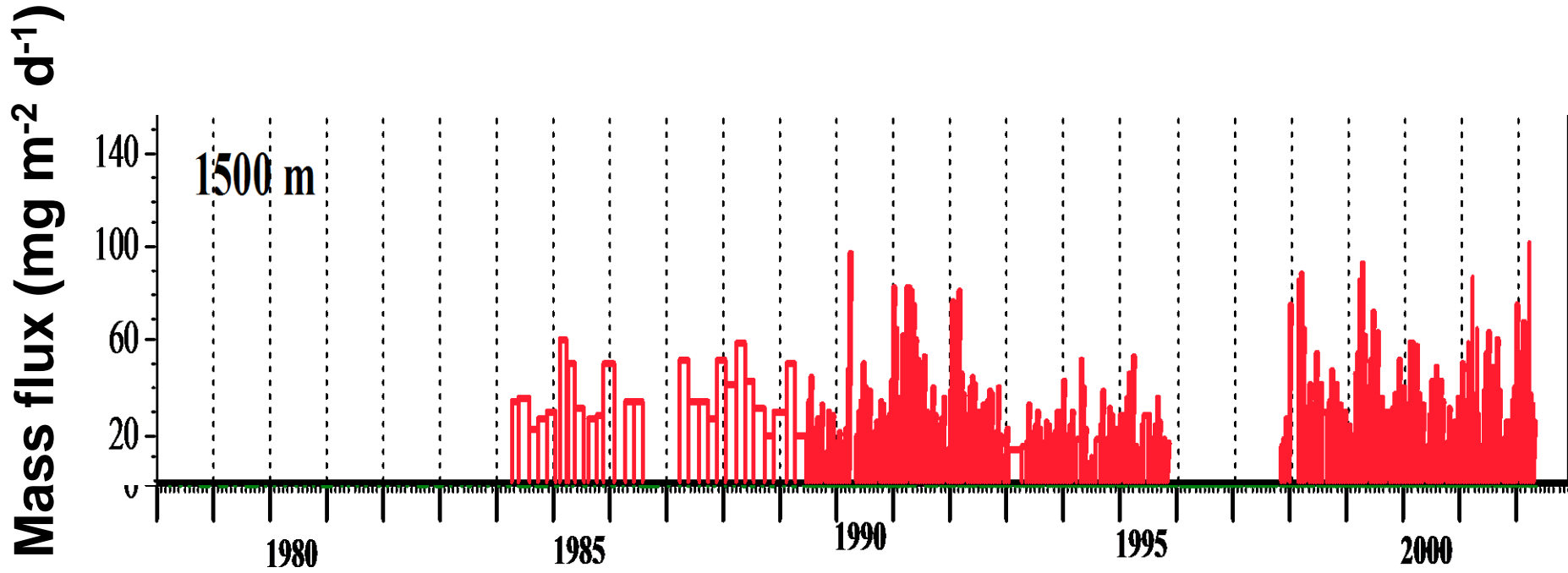
Particle transformations and foodwebs

Subsurface ocean – zooplankton & microbes



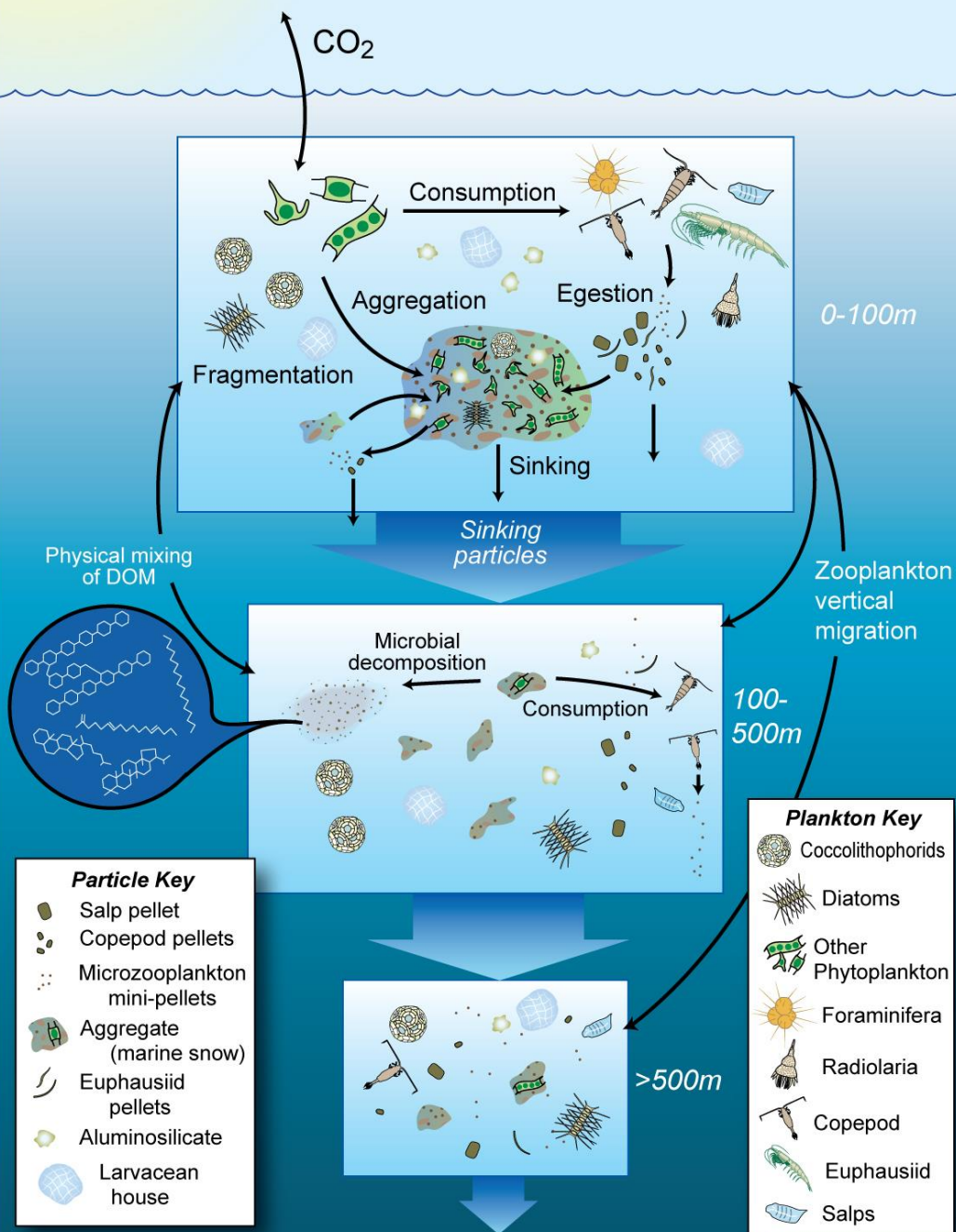
Images
Richard Lampitt
NOC UK

Seasonality in particle export is still evident in deep water



Courtesy Scott Nodder NIWA NZ

Factors controlling Export fluxes



Particle size

Particle composition

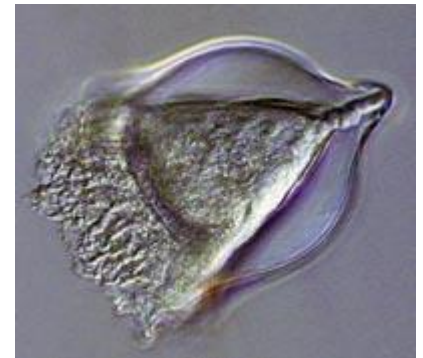
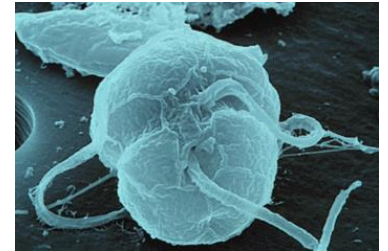
Particle geometry

Particle specific gravity

Particle sinking rate

Many of these processes are biologically-mediated

The grazers



Zooplankton and mesopelagic zone particle cycling



fixation of C,N
by phytoplankton

grazing

egestion

Base of euphotic

fragmentation

vertical migration

carnivory

respiration
excretion

detritivory

egestion

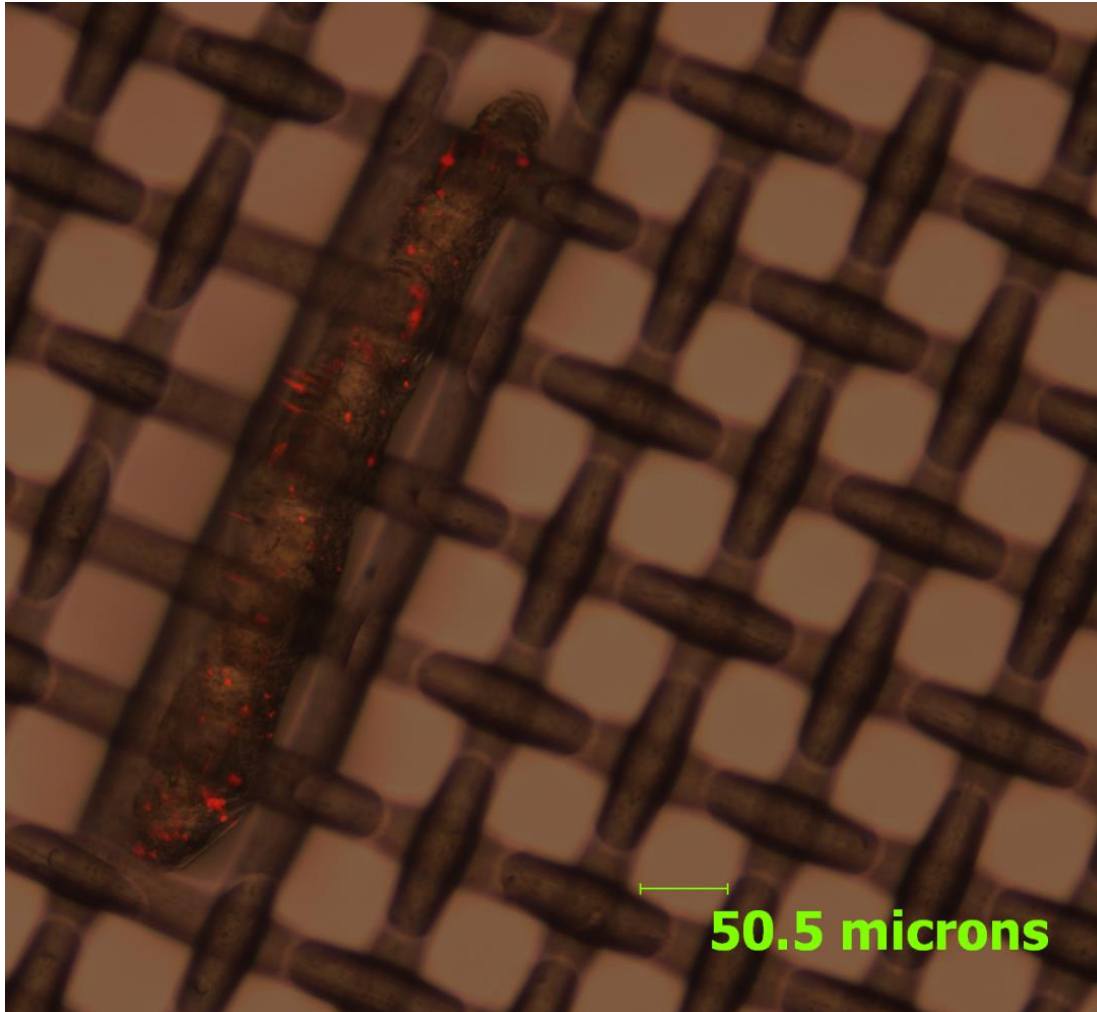
respiration
excretion

egestion

(death)

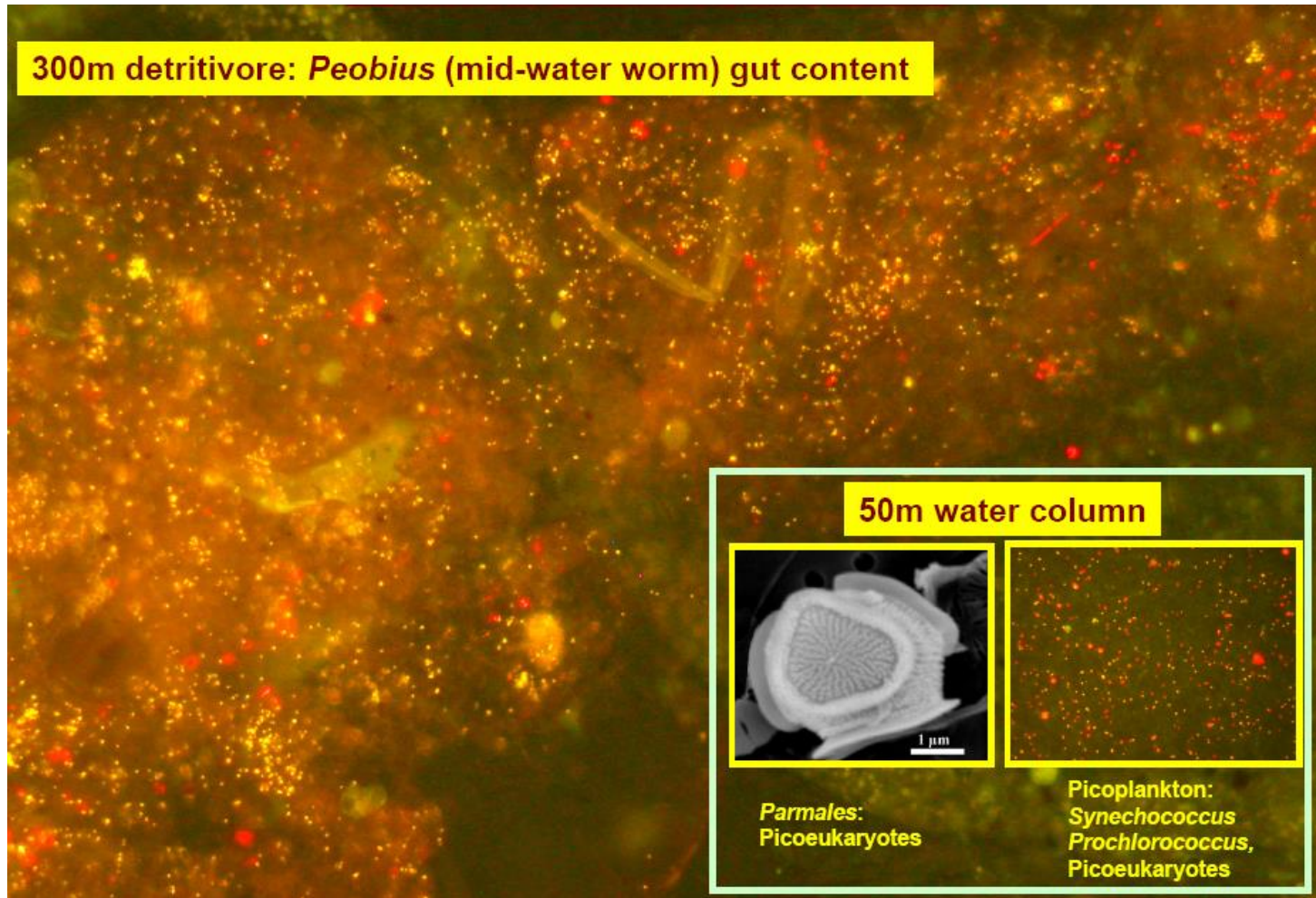


Particle repackaging by mesozooplankton

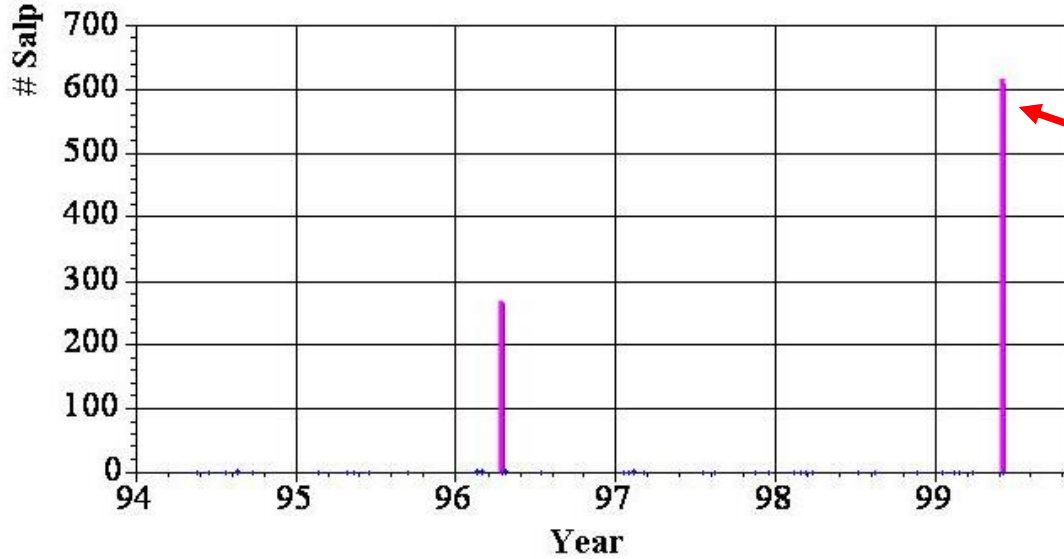
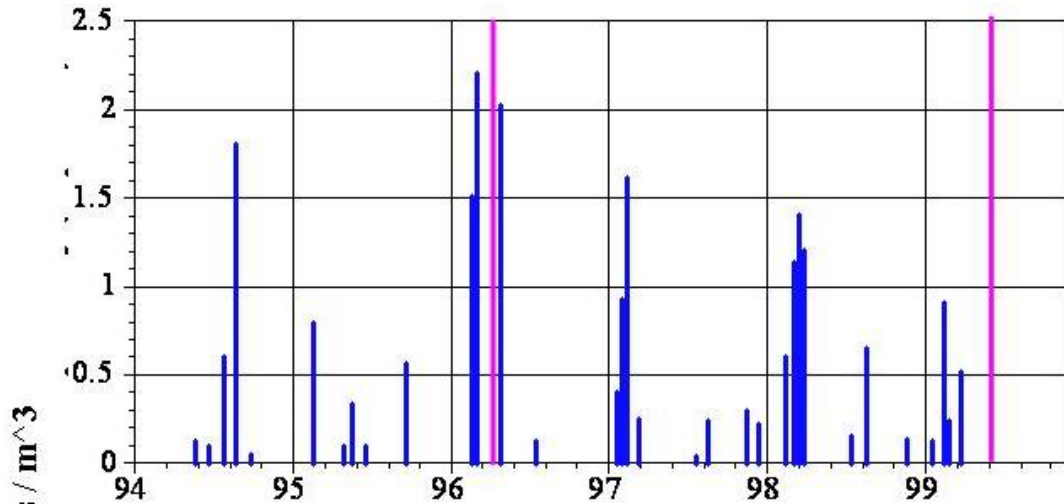


Boyd et al. (2012)

A mixed diet.....



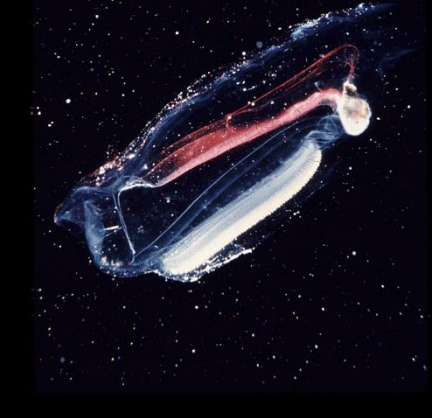
Salp blooms at BATS



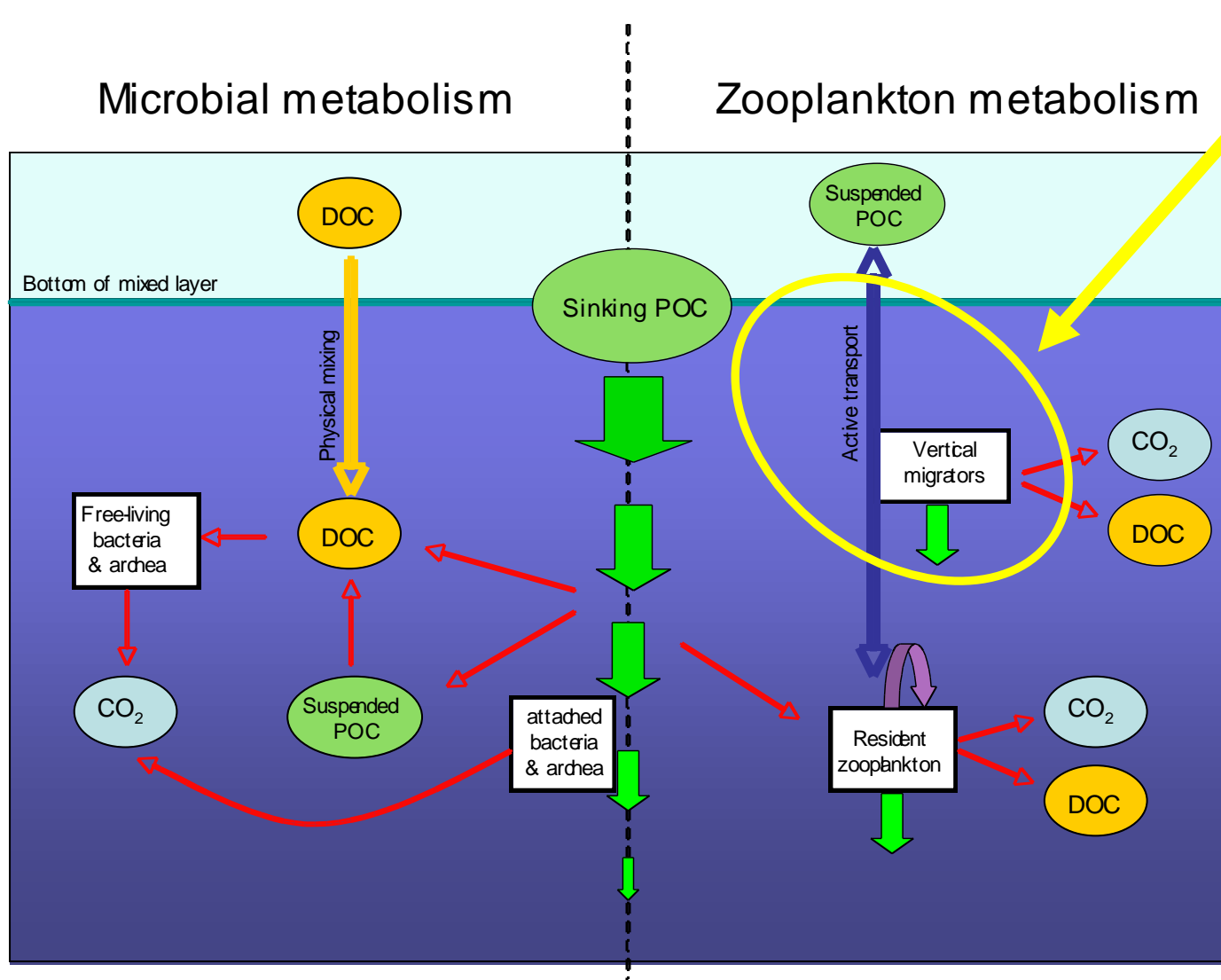
3200 m trap
Conte et al.



Steinberg & Madin



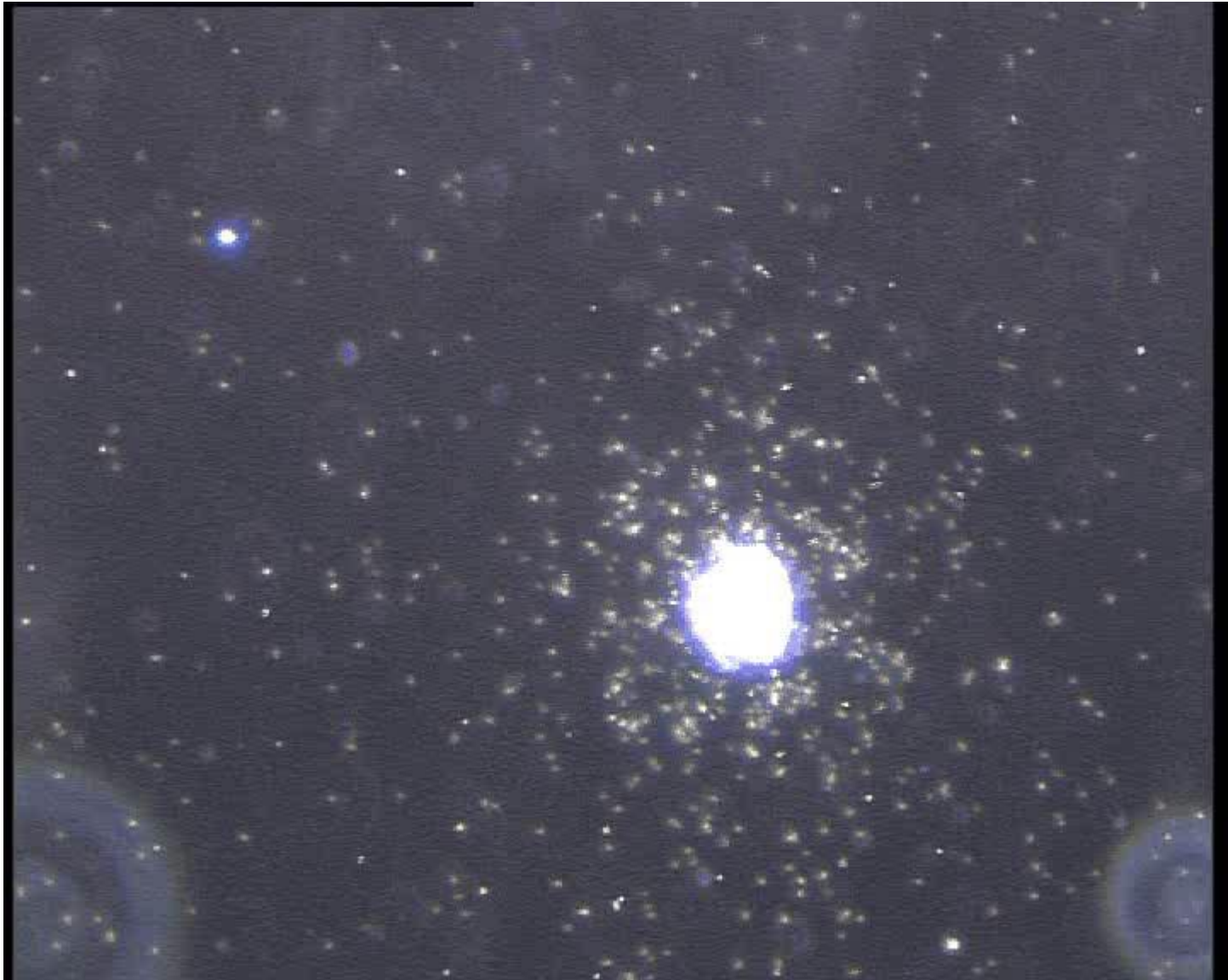
Surface feeding and zooplankton transport is large source of C required to feed mesopelagic bacteria & zooplankton carnivores



Steinberg et al., (2009)

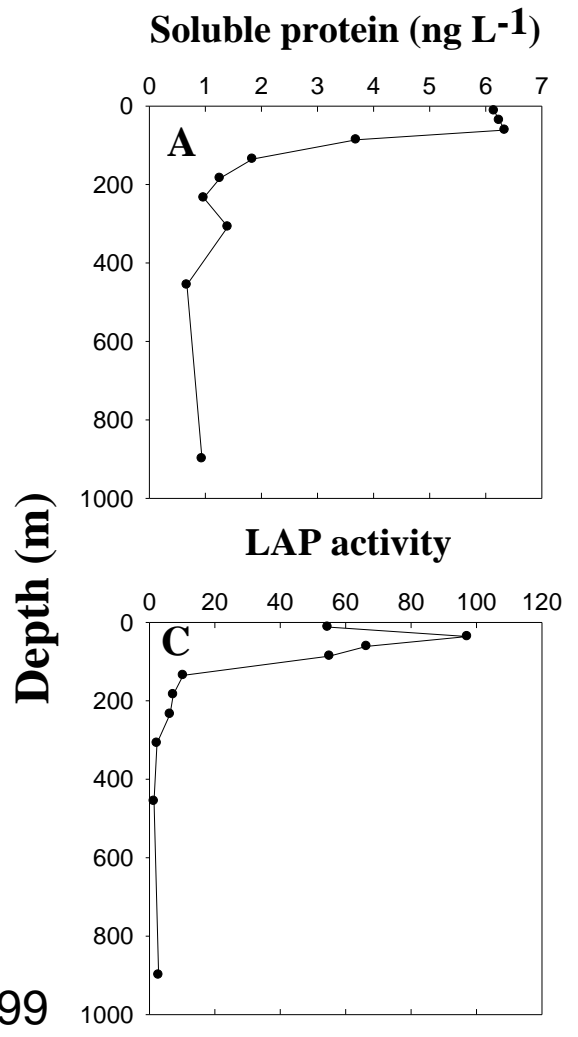
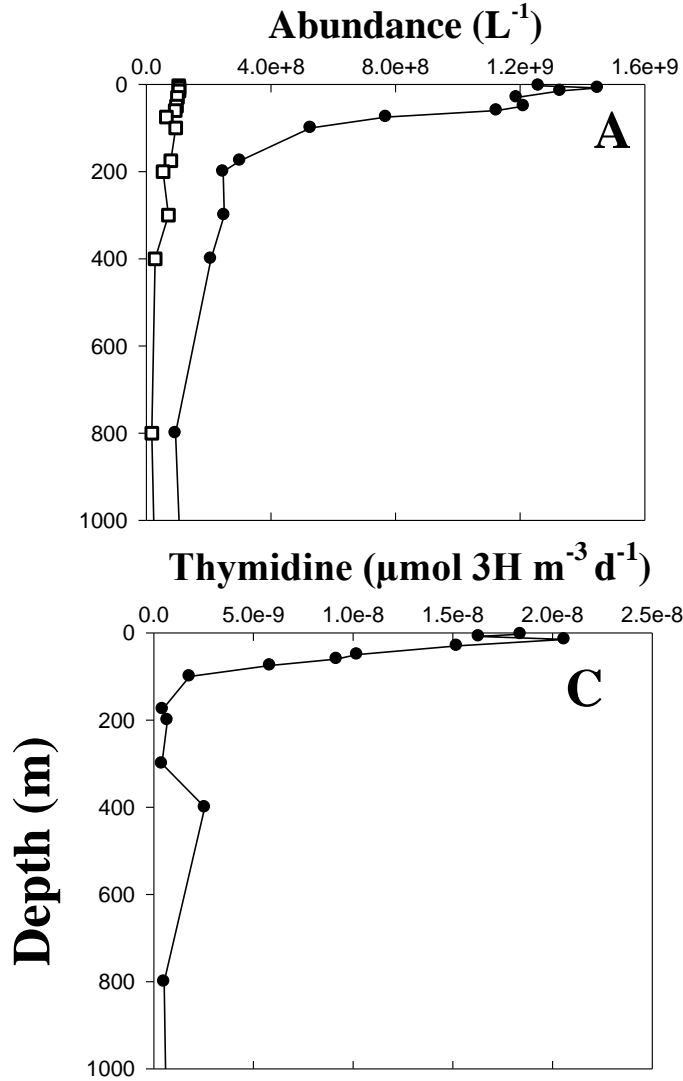
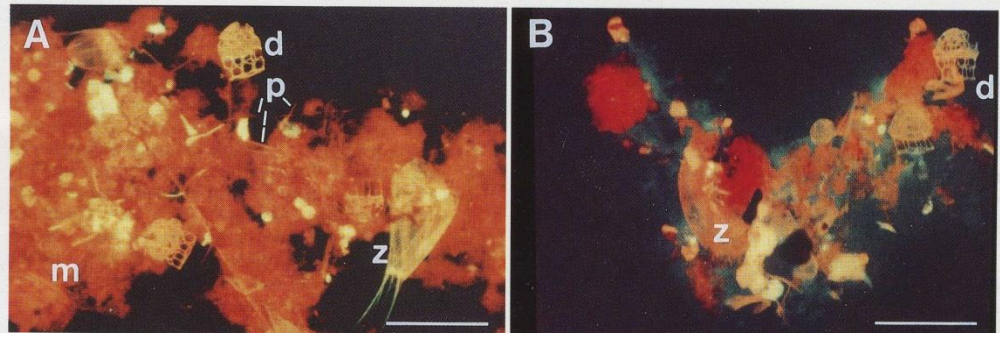
Particle transformations and foodwebs

Subsurface ocean - microbes



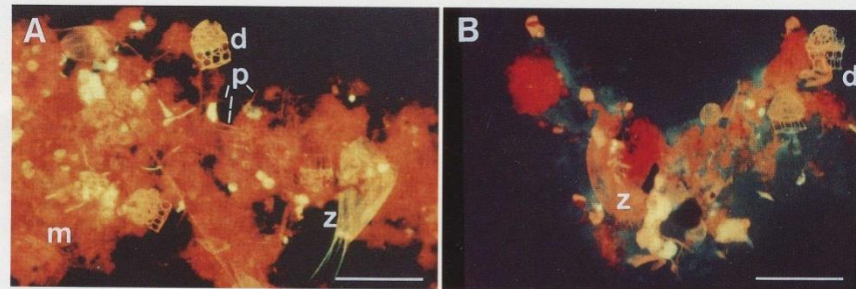
Courtesy Thomas Kiorboe Denmark

Bacteria solubilize particles



Boyd et al. 1999

Nutrient regeneration



N



C



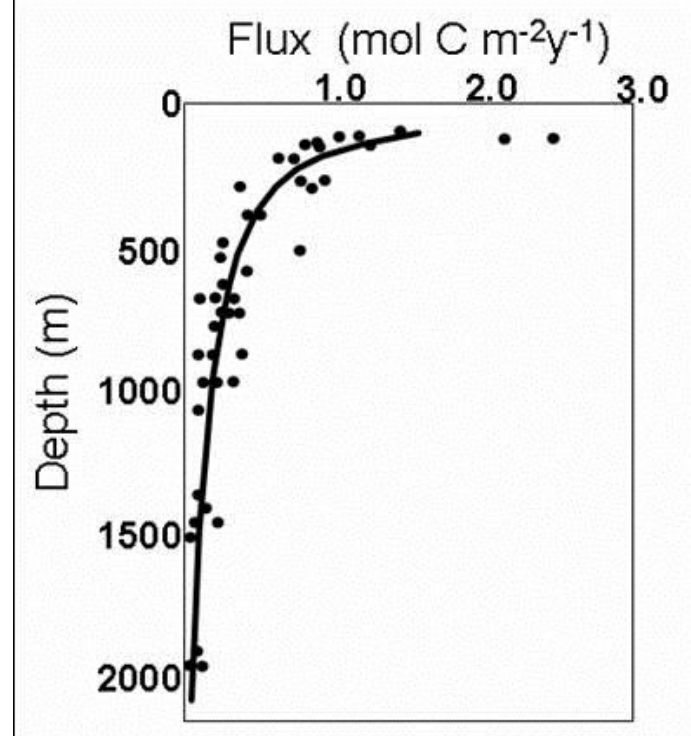
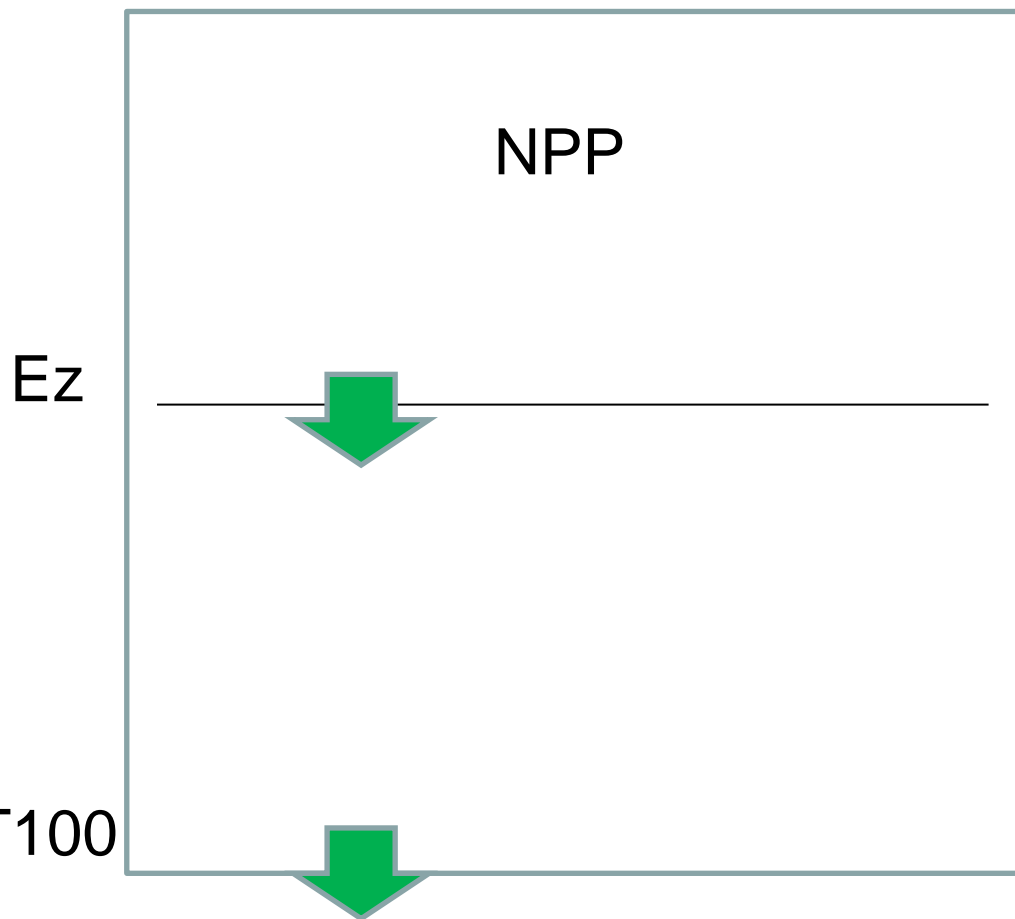
Si

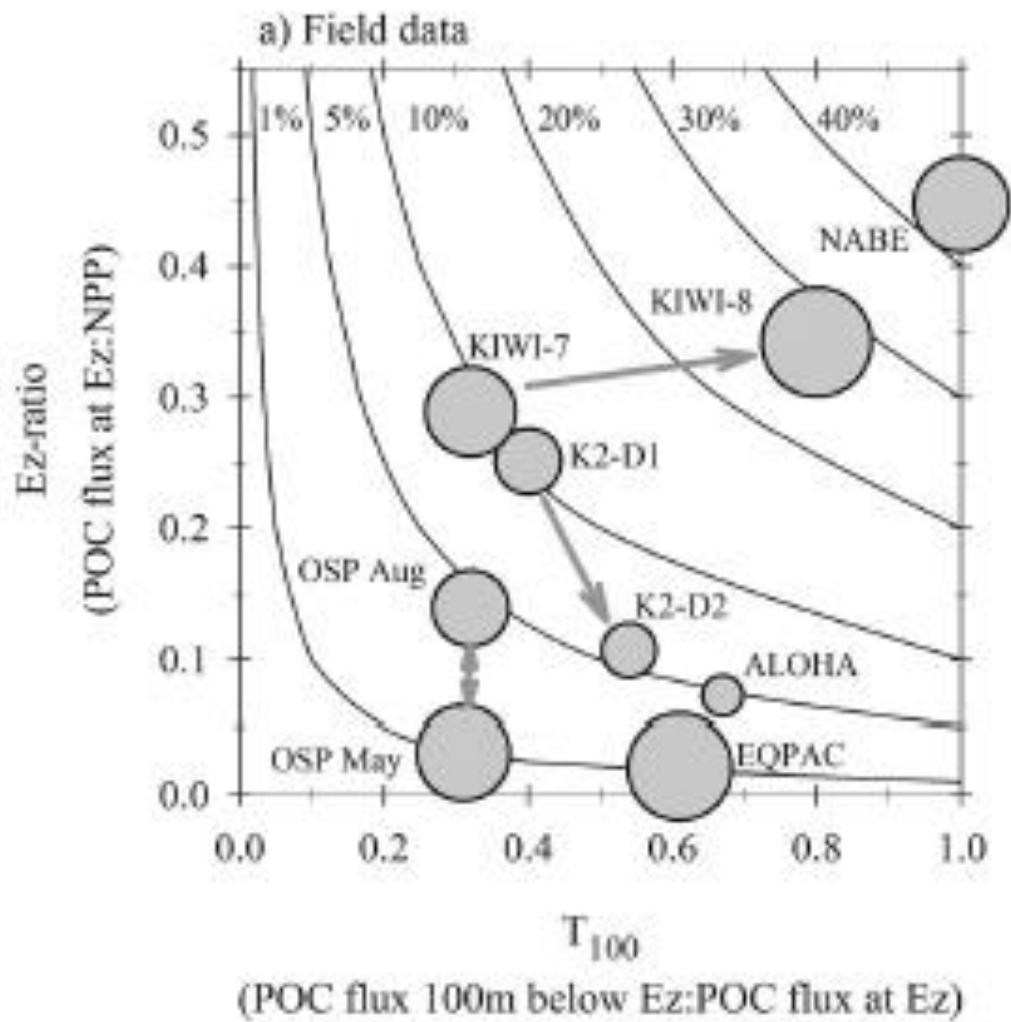


Fe

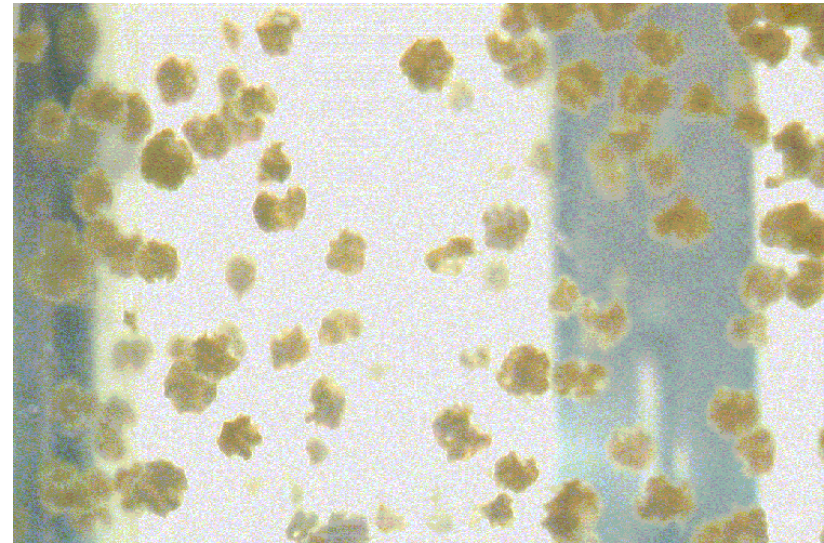
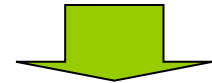
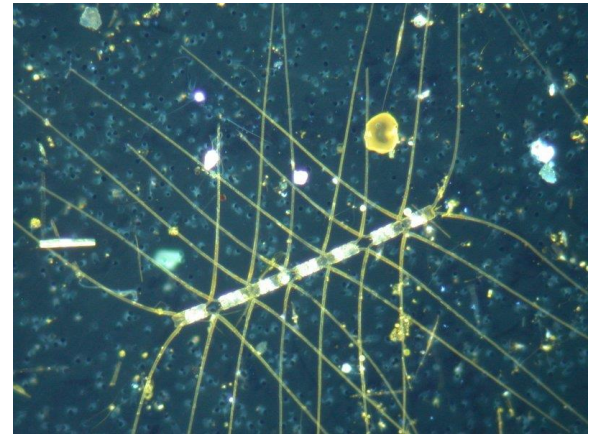
**Remineralization length scales
of major & trace elements are
microbially mediated**

Particle attenuation surface vs. subsurface waters

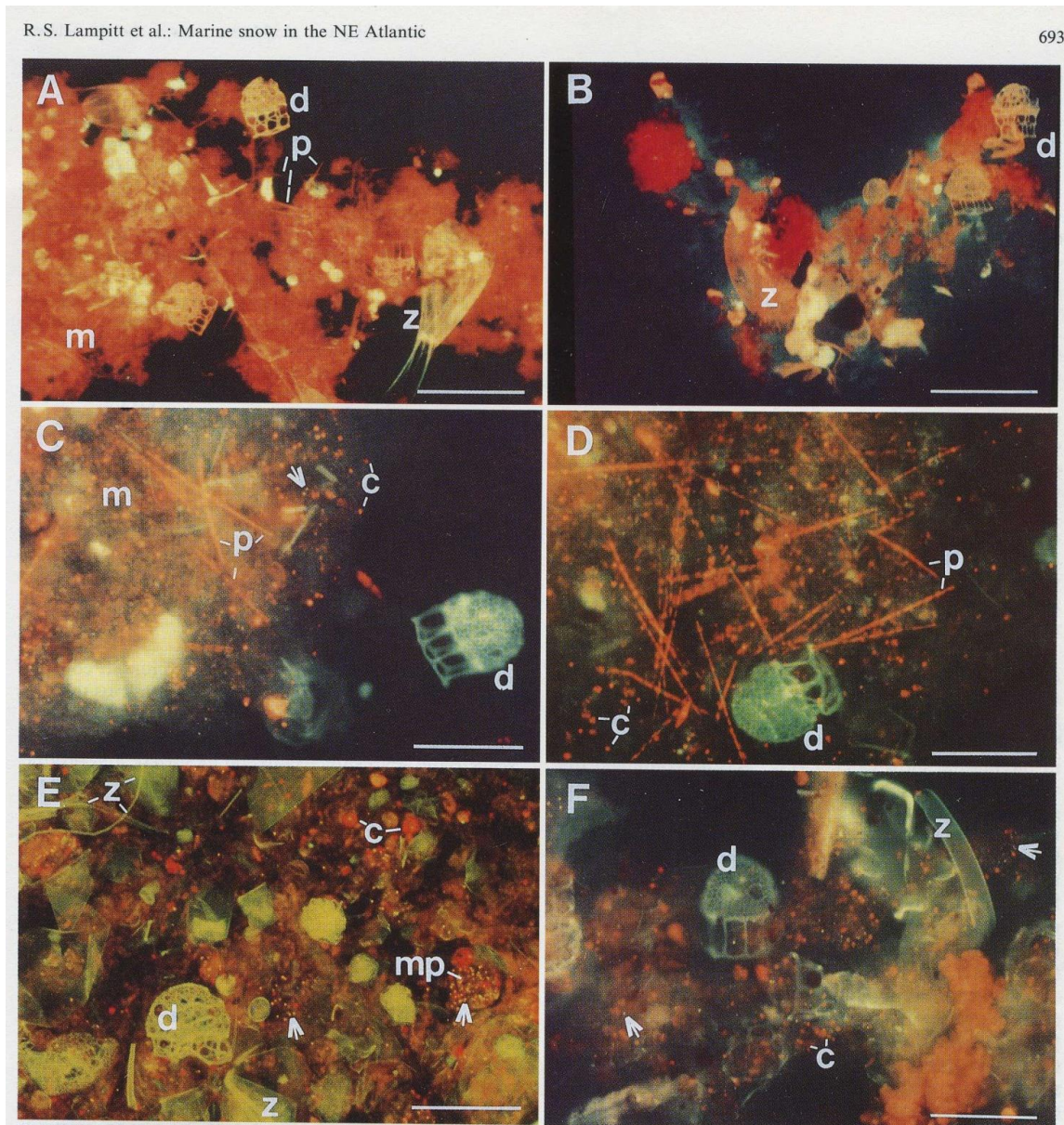


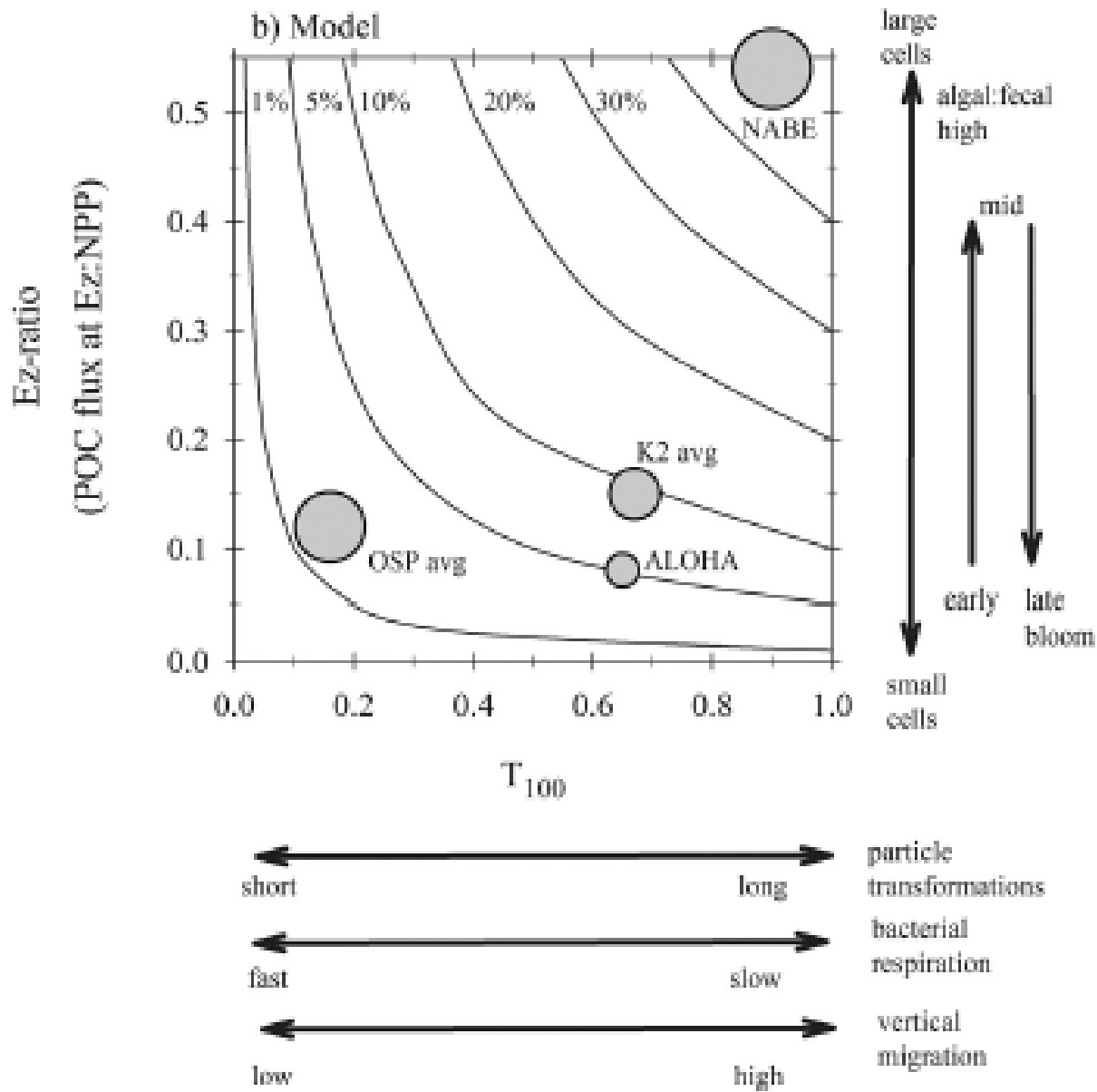


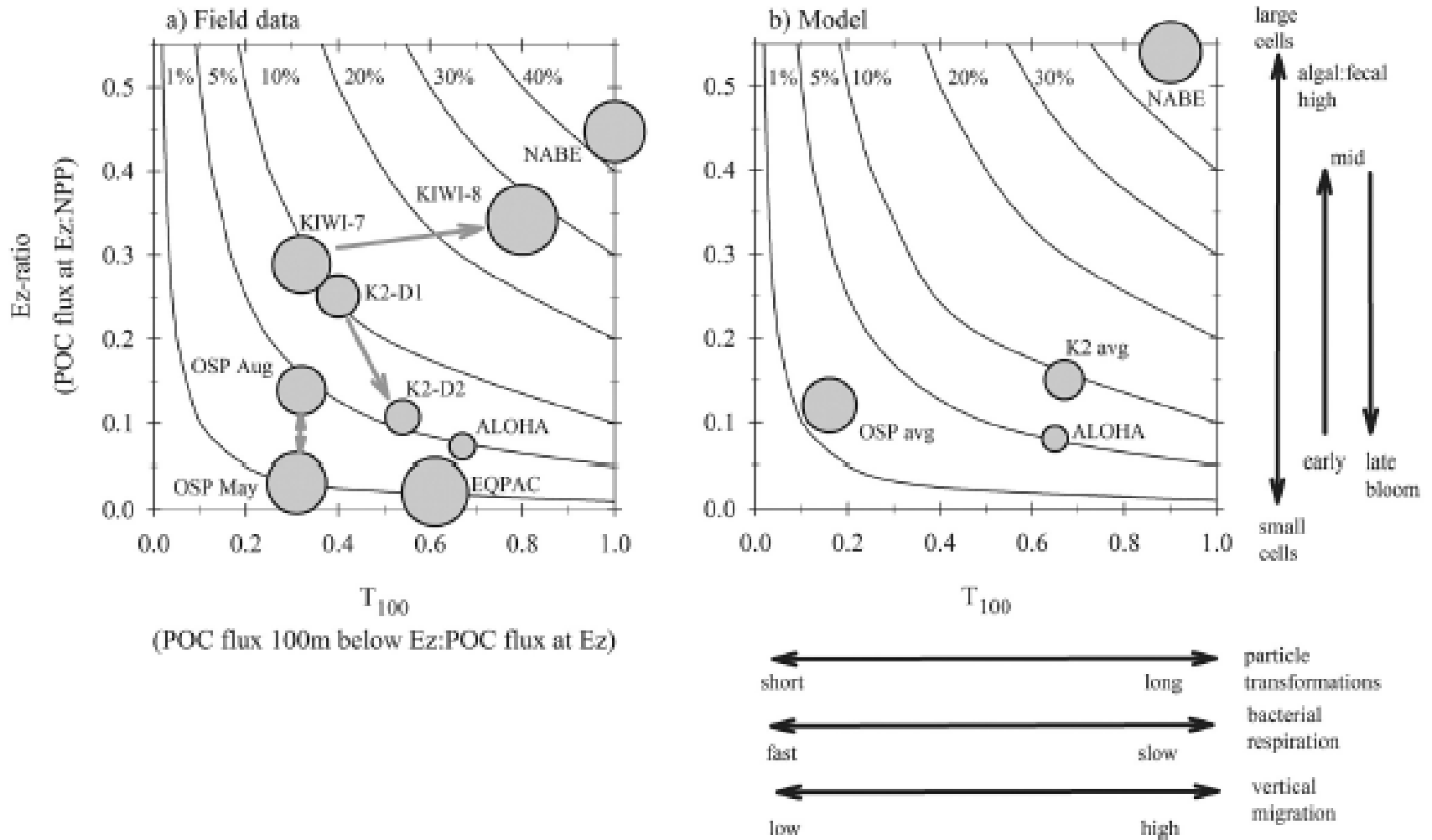
End-member – Direct algal sinking



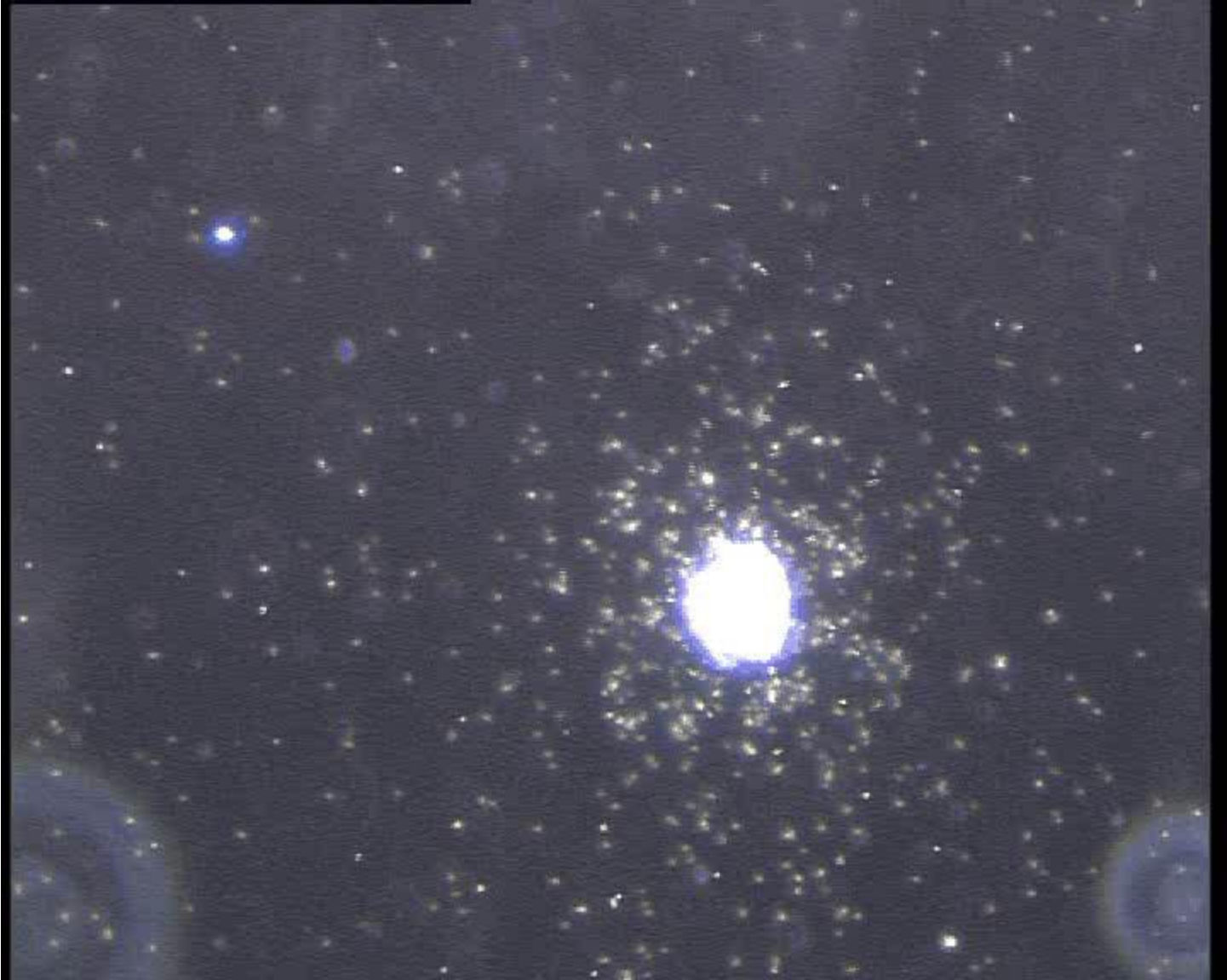
End-member – heavily transformed particles







Biological pump – at a cellular level



Courtesy Thomas Kiorboe Denmark

The path to preservation: Using proteomics to decipher the fate of diatom proteins during microbial degradation

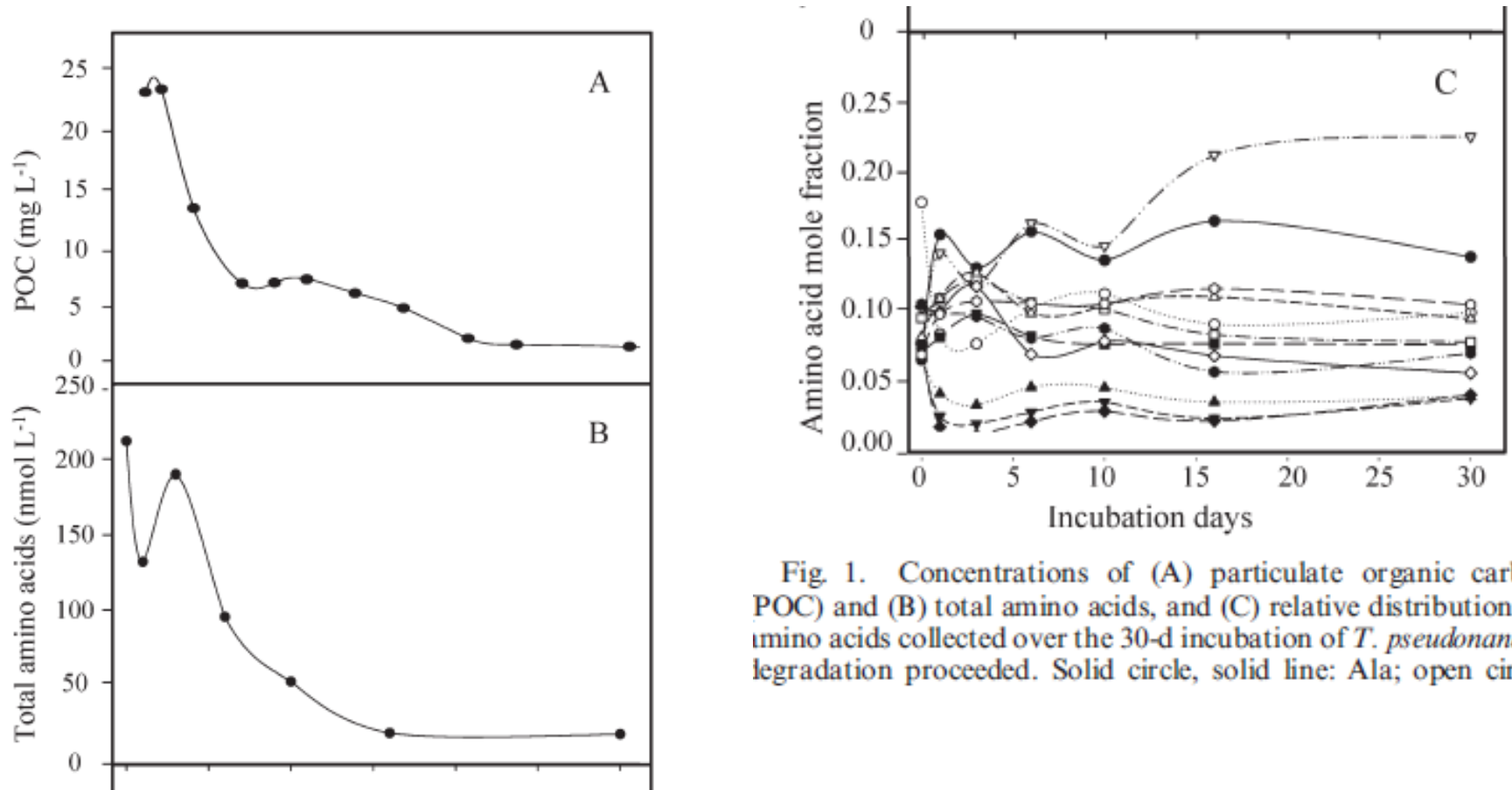


Fig. 1. Concentrations of (A) particulate organic carbon (POC) and (B) total amino acids, and (C) relative distributions of amino acids collected over the 30-d incubation of *T. pseudonana* as degradation proceeded. Solid circle, solid line: Ala; open circle,

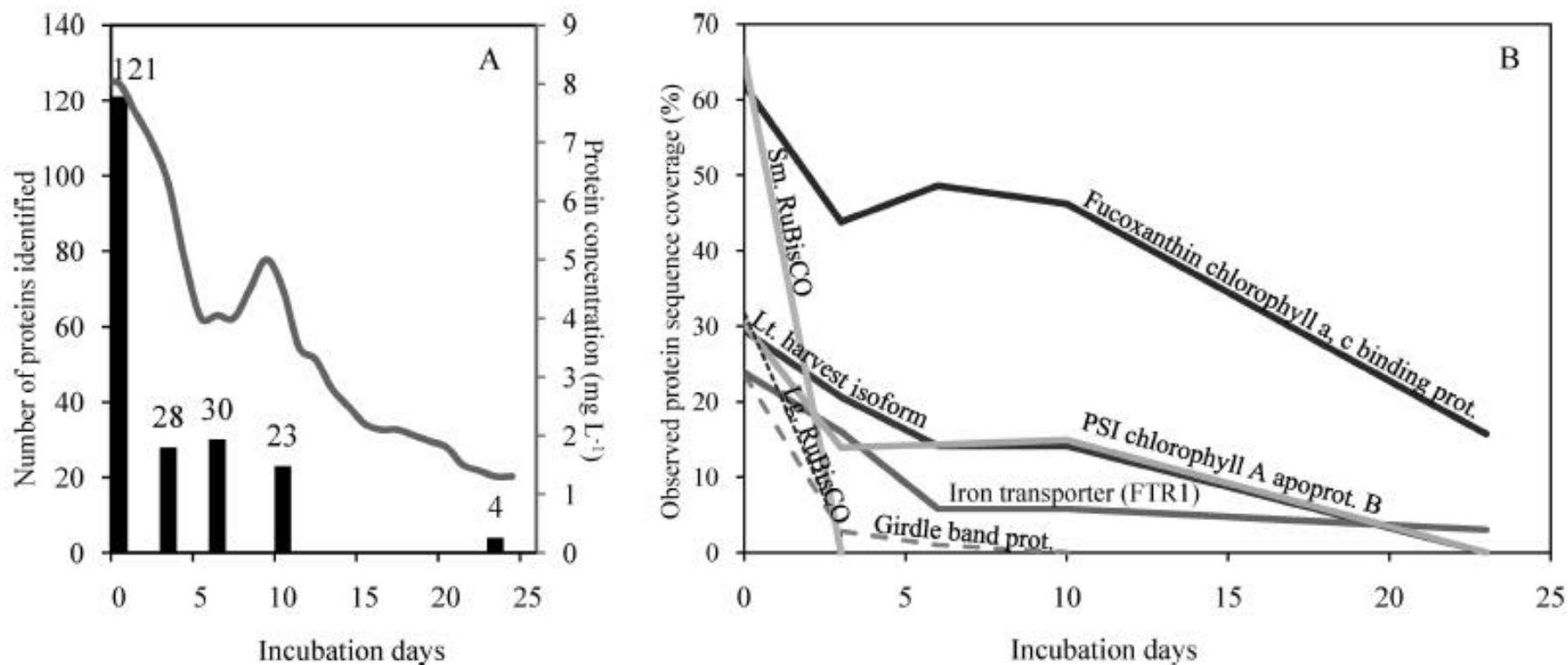
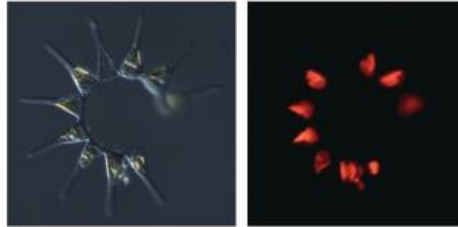


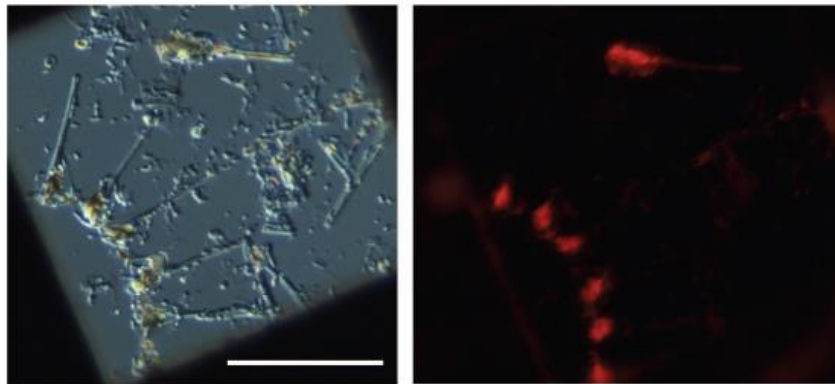
Fig. 2. (A) The time course of total particulate protein present during diatom degradation. Histogram: total number of proteins identified using LC-MS/MS; solid gray line: concentration of protein as measured by the Bradford assay. (B) Line graph tracking the percent of protein sequence detected using LC-MS/MS of seven different proteins through the degradation experiment.

Differential remineralization of major and trace elements in sinking diatoms

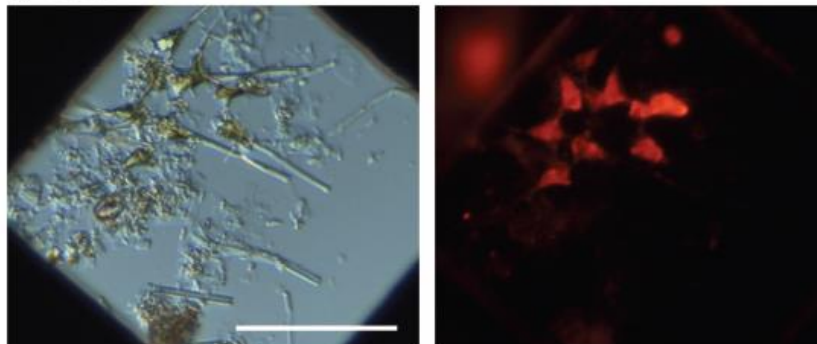
A) Surface



B) 100 m



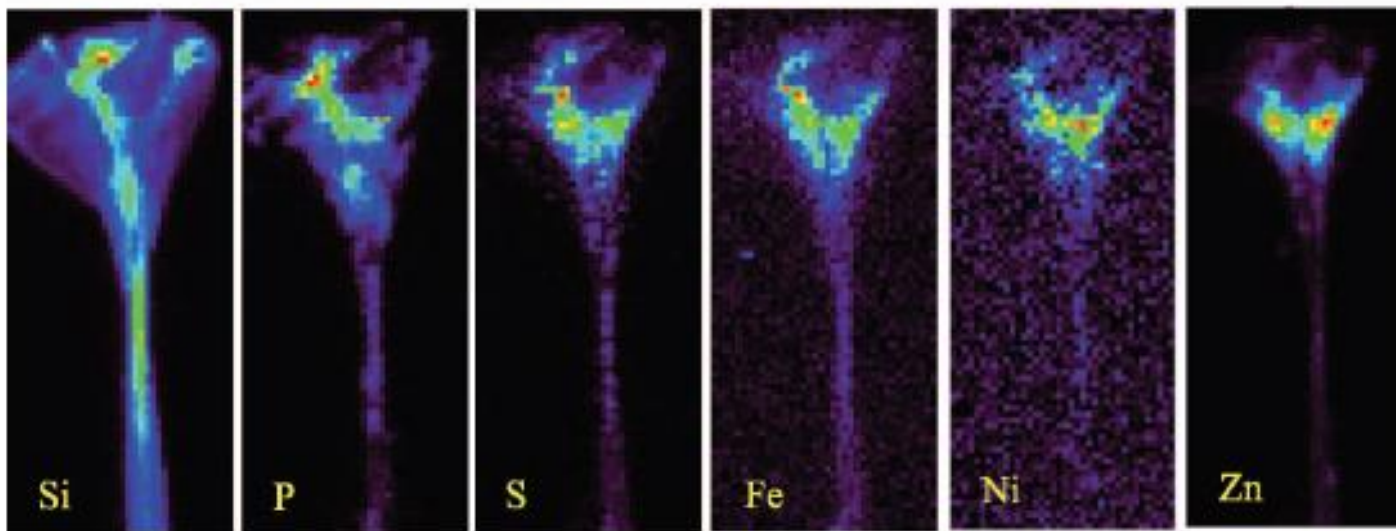
C) 200 m



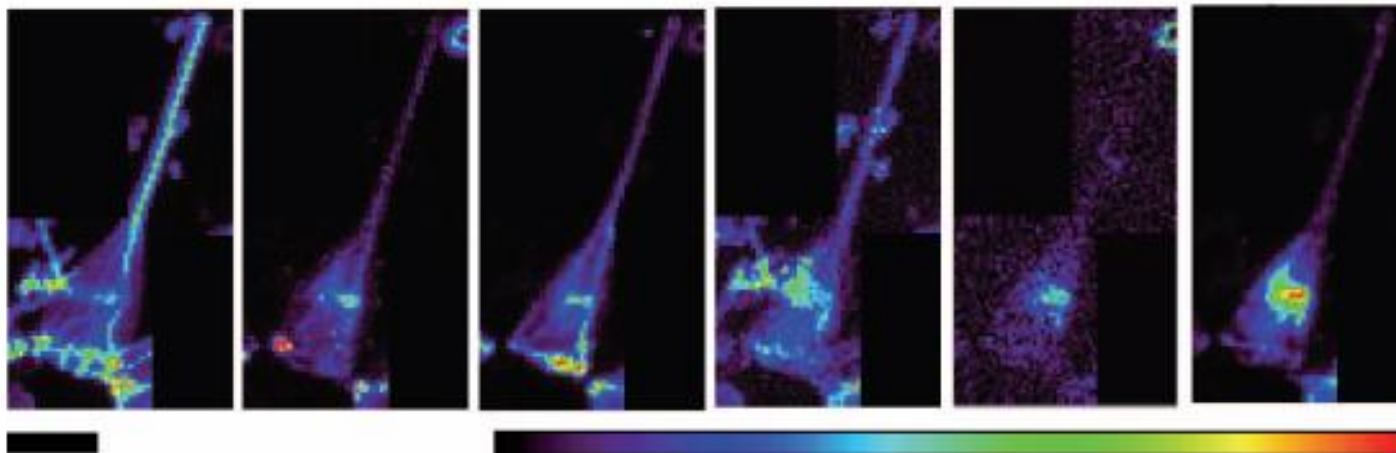
Twining et al.
2014

Fig. 3. Light (differential interference contrast) and epifluorescence (480 nm excitation)

A) 30 m



C) 200 m



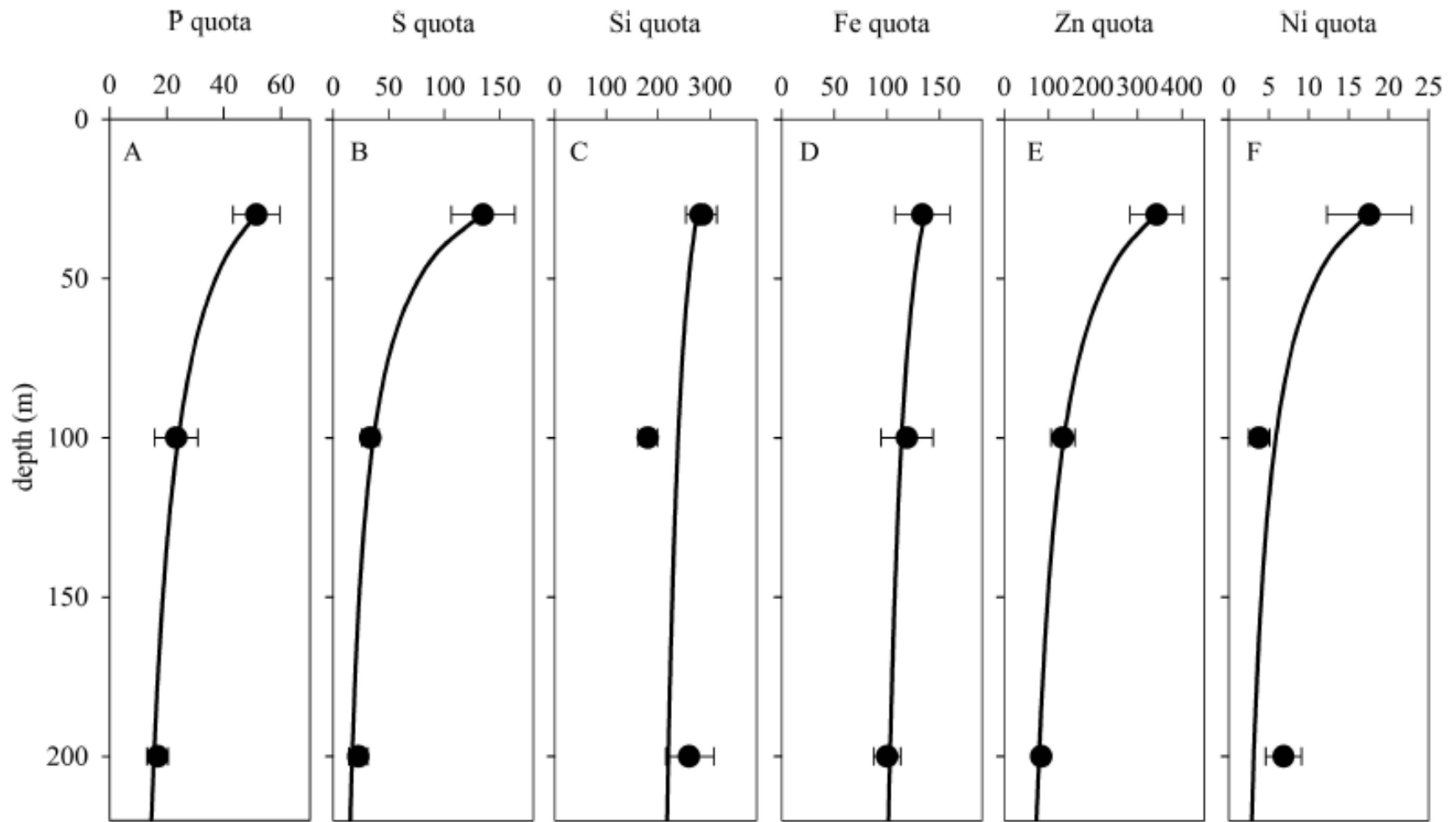
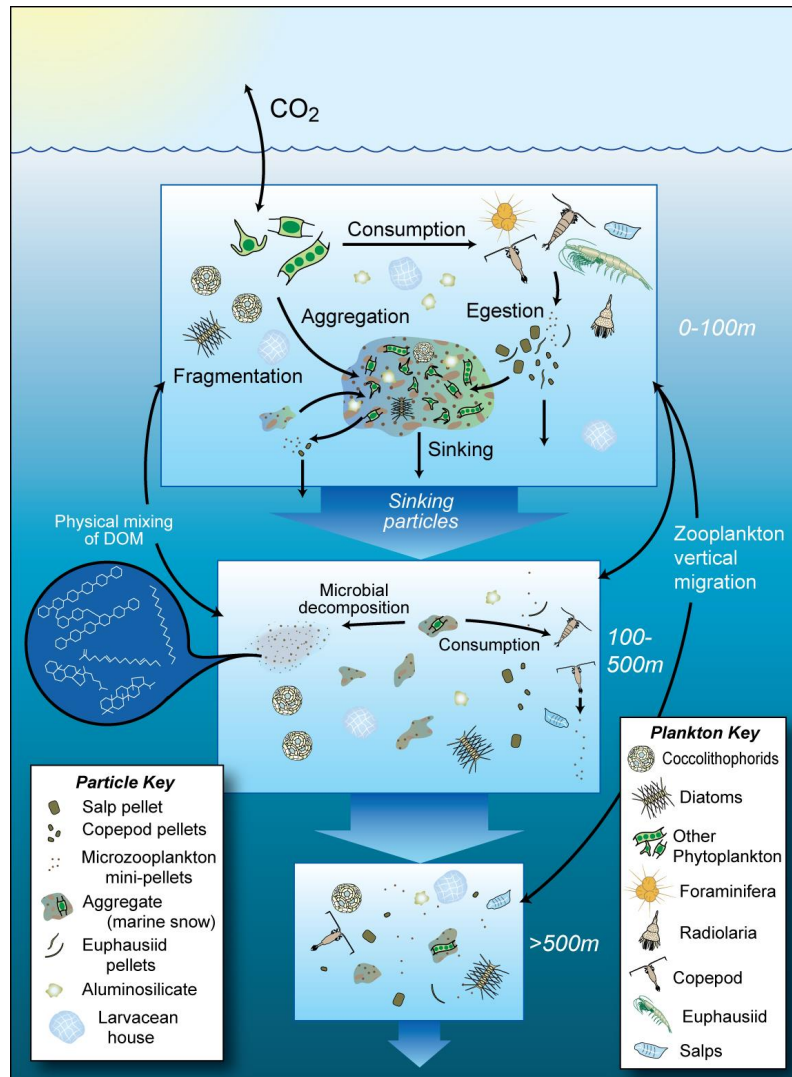


Fig. 4. Depth profiles of *Asterionellopsis glacialis* cellular element quotas for (A) P, (B) S, (C) Si, (D) Fe, (E) Zn, and (F) Ni. The quotas are means \pm SE ($n = 5-14$). A power-law function ($\text{Flux} = a(\text{depth})^{-b}$) has been fitted to each data set and the resulting attenuation coefficients (b) for the cells are shown in Table 1. P, S, and Si quotas are presented as fmol cell^{-1} , while Fe, Zn, and Ni quotas are presented as amol cell^{-1} .

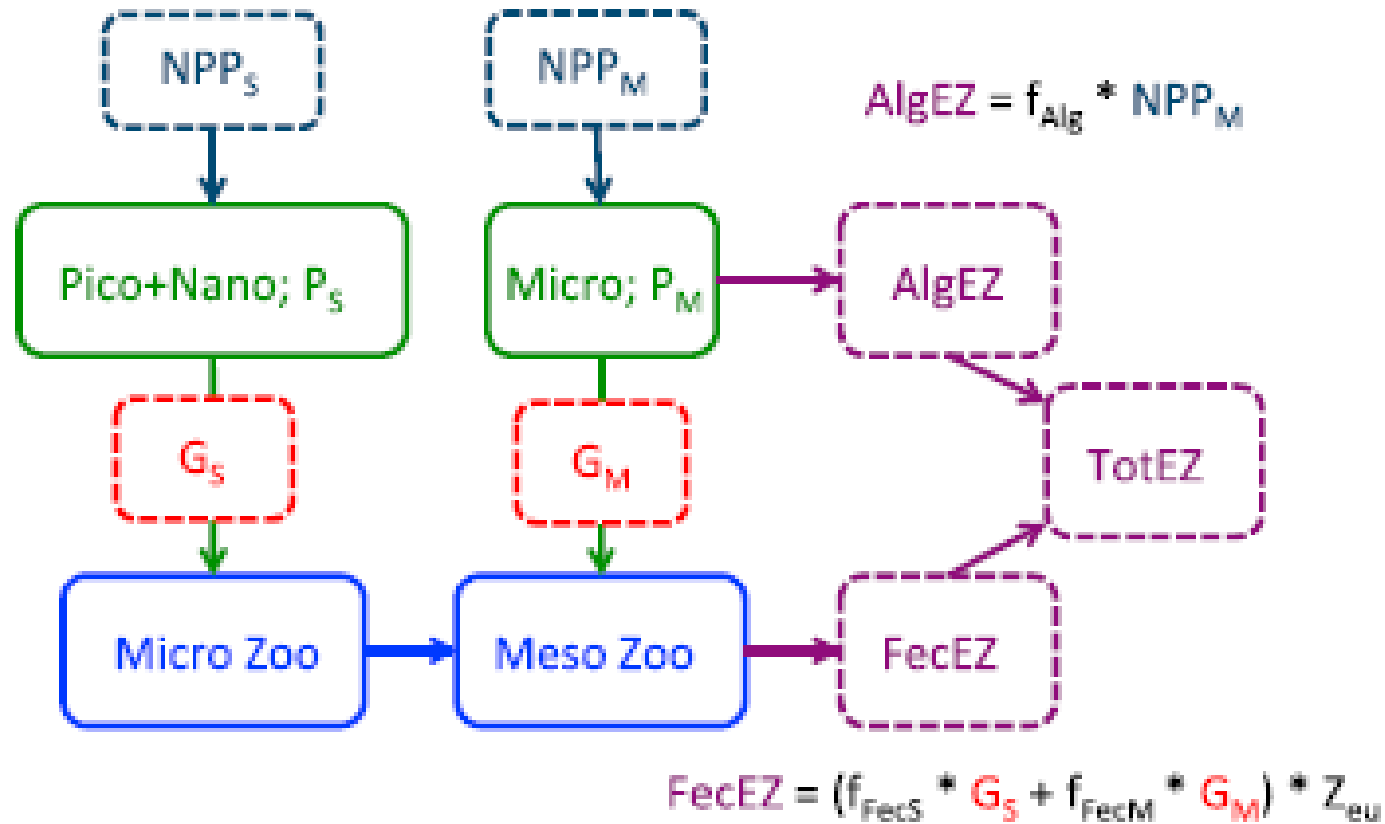
The Future

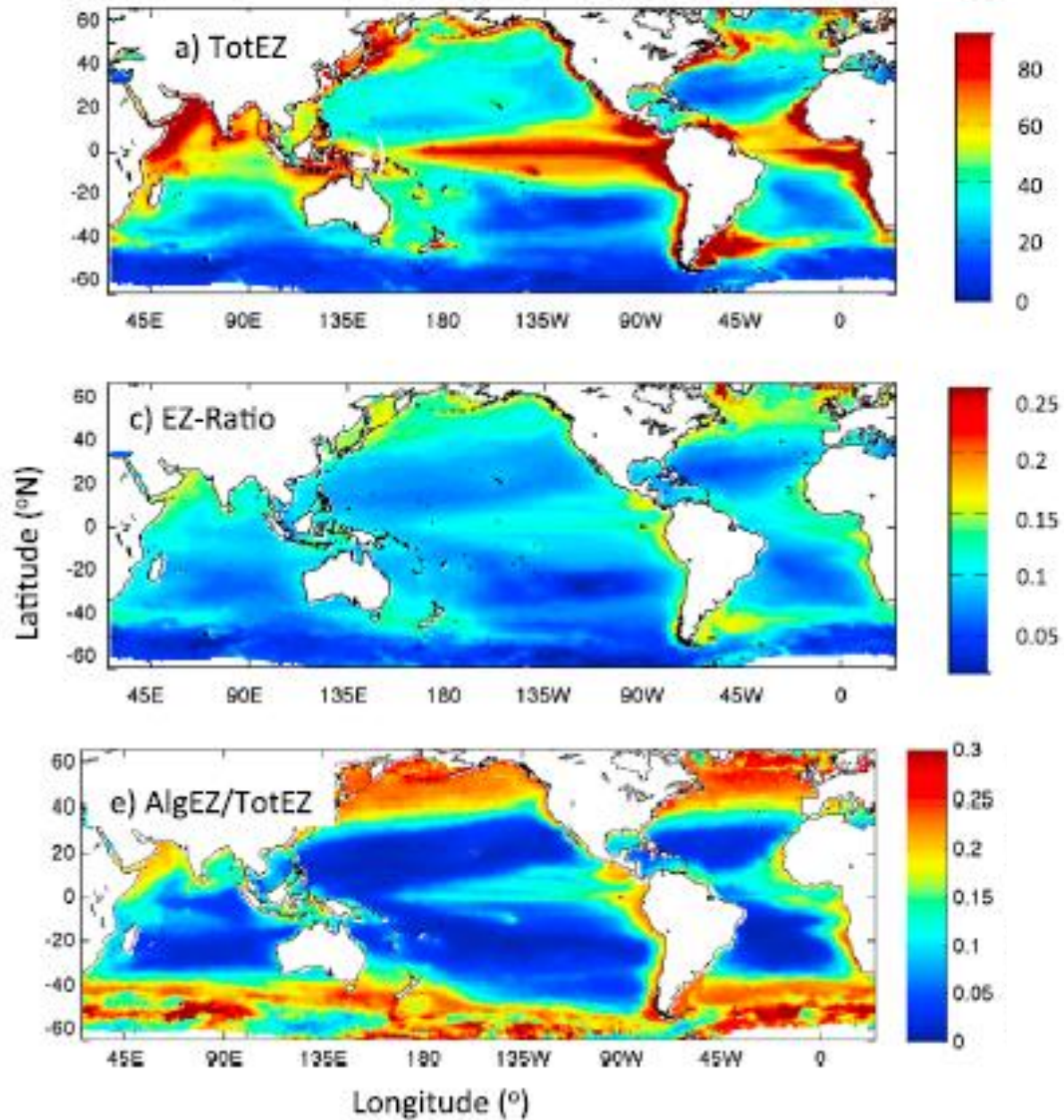
Detection and attribution of change in export



Global assessment of ocean carbon export by combining satellite observations and food-web models

D. A. Siegel¹, K. O. Buesseler², S. C. Doney², S. F. Sailley³, M. J. Behrenfeld⁴, and P. W. Boyd⁵



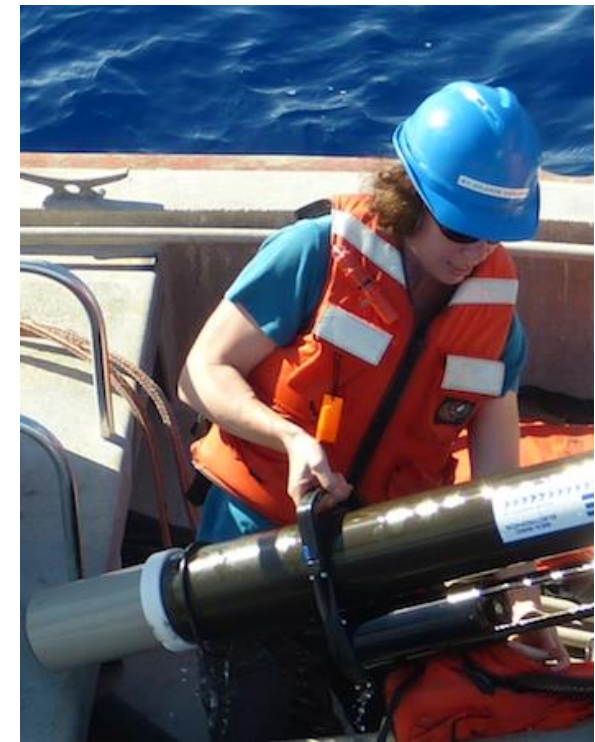
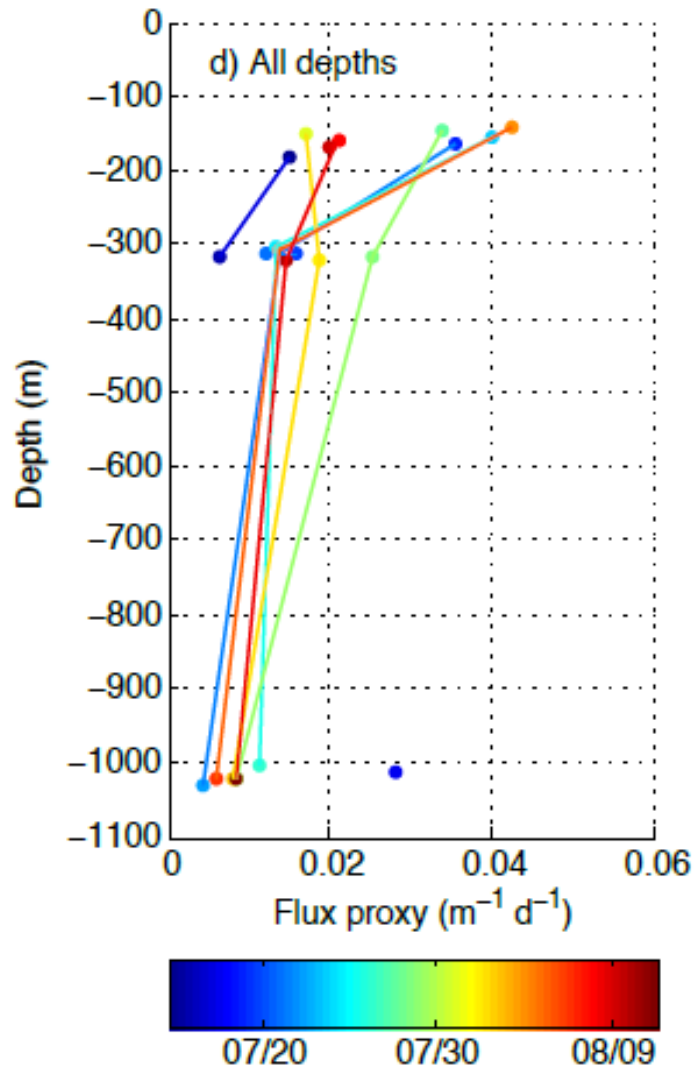


Log 10
 $\text{mg C m}^{-2} \text{d}^{-1}$

Autonomous, high-resolution observations of particle flux in the oligotrophic ocean

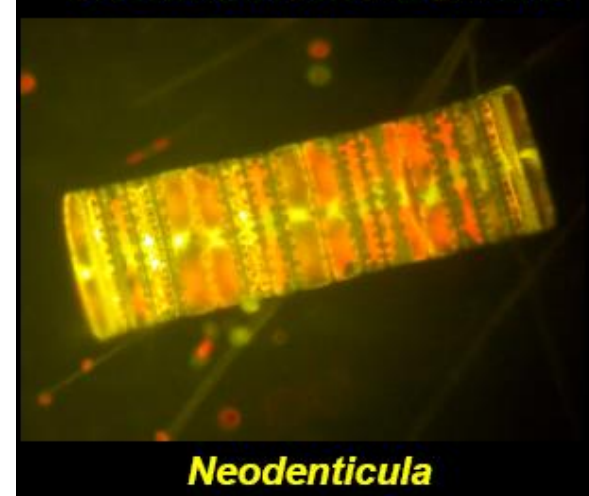
M. L. Estapa¹, K. Buesseler¹, E. Boss², and G. Gerbi³

2014



APEX float
With BOSS bio-optics

The biological pump and planktonic community structure



Conclusions

The pump drives multiple processes – C sequestration, nutrient regeneration, particle scavenging, nutrition of the deep ocean

The export of carbon from the surface is driven by foodweb transformations – from aggregation to faecal pellets

Carbon export in the subsurface ocean is controlled by both animal and microbial activity on particles

Export efficiency varies regionally and seasonally

Models are now providing the means to obtain global estimates of export from the surface ocean