

BIOGEOCHEMICAL AND ECOLOGICAL RESEARCH AT THE PORCUPINE ABYSSAL PLAIN

HA Ruhl*, SE Hartman, &
RS Lampitt *et al.*

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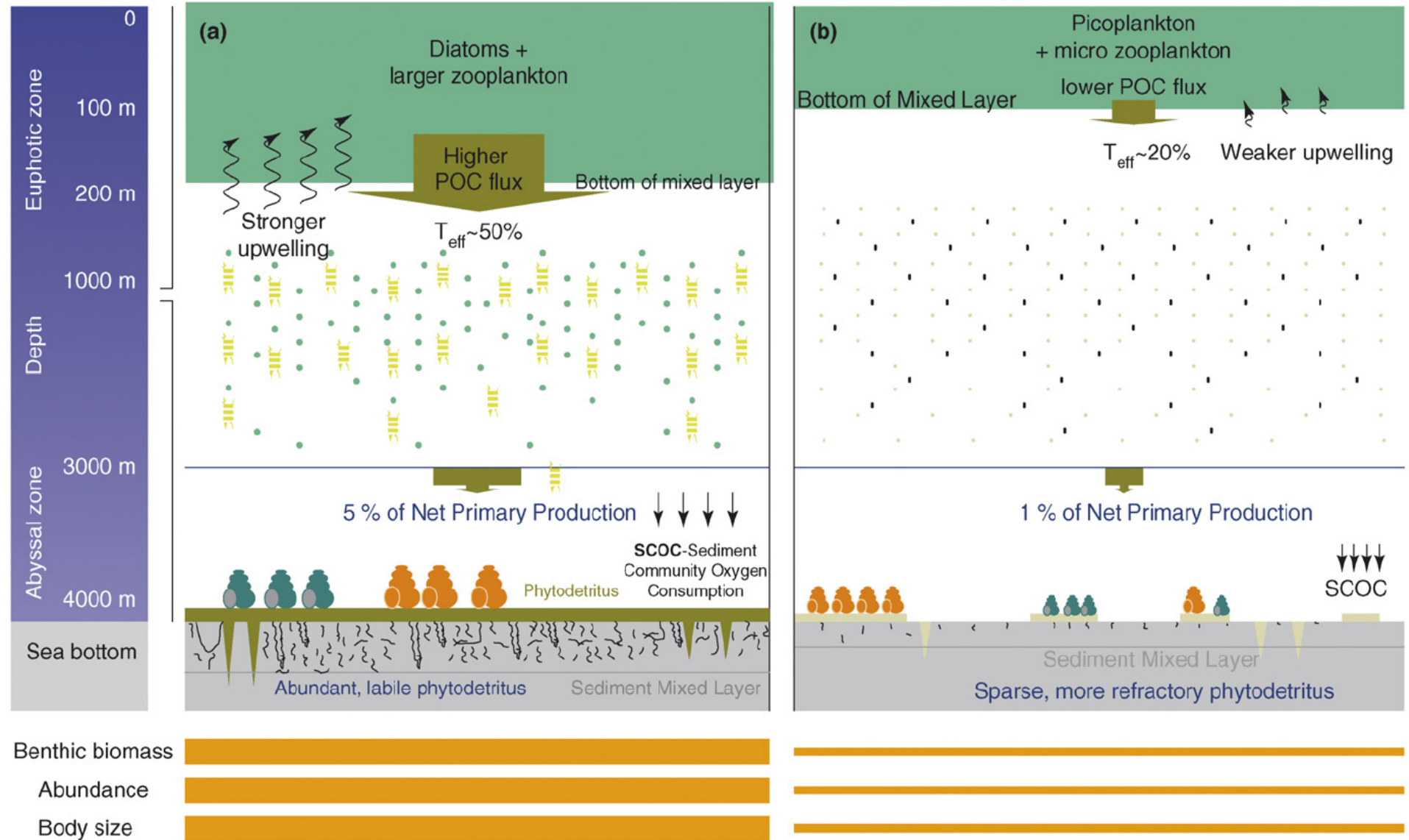
Atmosphere
 $p\text{CO}_2$

$P\text{CO}_2 = 384\text{ ppm}$ -today's concentration

$P\text{CO}_2 = 540\text{ ppm}$ -predicted by 2100

Lower sea surface temperature

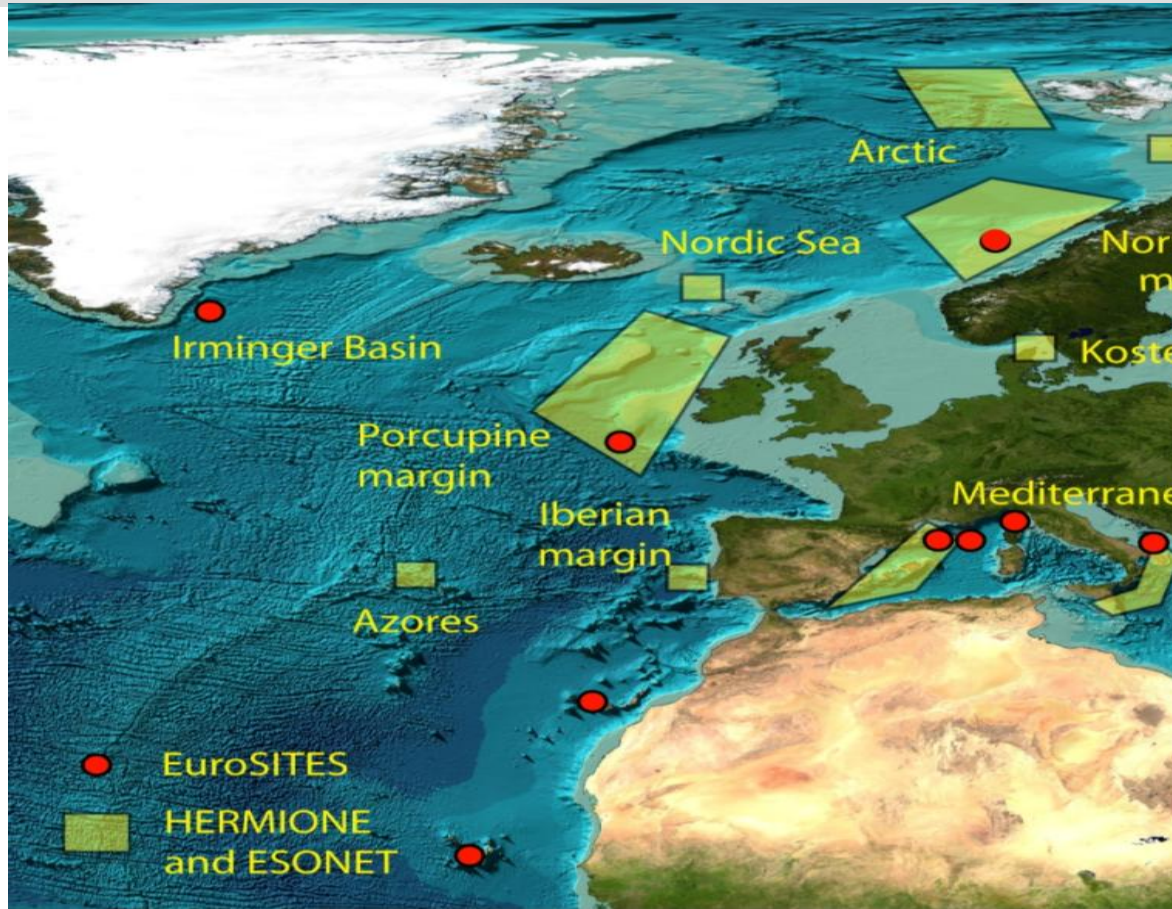
Higher sea surface temperature



Outline

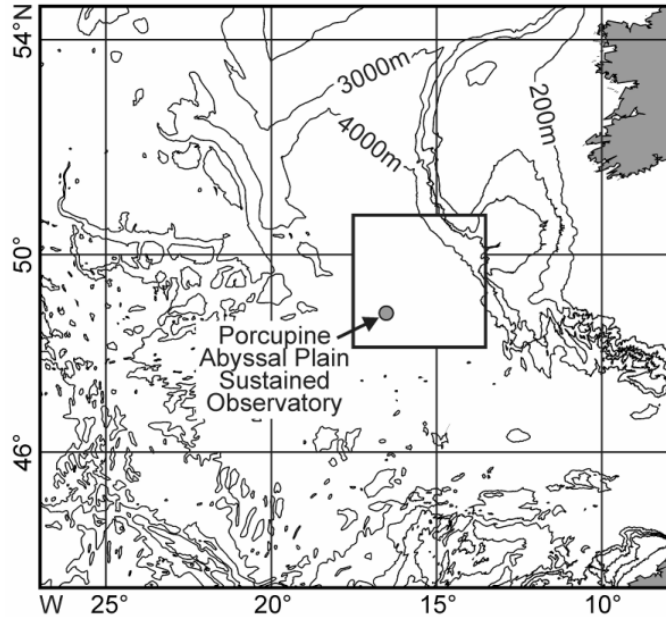
- The setting at Porcupine Abyssal Plain
- Selected time series measurements
- Biophysical interactions
- PP and CO₂ uptake
- Transfer of C into the deep ocean
- Seafloor life and the fate of C
- Future directions

Observatories in Europe



- NOC operates an observatory @ Porcupine Abyssal Plain.

Porcupine Abyssal Plain (PAP) - SO



- Research @ ~4,850m depth in NE Atlantic since 1989
- Evidence suggests POC flux at the site has links to the NAO & surface export flux
- Located in transition zone where MLD has dynamic links to PP and NAO variation
- UK contribution to OceanSITES and GOOS
- Major European open ocean observatory delivering real-time data, 'pre-operational' use
- Collaboration with MetOffice and many others



The PAP setting

North Atlantic Oscillation



Positive Phase



Negative Phase

The PAP setting

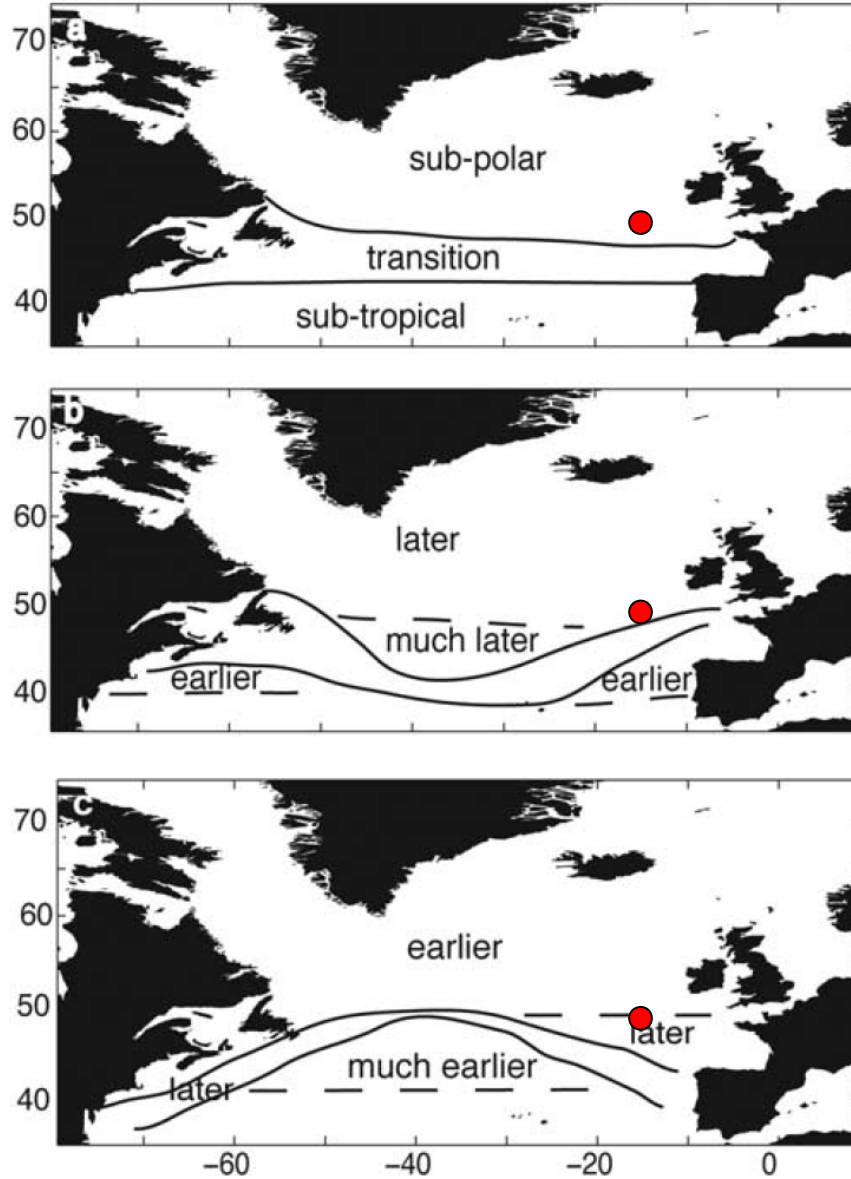


Figure 8. Sketch of the locations of the three bloom regions under (a) mean conditions, (b) in a positive NAO phase, and (c) in a negative NAO phase (dashed line marks position of transition region under mean conditions). The nature of the response in bloom timing is marked.

The PAP setting

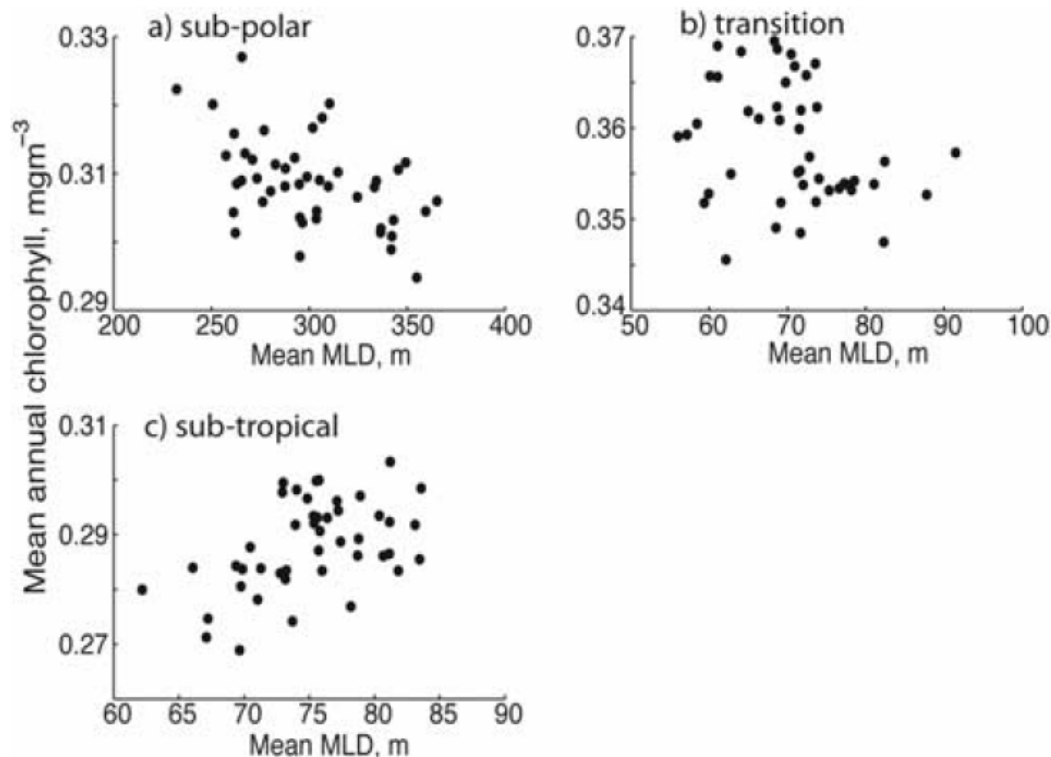
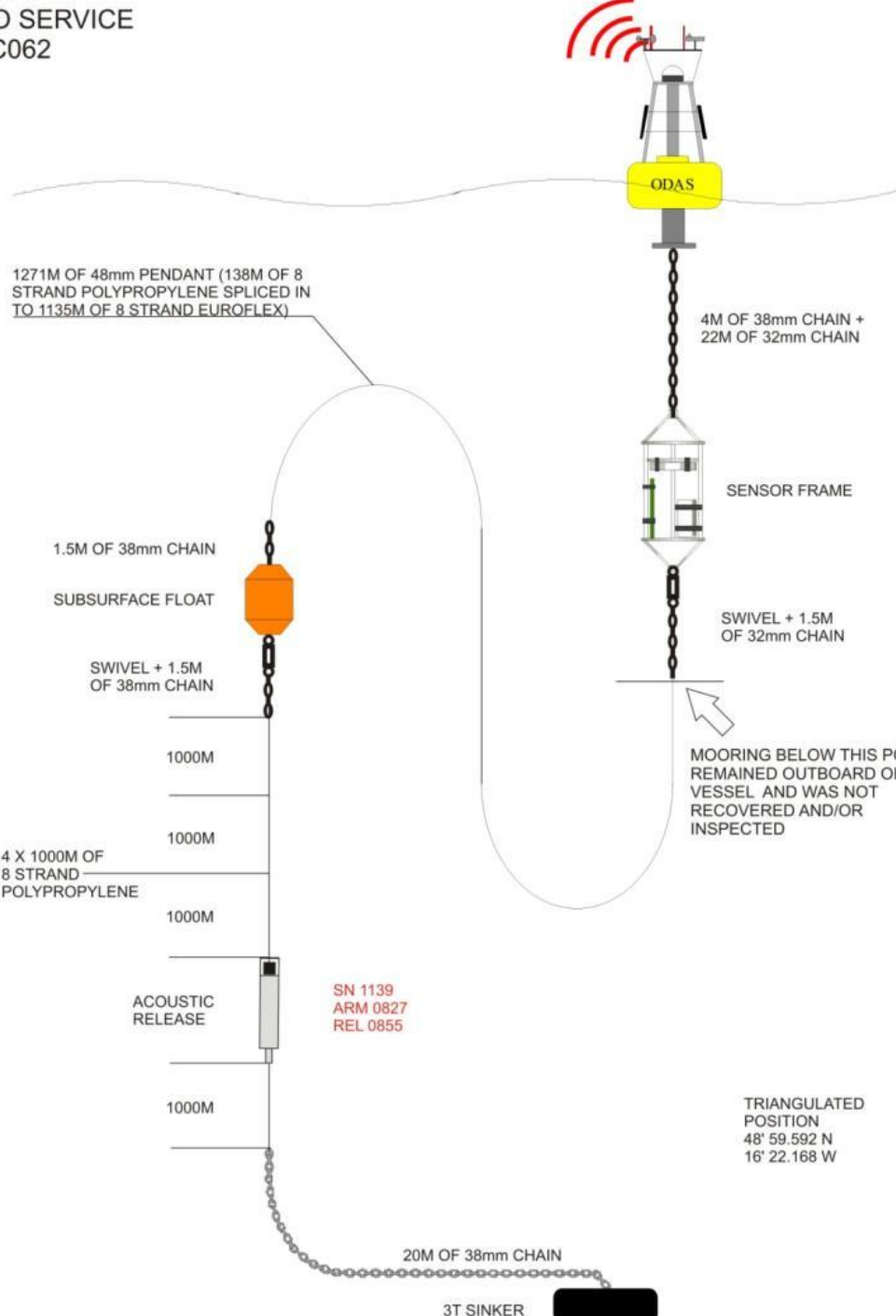


Figure 9. Mean annual modeled chlorophyll plotted as a function of mean annual MLD for the (a) subpolar, (b) transition, and (c) subtropical regions. Linear correlation coefficients are -0.48 ($p < 0.05$) and 0.49 ($p < 0.05$) for the subpolar and subtropical regions, respectively. Correlation coefficient for the transition region is not statistically significant.

PAP1
TO SERVICE
JC062

PAP 3
Deployed JC062
27/7/2011
48°59.382' N
16°29.585' W

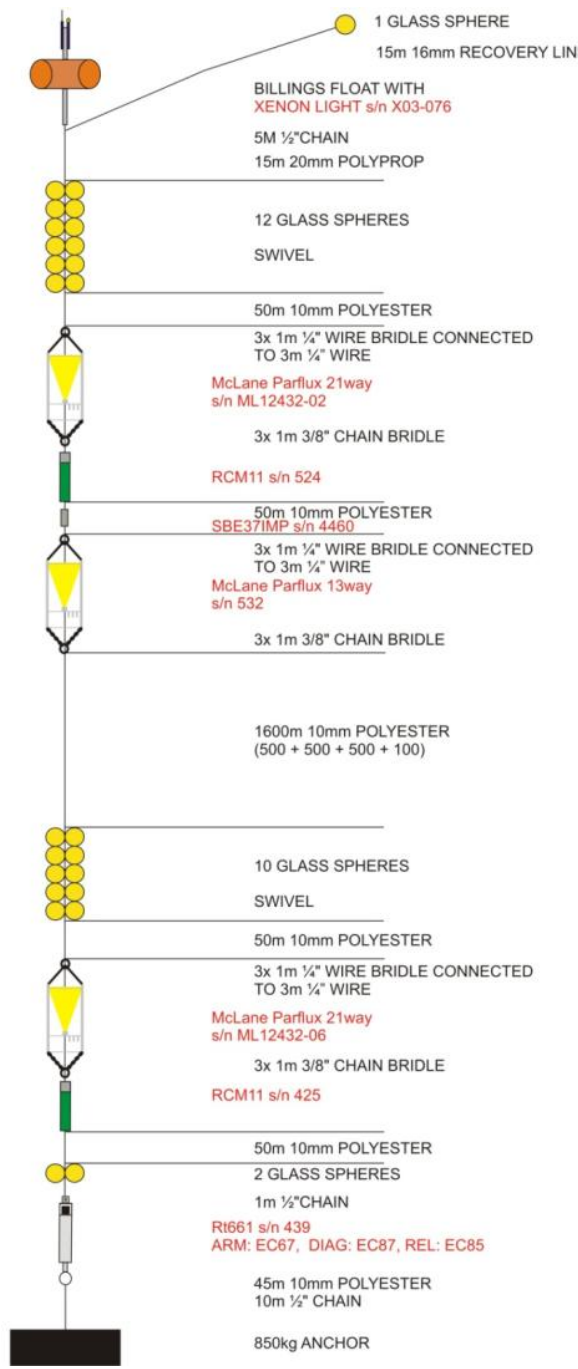


SEDIMENT TRAP
1800m from seabed

SEDIMENT TRAP
1750m from seabed

SEDIMENT TRAP
100m from seabed

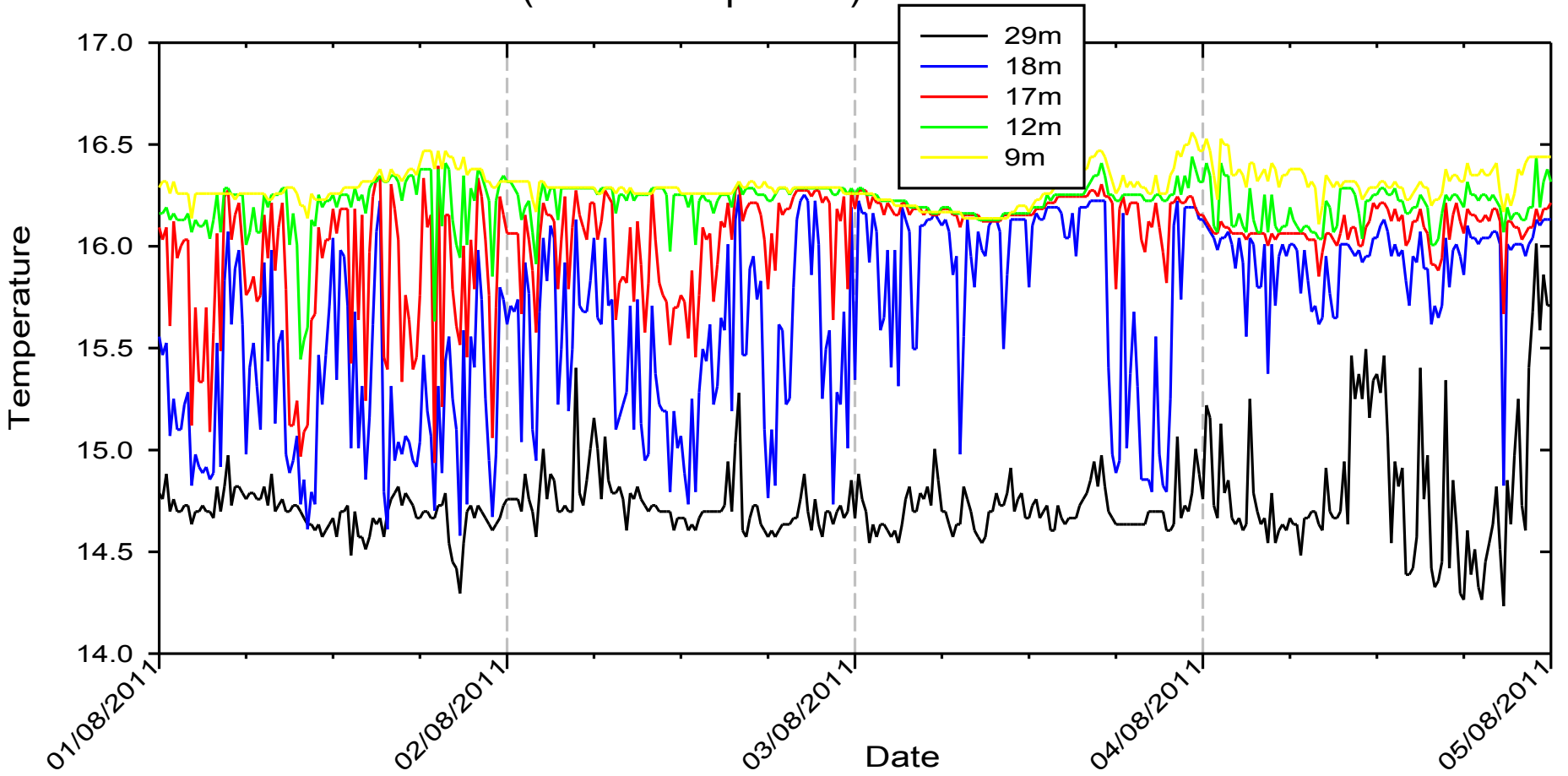
WATER DEPTH 4814m



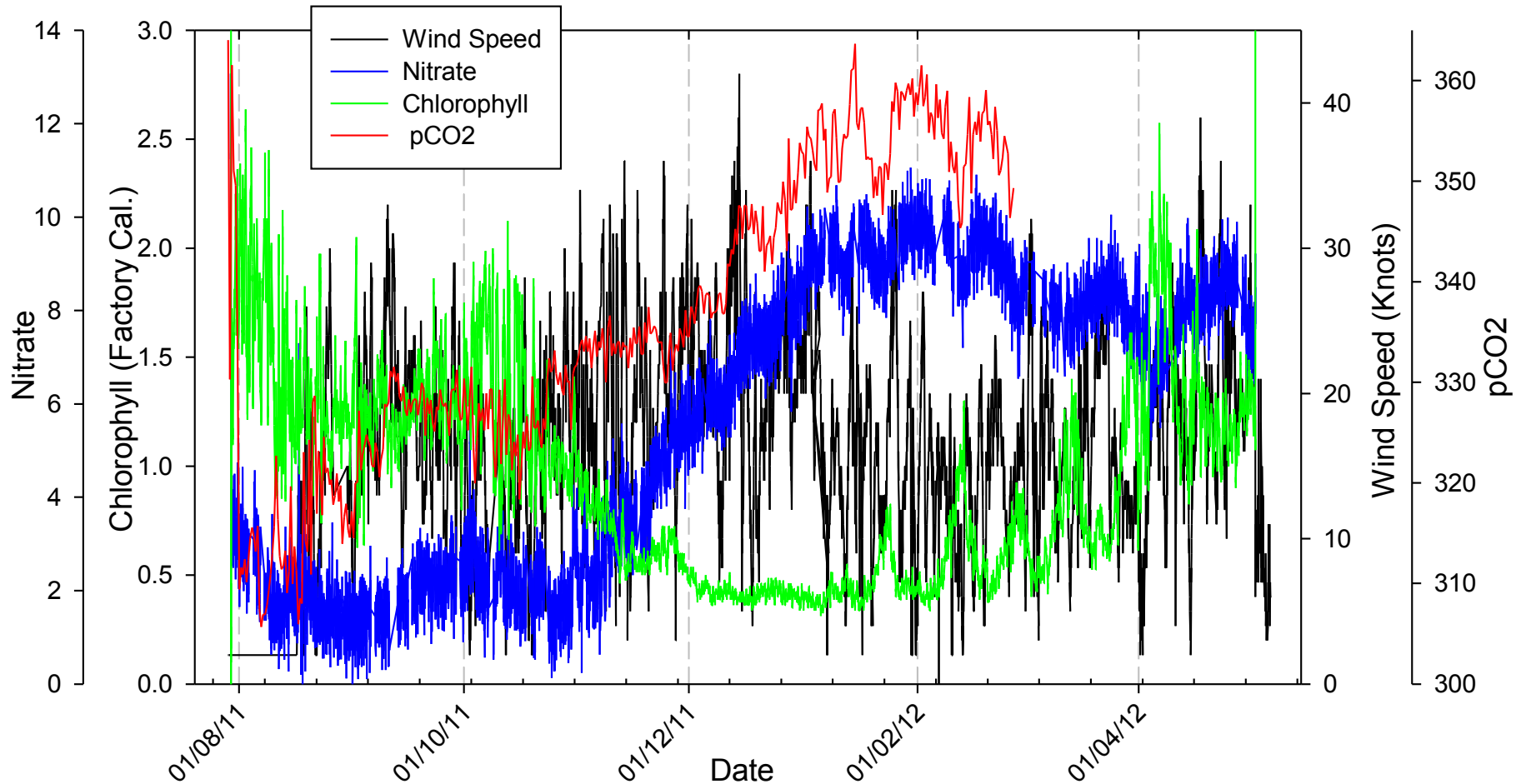


MOORED DATA BUOY
ID 4413

Variation in sea surface temperature (over the top 30 m) at the PAP-SO



Time-series data from PAP



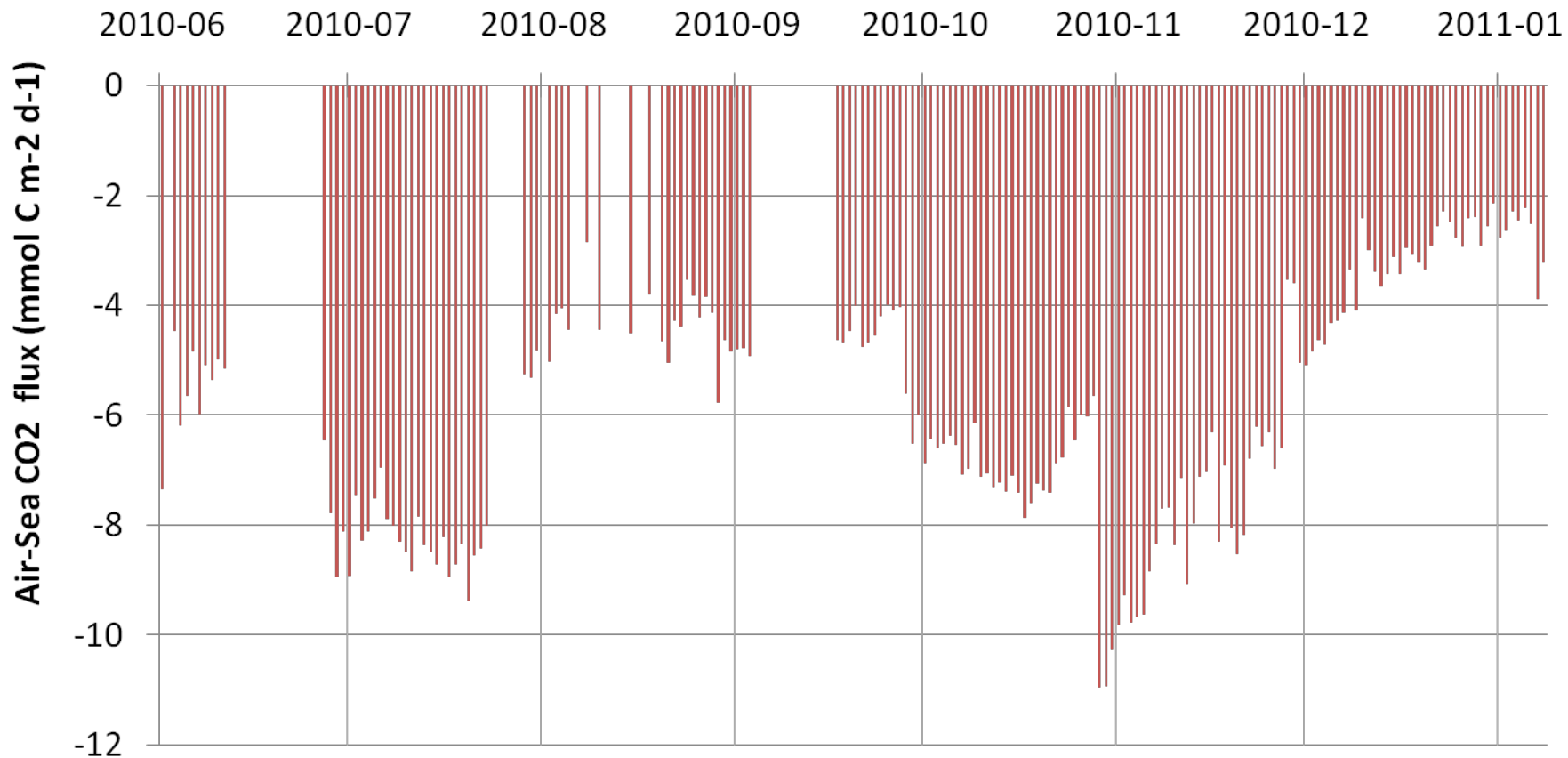
Aug. 2011 – May 2012

Data from 30 m depth from recent deployment of the PAP-SO:

Nitrate & pCO₂ increases with convective mixing in the winter months as chlorophyll-fluorescence decreases.

PAP-SO Air/sea CO₂ flux 2010, calculated using Nightingale parameterization

Illustrates that PAP is a CO₂ 'sink'



Variability at the PAP site

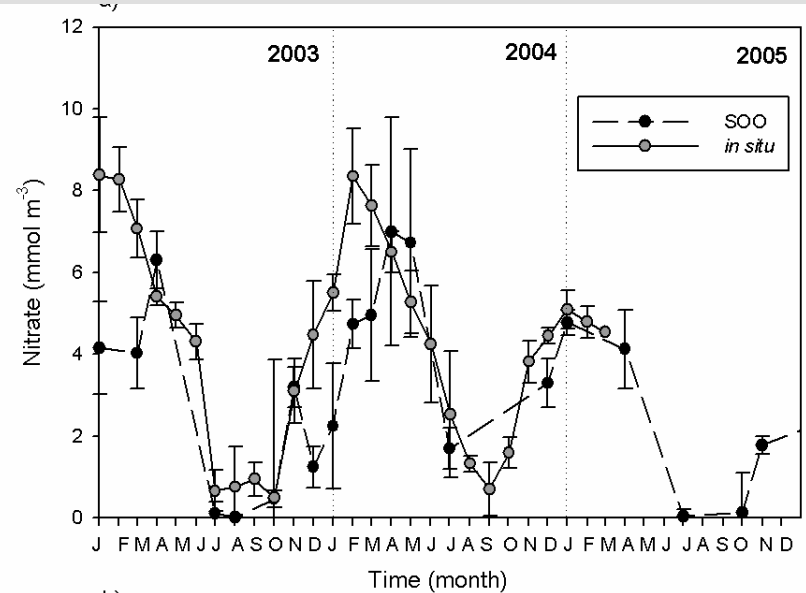
Decrease in nitrate

Not due to winter
mixing

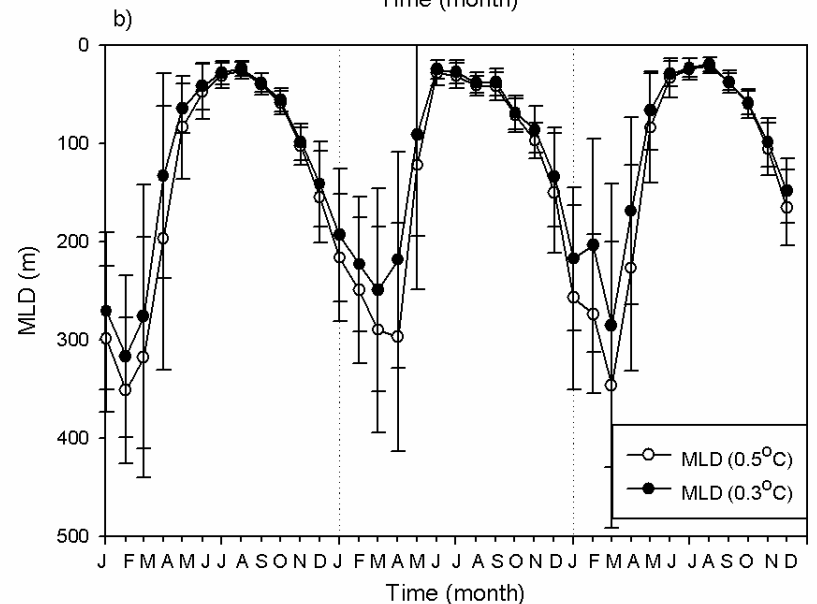


Hartman et al., 2010

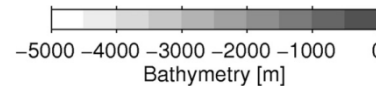
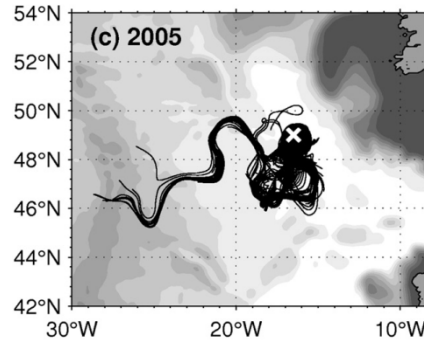
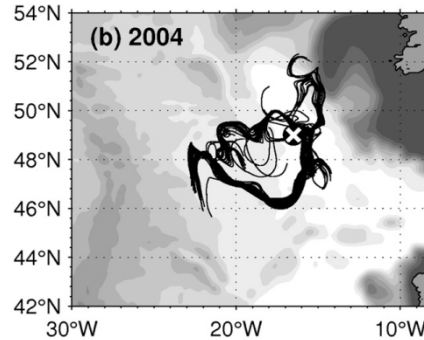
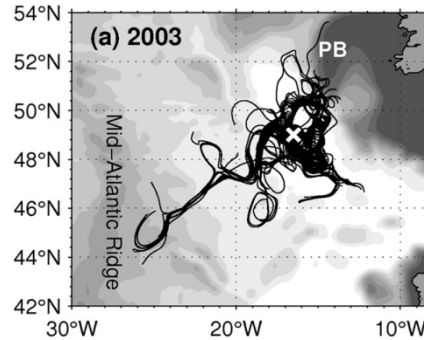
Nitrate



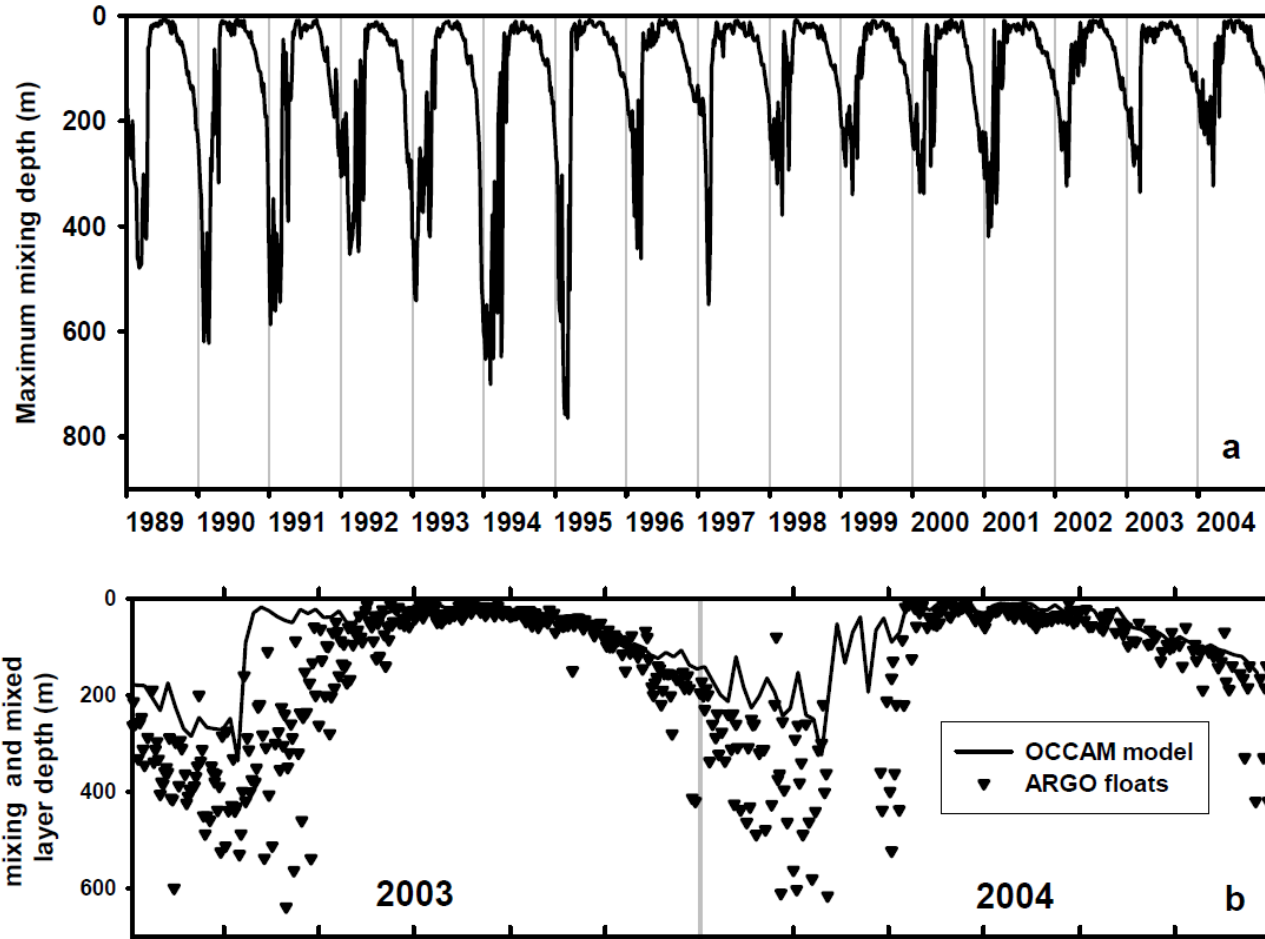
Mixed Layer Depth



Progressive changes 2003-2005 Variability in source waters

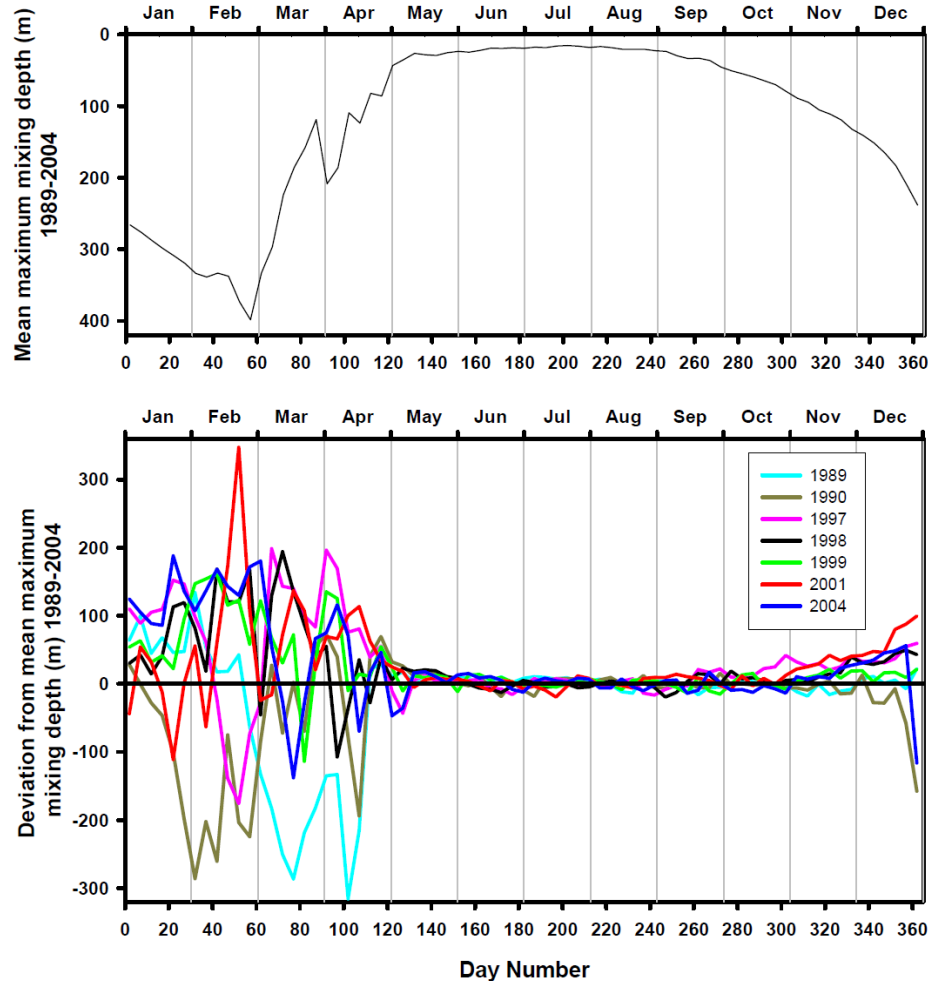


Mixed Layer Depth



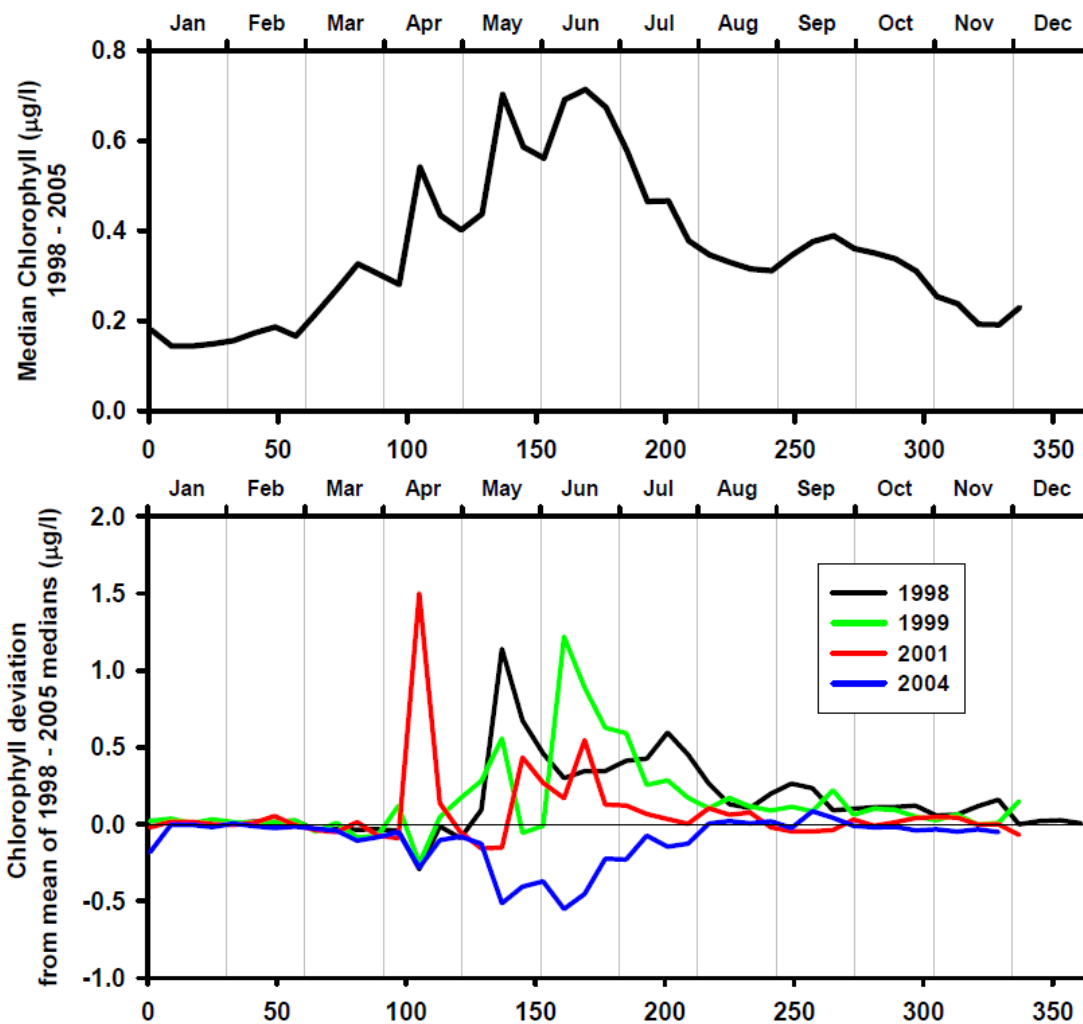
Maximum mixing depth derived from the $\frac{1}{4}$ degree OCCAM model in a circle around the PAP site with radius of 200km.(b)Expansion of data in part a(above) with the addition of plus data derived from ARGO floats which became sufficiently abundant after 2002.

Mixed Layer Depth



Mean maximum mixing depth determined from the 200km radius area around PAP by the $\frac{1}{4}$ degree OCCAM model. Lower: Deviation from the mean mixing depth for each of the 7 years for which there are adequate particle flux data.

Chl-a



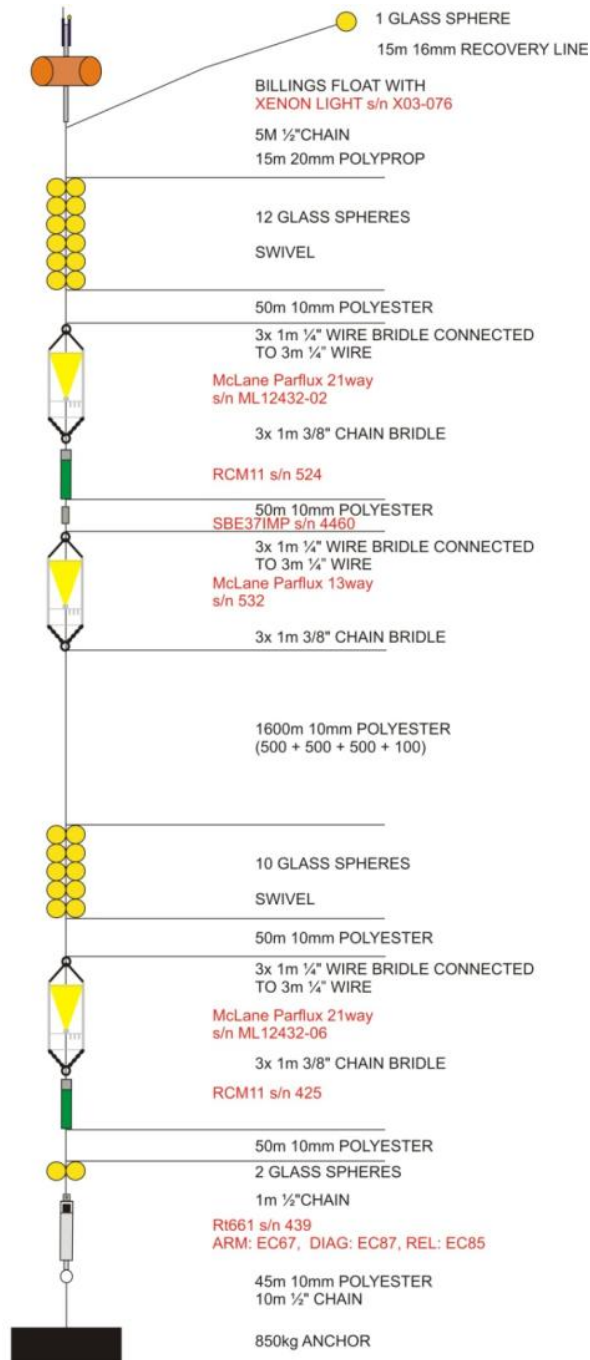
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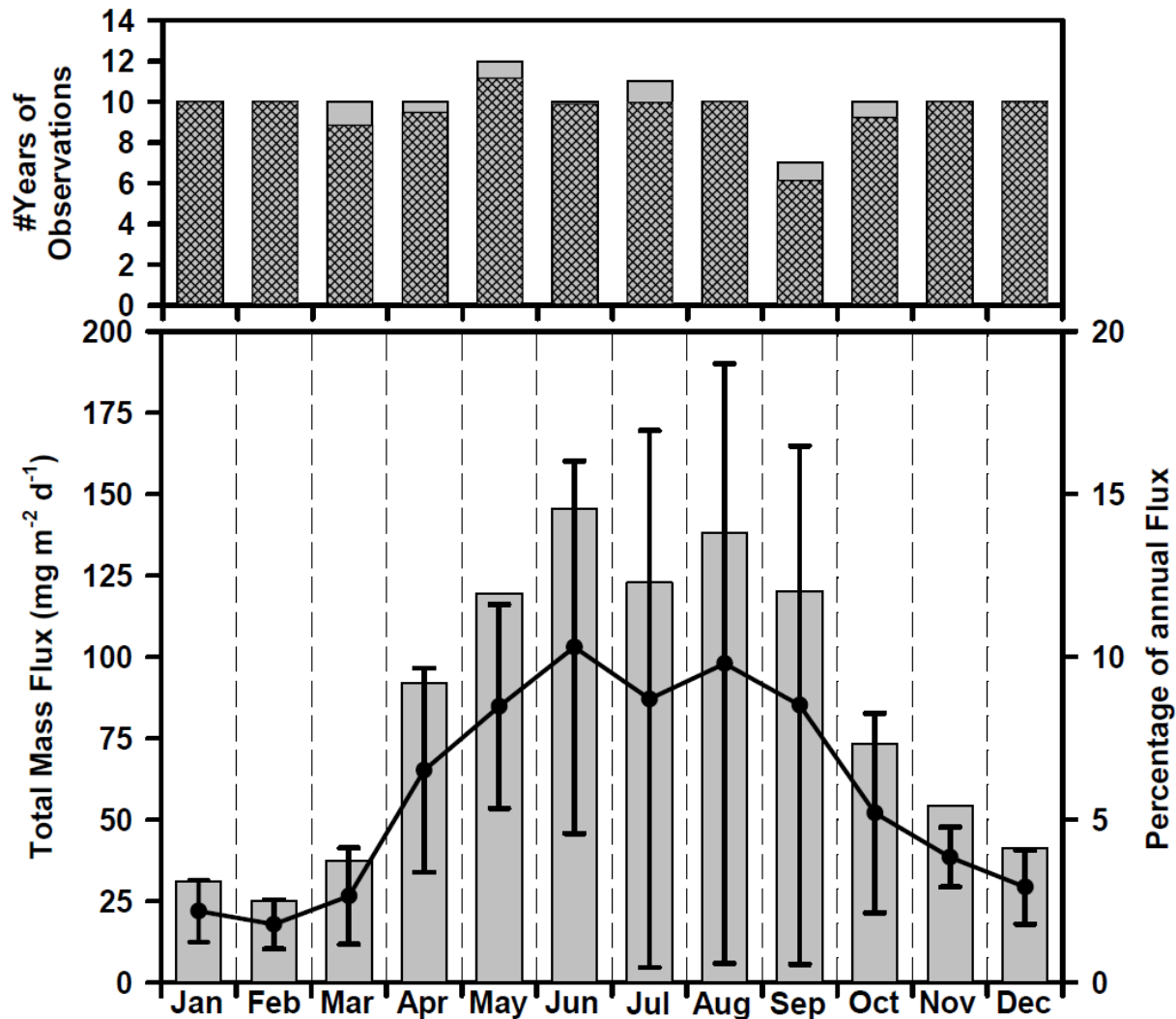
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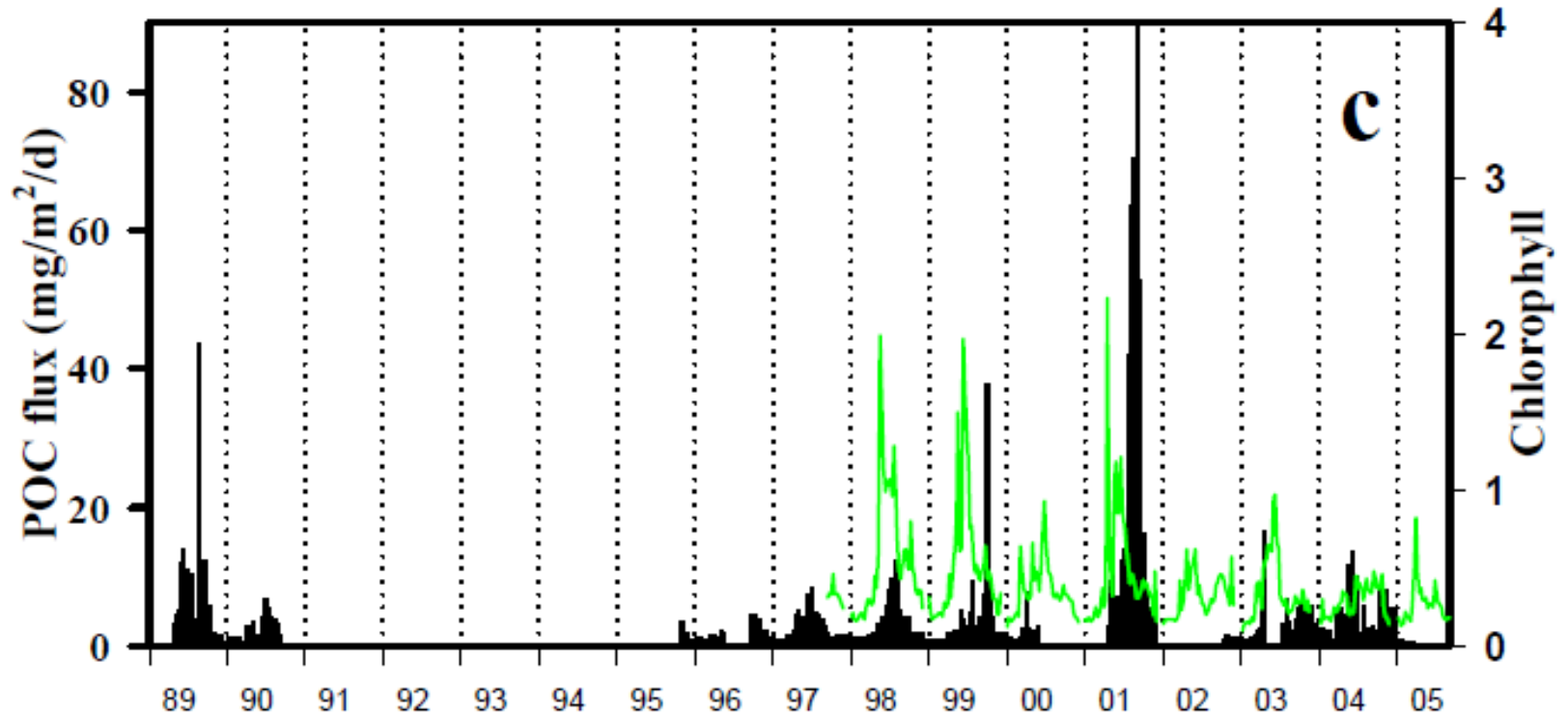
WATER DEPTH 4814m





Seasonal cycle of mass flux at 3000 m based on averages of all measurements made between 1989 and 2005. In top panel bars represent the number of years for which there is greater than 15 days of flux data per month. The hatched area is the coverage co-efficient. In the bottom panel grey bars represent the percentage of annual flux each month contributes. Line plot is the average total mass flux for each individual month with error bars representing ± 1 standard deviation.

What are the levels and long term trends in particle flux?

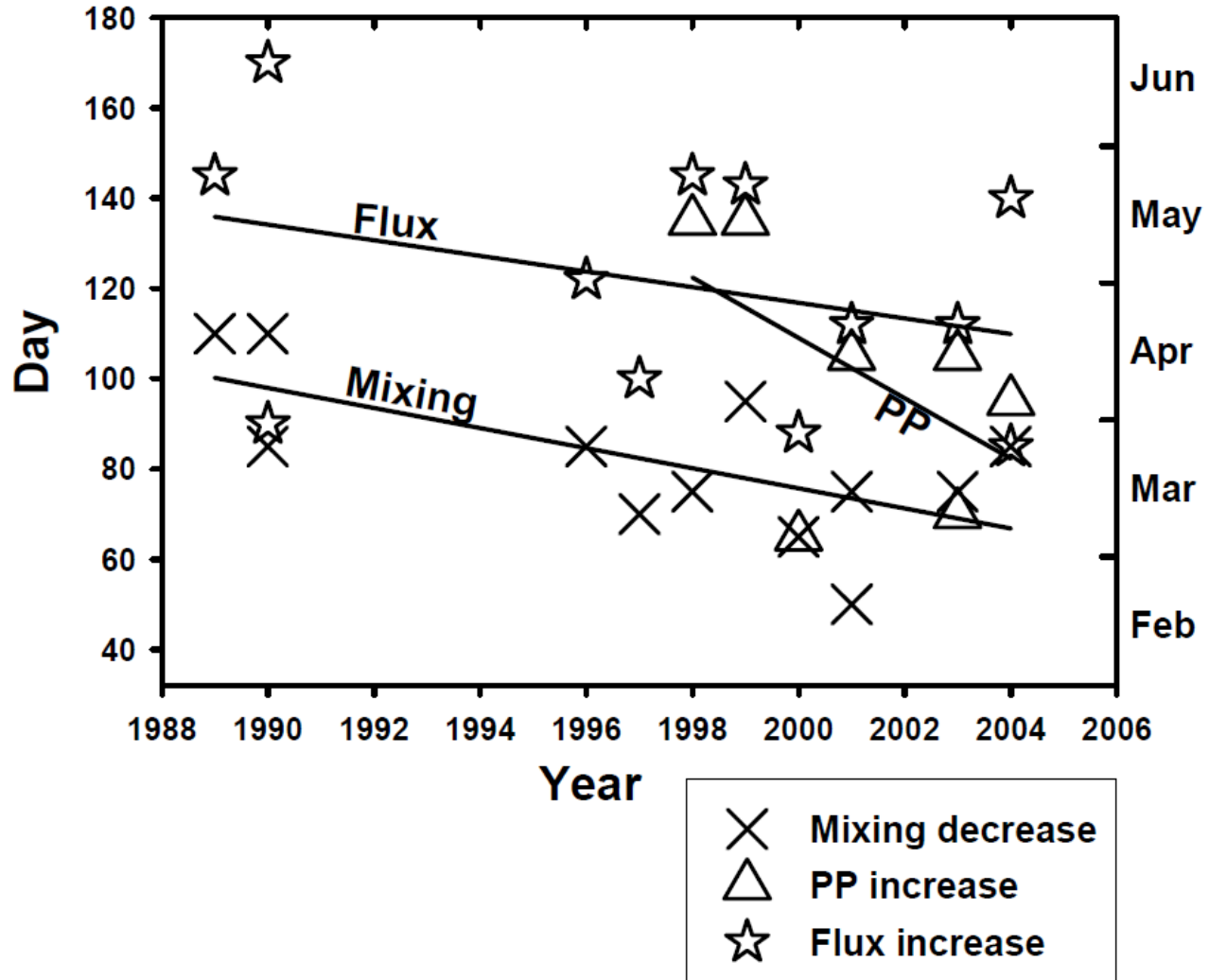


Particulate downward flux at 3000 m depth in terms of

(A) dry weight, (B) biogenic silica, (C) organic carbon and (D) inorganic carbon.

(B) Also shown (green) are the surface chlorophyll concentrations averaged over a circle of 200 km radius around the PAP site.

Phenology



Are spring-time changes at the PAP site as reflected in the time at which mixing decreases, productivity increases and deep ocean flux increases?

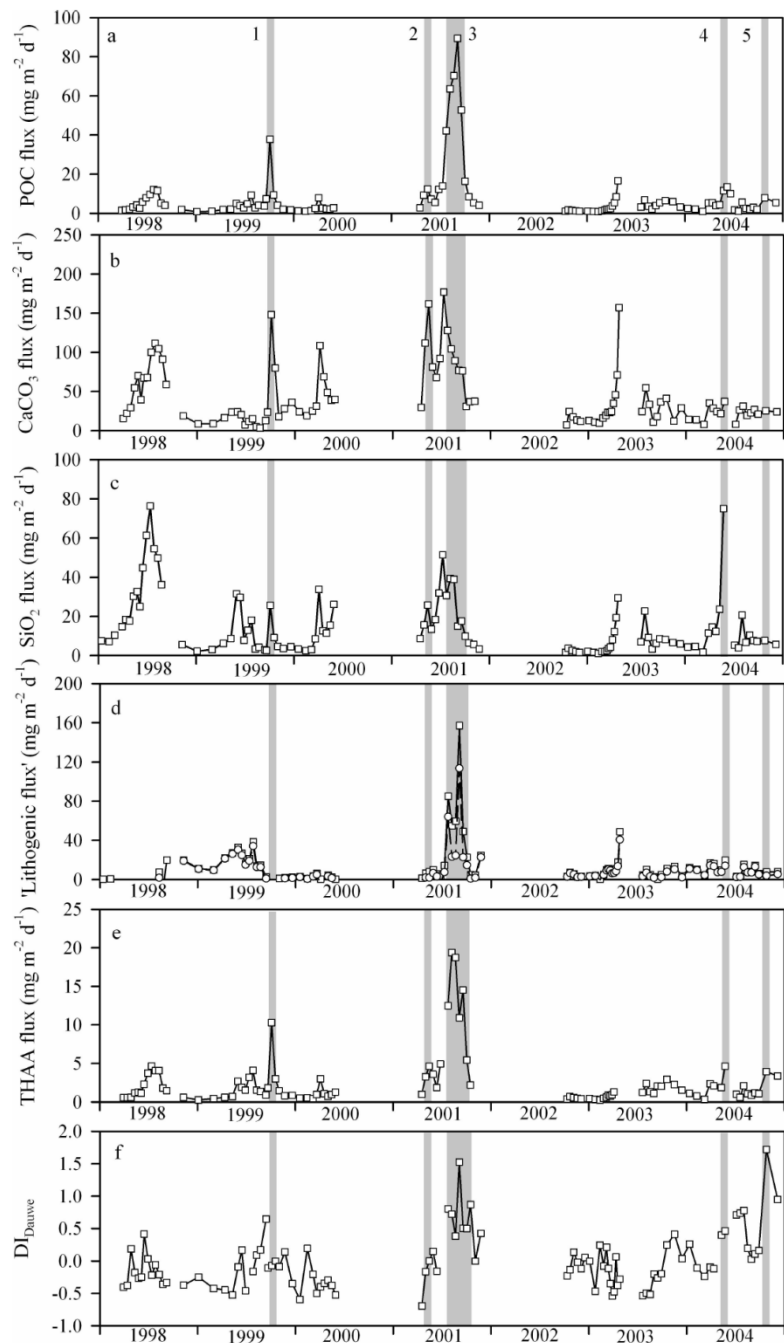
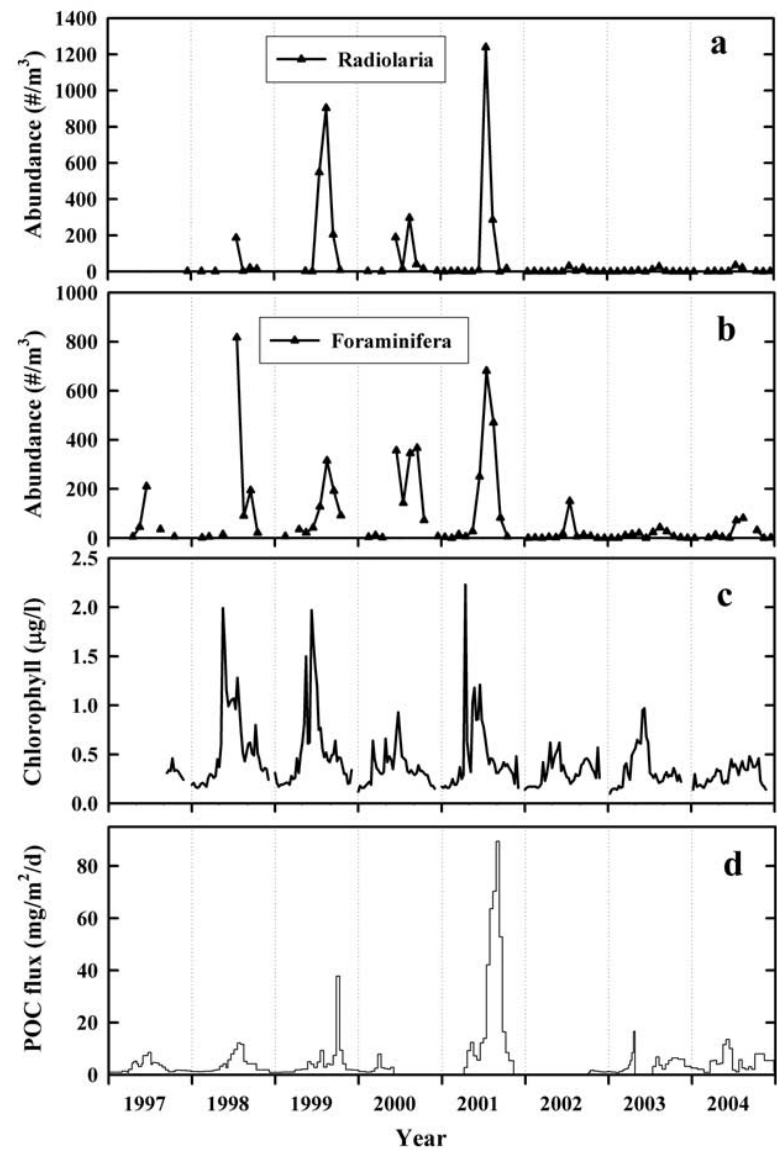


Fig. 2. Interannual variability in the fluxes of (a) POC, (b) CaCO_3 , (c) SiO_2 , (d) lithogenic matter, (e) total hydrolysable amino acids (THAA), and (f) calculated degradation index. The lithogenic flux (a) has been calculated according to Eq. 1 using POC to POM scaling factors (S) of



Temporal variation in (a) abundance of radiolarian and (b) abundance of foraminifera, (c) mean chlorophyll concentration around the PAP site, and (d) POC flux at 3000 m.

Faecal Pellet Fluxes

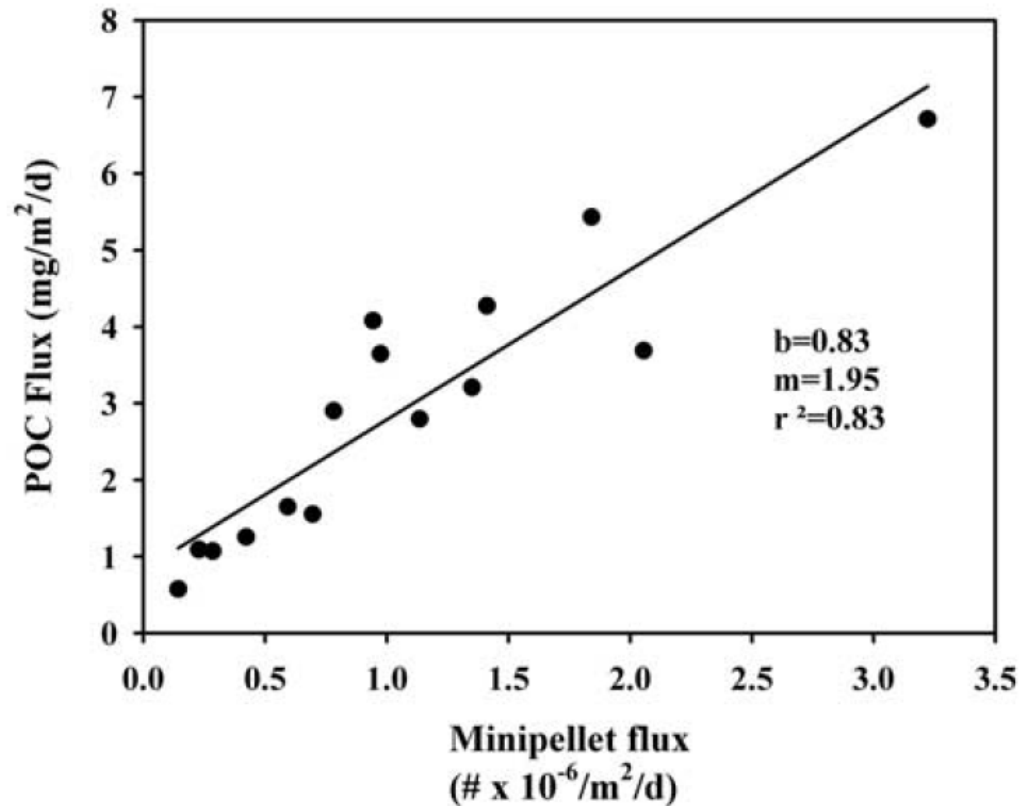
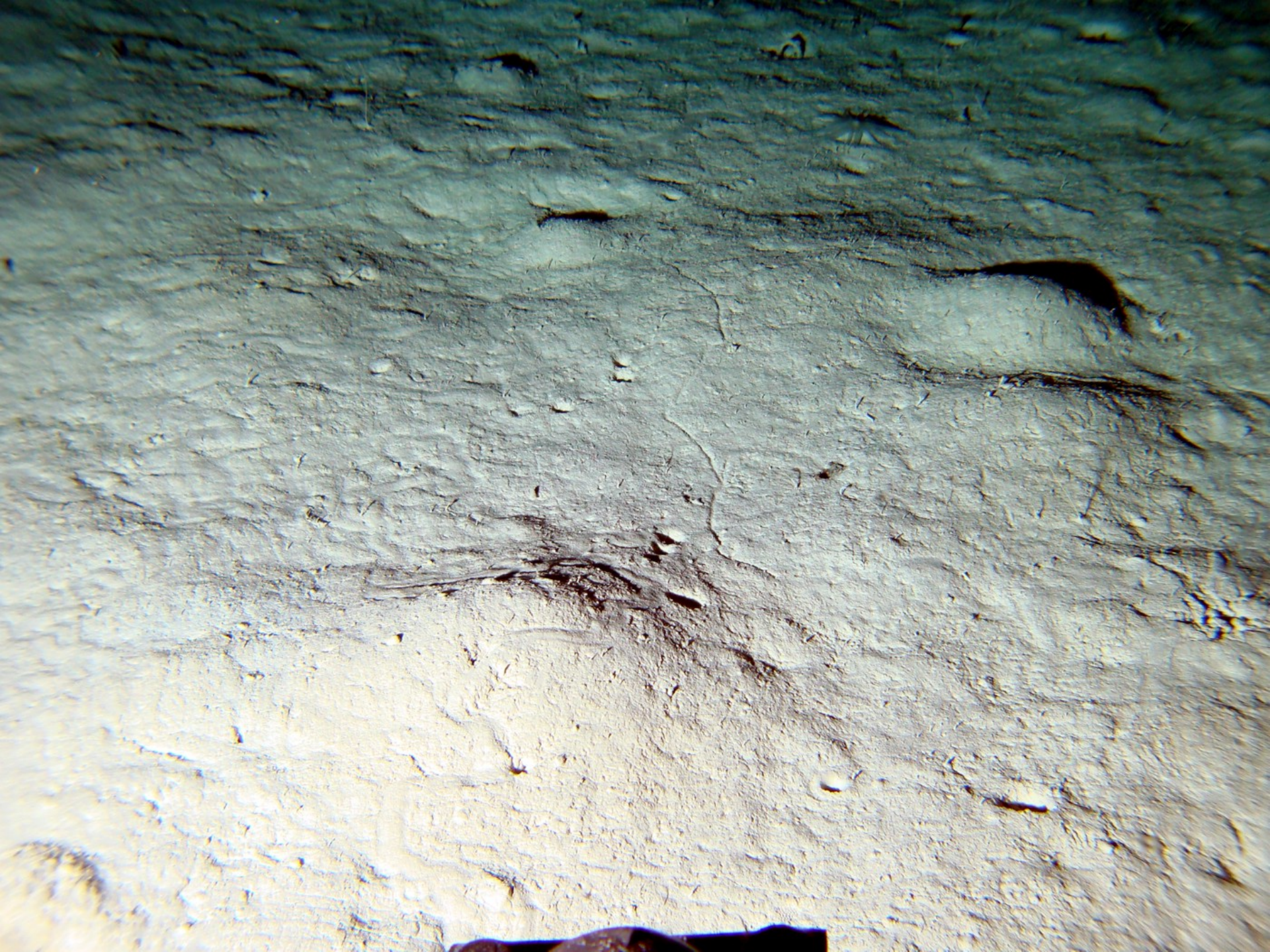
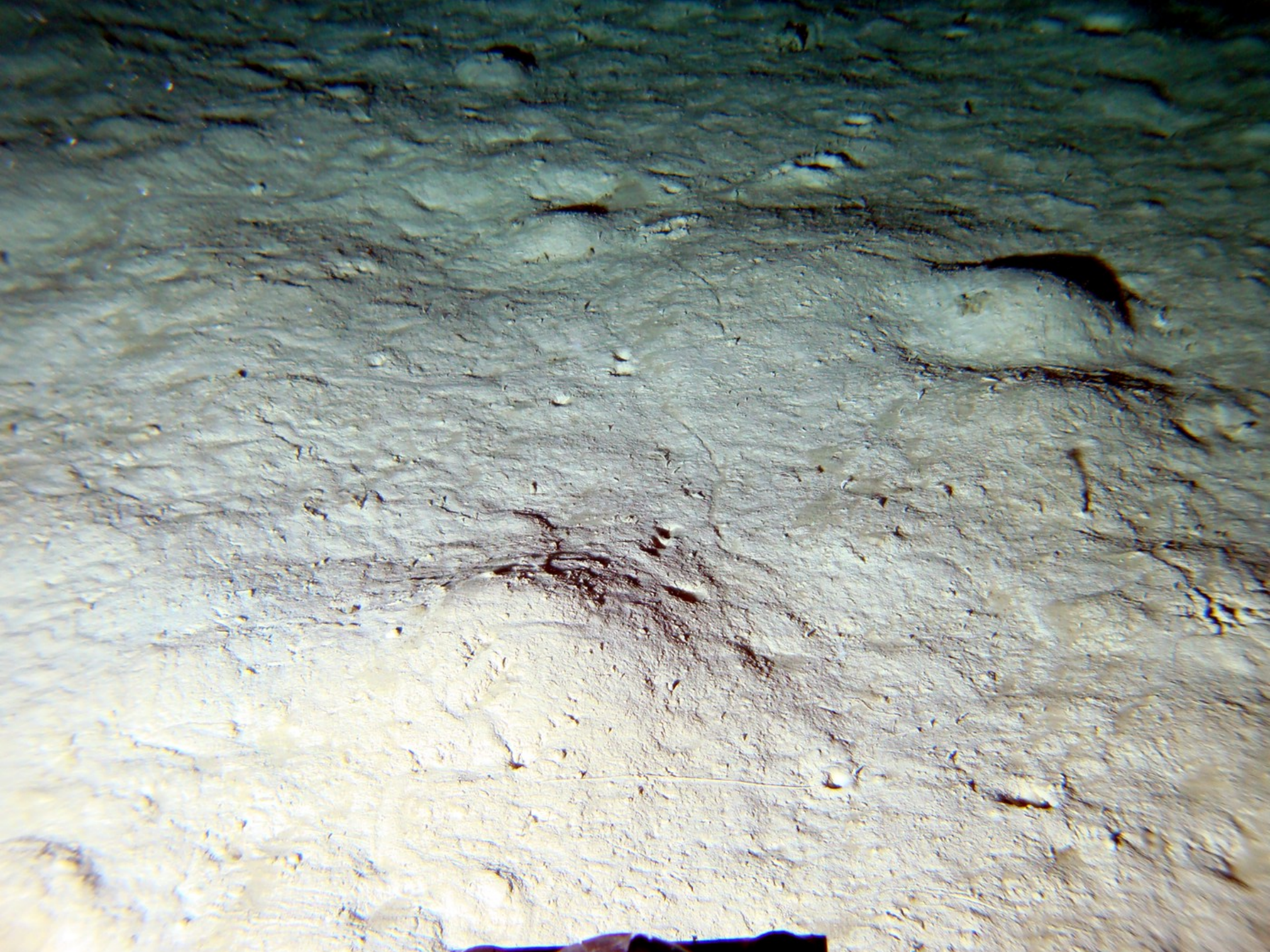
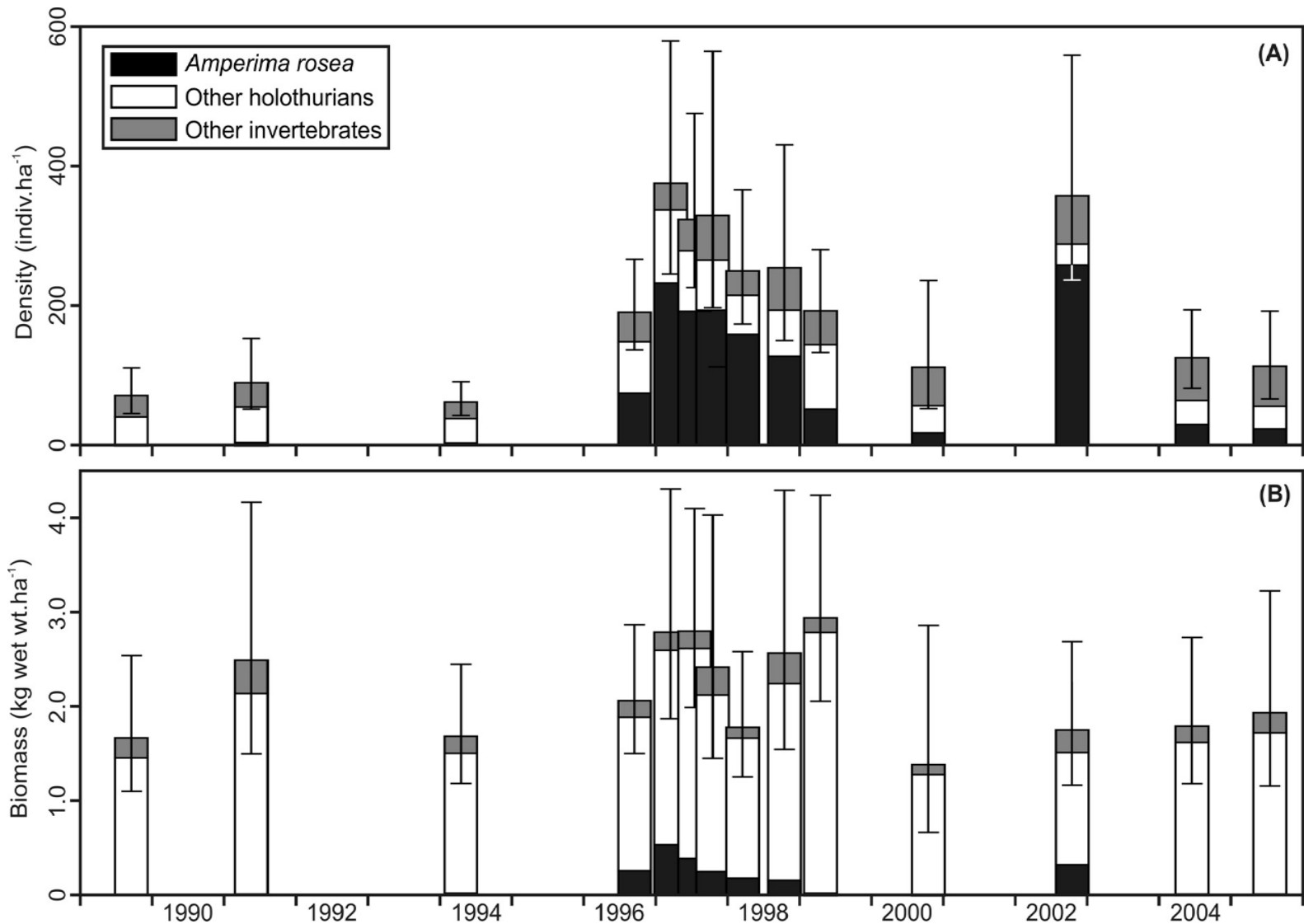


Figure 6. Relationship between downward flux of mini-pellets and POC during 1990 at 3000 m depth.



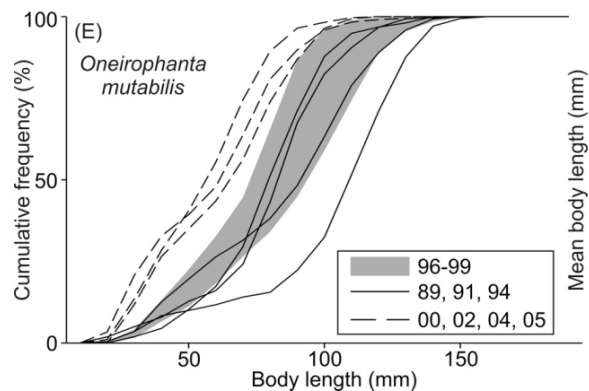
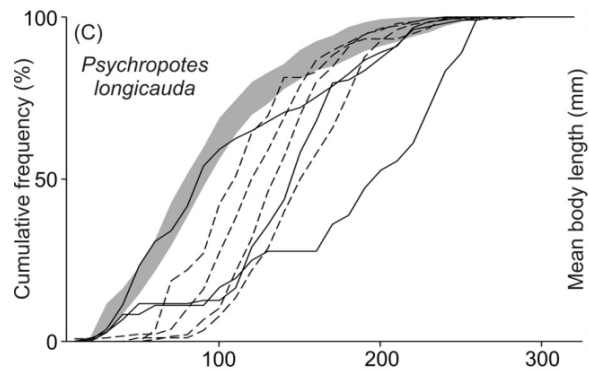
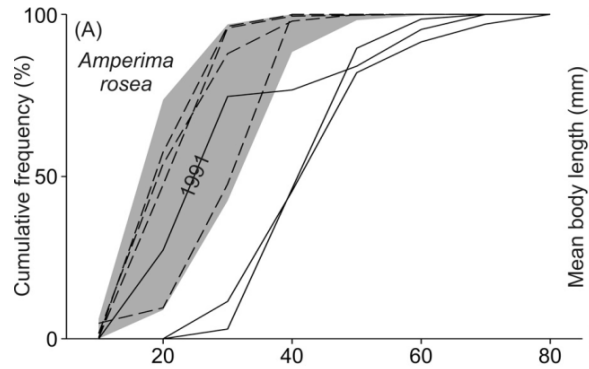






(A) density and (B) biomass of 8 selected megafaunal groups at the Porcupine Abyssal Plain Sustained Observatory site 9 (means and 95% confidence intervals are shown).

Size Distribution Variation



Cumulative frequency
distributions of body length;

Size Distribution Variation & Respiration

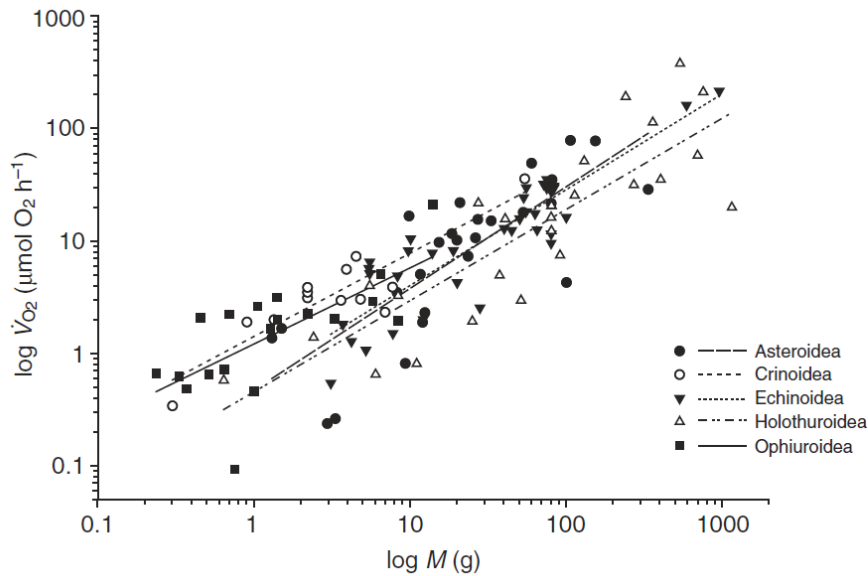
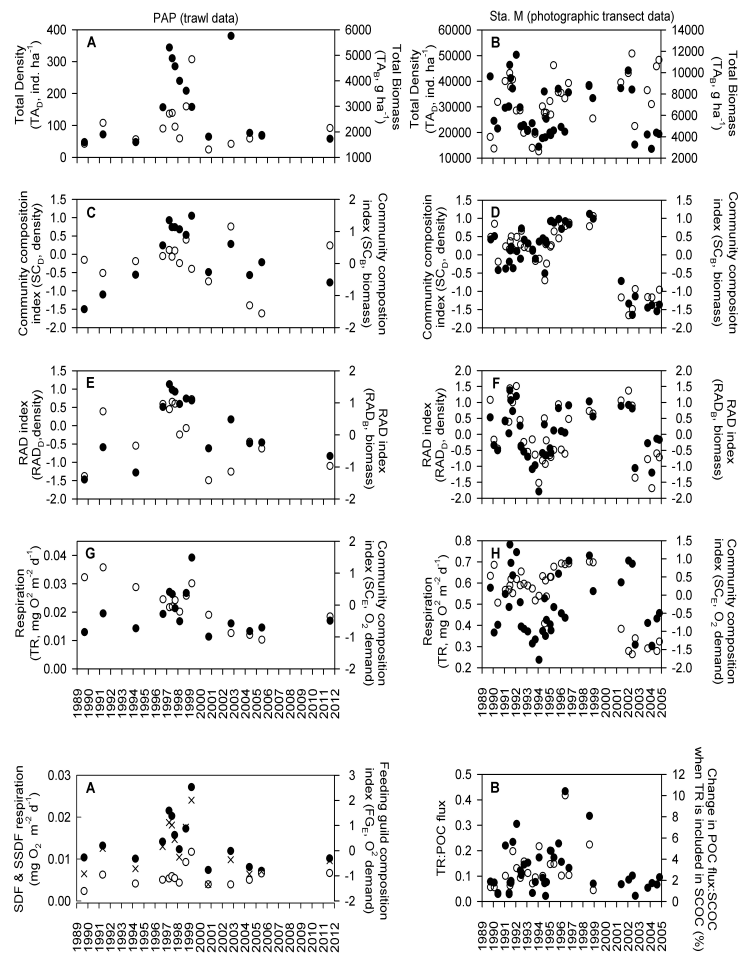


Fig. 2. Metabolic rates (\dot{V}_{O_2}) of the five echinoderm classes as a function of wet mass (M) as collated in the final echinoderm metabolic rate data set (see text). Scaling relationships are in the form $\dot{V}_{O_2}=aM^b$ where a is a normalisation constant and b is a scaling coefficient representing the slope of the relationship between \dot{V}_{O_2} and M . The metabolic rates of all five echinoderm classes are highly significantly correlated with mass ($P<0.001$): Asteroidea ($0.472M^{0.90}$, $R^2=0.636$), Crinoidea ($1.414M^{0.74}$, $R^2=0.792$), Echinoidea ($0.564M^{0.85}$, $R^2=0.796$), Holothuroidea ($0.456M^{0.81}$, $R^2=0.730$), Ophiuroidea ($1.215M^{0.68}$, $R^2=0.481$). All metabolic rates have been normalised to 12°C using a Q_{10} adjustment of 2.15. Sources of all data can be found in supplementary material Table S1.

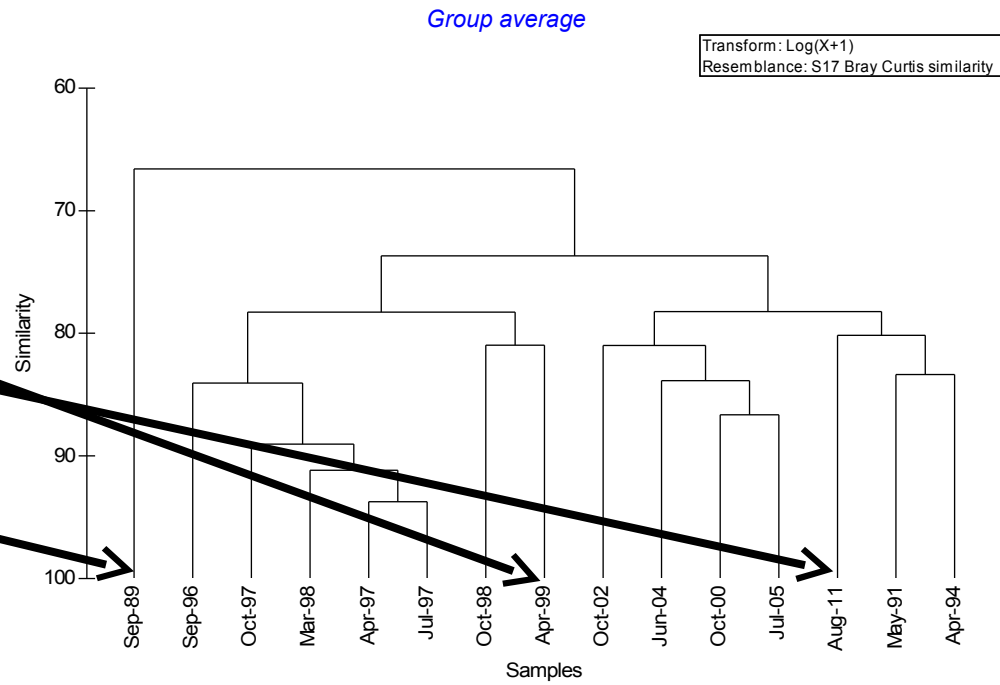
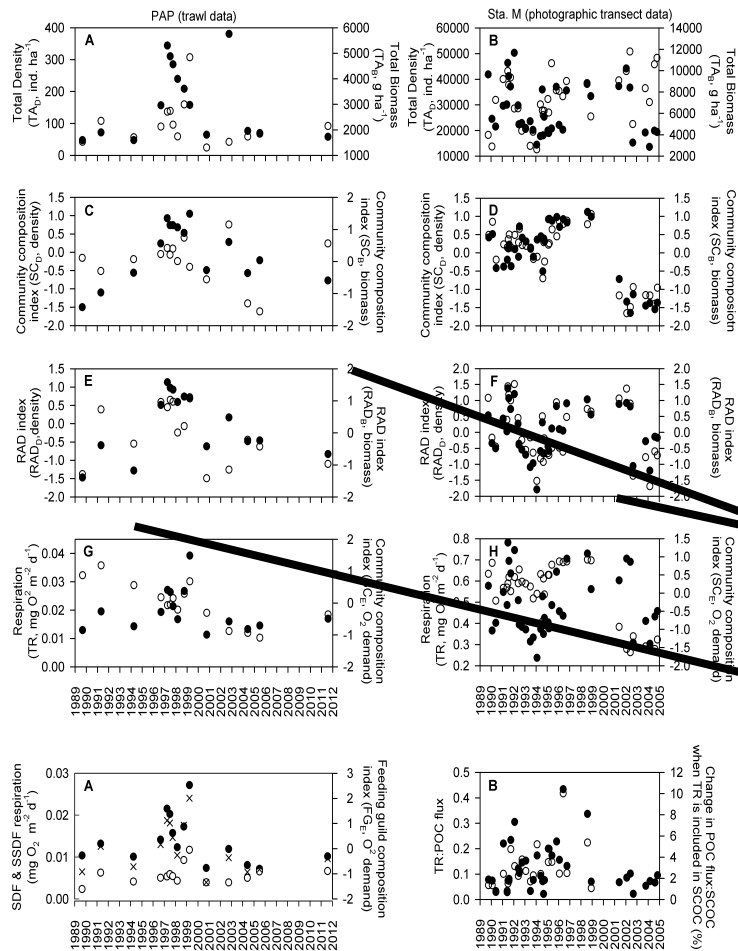
We then took our avg. body size data for each taxon and converted that to individual respiration rates.

We then multiplied the individual respiration rates by animal density for all the dominant taxa to get community respiration rates.

Which then gave us...

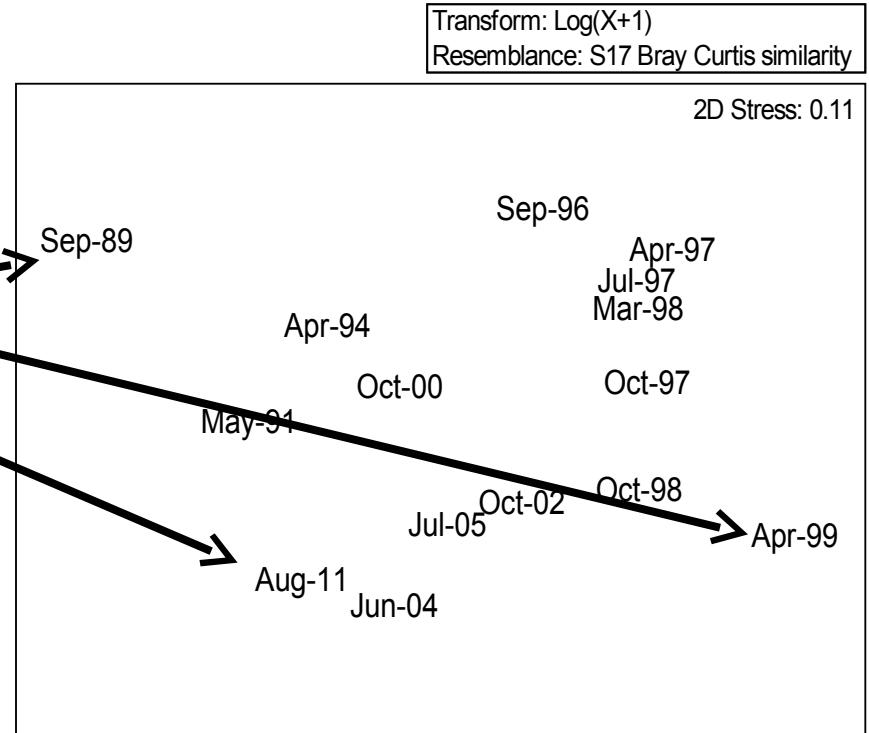
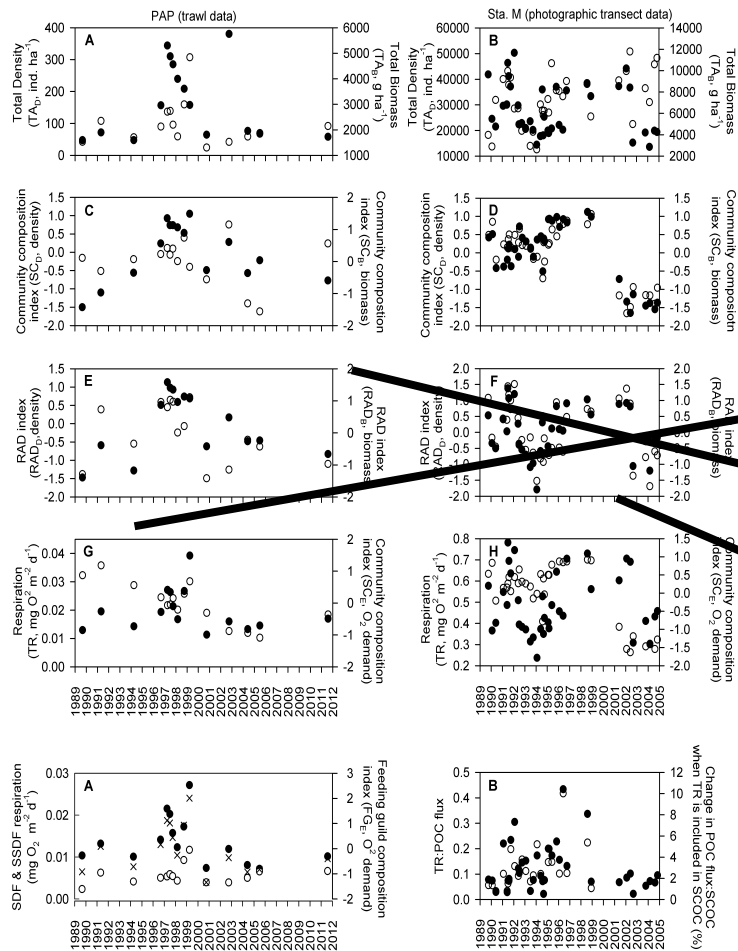


Time series of echinoderm megafauna community dynamics for the PAP and Sta. M research sites including A and B) density (\bullet , TA_{D_i}), biomass (\circ , TA_{B_i}); C and D) indices of species composition similarity based on density data (\bullet , SC_{D_i}) and biomass (\circ , SC_{B_i}); E and F) indices of rank abundance distribution similarity based on density (\bullet , RAD_{D_i}) and biomass (\circ , RAD_{B_i}); and G and H) total respiration of the studied megafauna (\bullet , TR) and an index of species composition similarity based on energetic demand (\circ , SC_{E_i}).



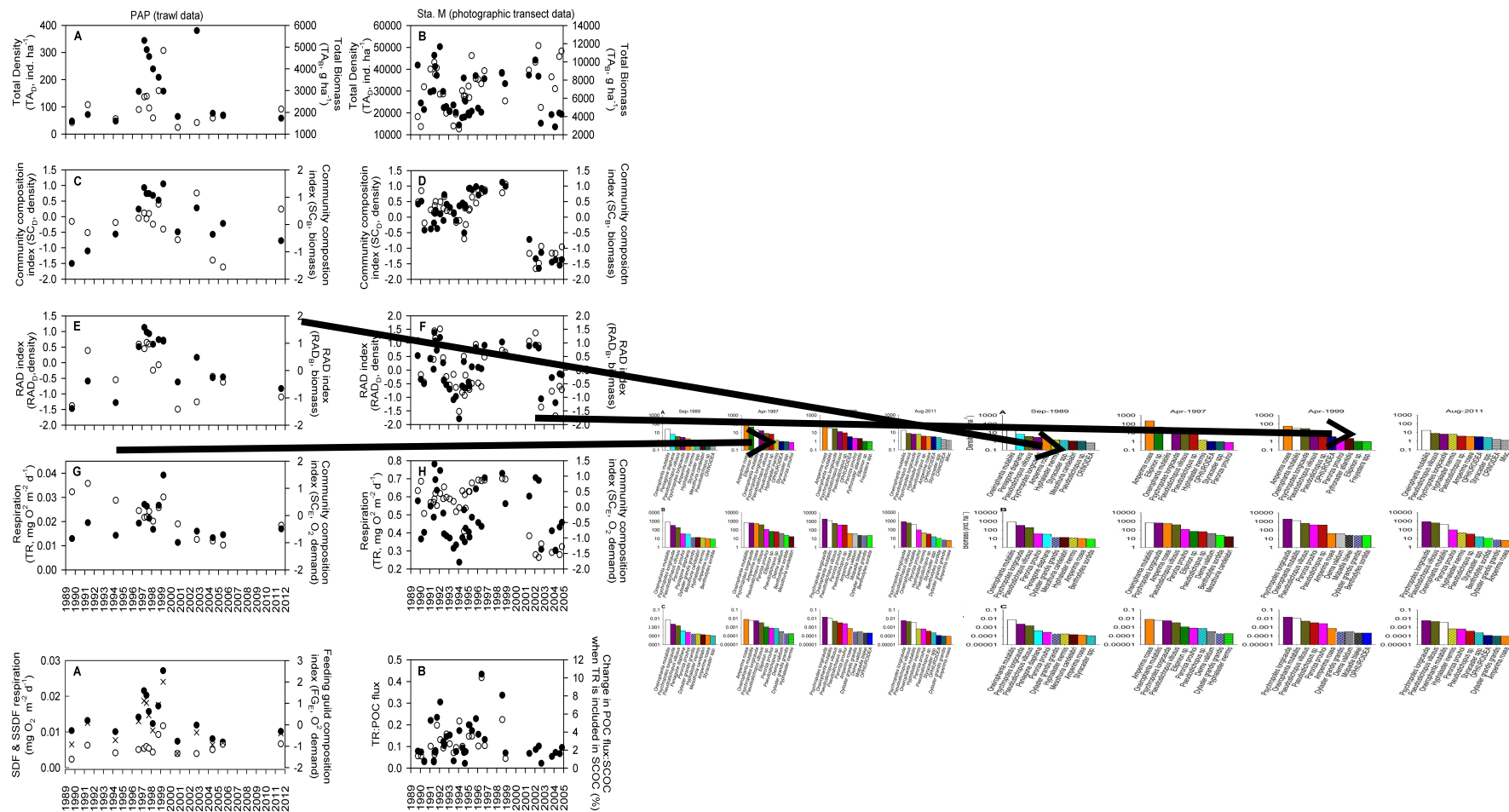
Cluster diagram of community composition (density)

Time series of echinoderm megafauna community dynamics for the PAP including A) density (\bullet , TA_D), biomass (\circ , TA_B); C) indices of species composition similarity based on density data (\bullet , SC_D) and biomass (\circ , SC_B); E) indices of rank abundance distribution similarity based on density (\bullet , RAD_D) and biomass (\circ , RAD_B); and G) total respiration of the studied megafauna (\bullet , TR) and an index of species composition similarity based on energetic demand (\circ , SC_E).



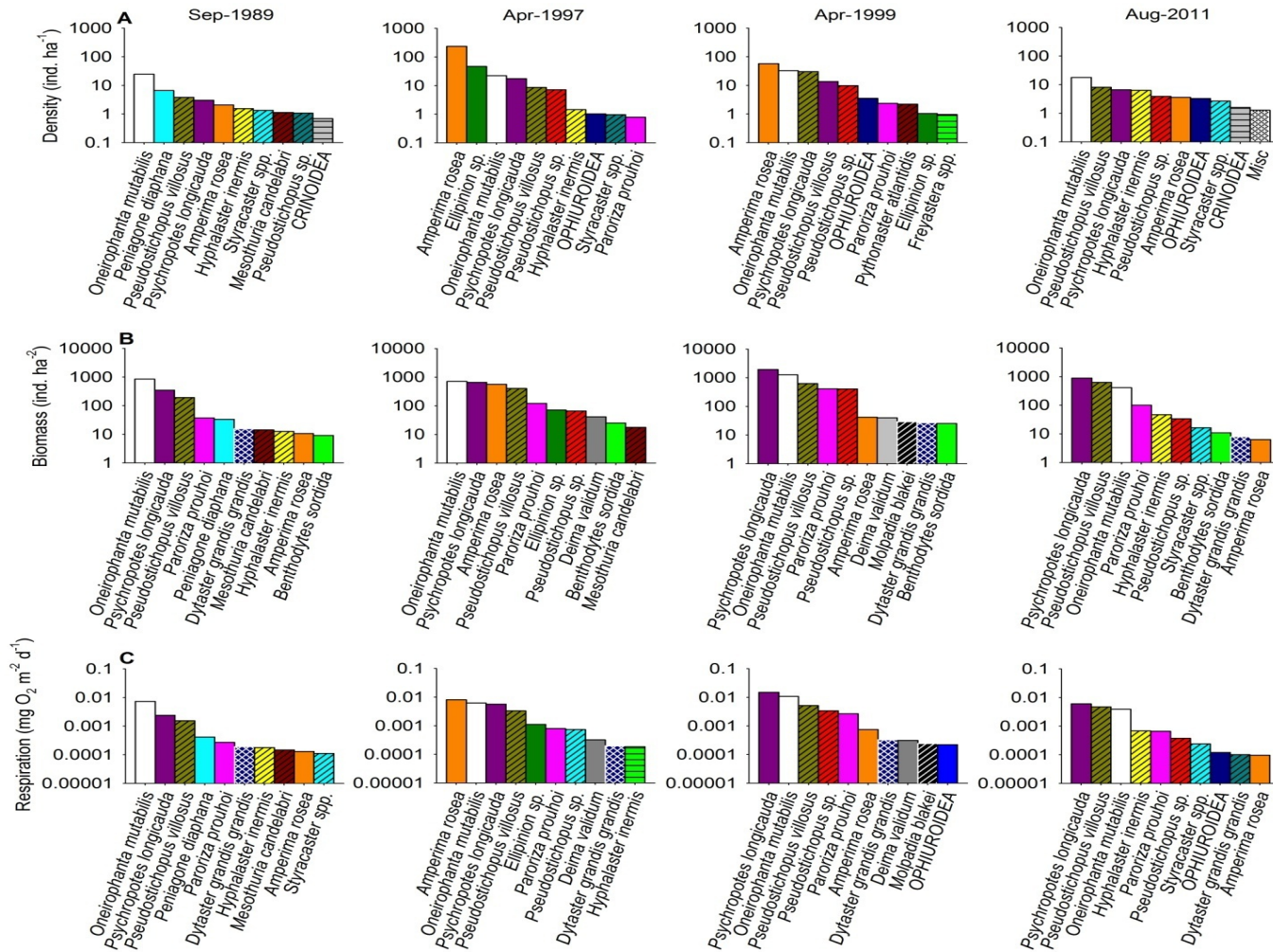
MDS plot of community composition (density), x-axis used as y axis for time series

Time series of echinoderm megafauna community dynamics for the PAP and Sta. M research sites including A and B) density (\bullet , TA_{D_B}), biomass (\circ , TA_{B_B}); C and D) indices of species composition similarity based on density data (\bullet , SC_{D_B}) and biomass (\circ , SC_{B_B}); E and F) indices of rank abundance distribution similarity based on density (\bullet , RAD_{D_B}) and biomass (\circ , RAD_{B_B}); and G and H) total respiration of the studied megafauna (\bullet , TR) and an index of species composition similarity based on energetic demand (\circ , SC_{E_D}).



Rank abundance distribution plots

Time series of echinoderm megafauna community dynamics for the PAP and Sta. M research sites including A and B) density (\bullet , TA_{D_i}), biomass (\circ , TA_{B_i}); C and D) indices of species composition similarity based on density data (\bullet , SC_{D_i}) and biomass (\circ , SC_{B_i}); E and F) indices of rank abundance distribution similarity based on density (\bullet , RAD_{D_i}) and biomass (\circ , RAD_{B_i}); and G and H) total respiration of the studied megafauna (\bullet , TR) and an index of species composition similarity based on energetic demand (\circ , SC_{E_i}).

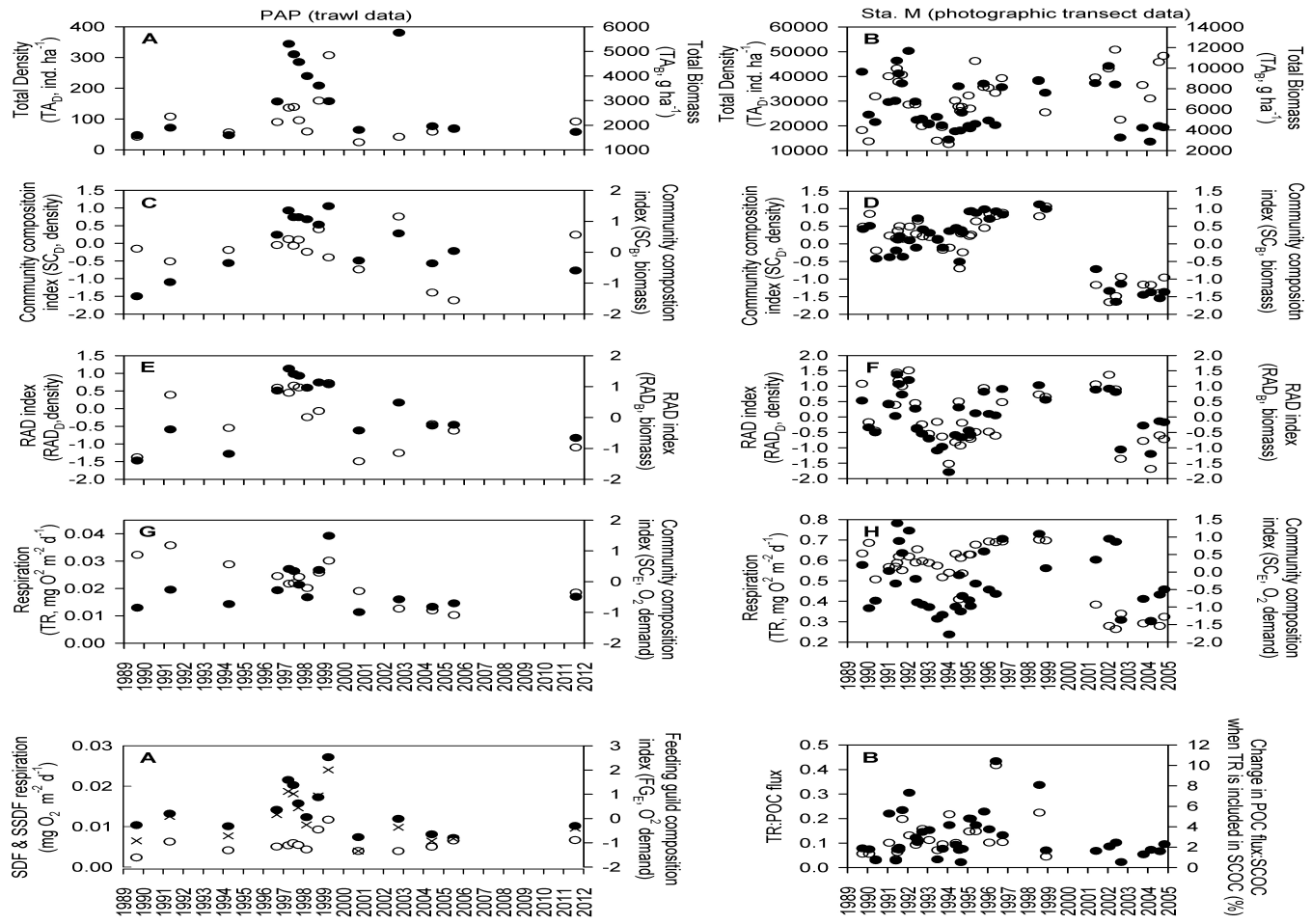


Rank abundance distribution plots for PAP data for selected times during the time series based on:

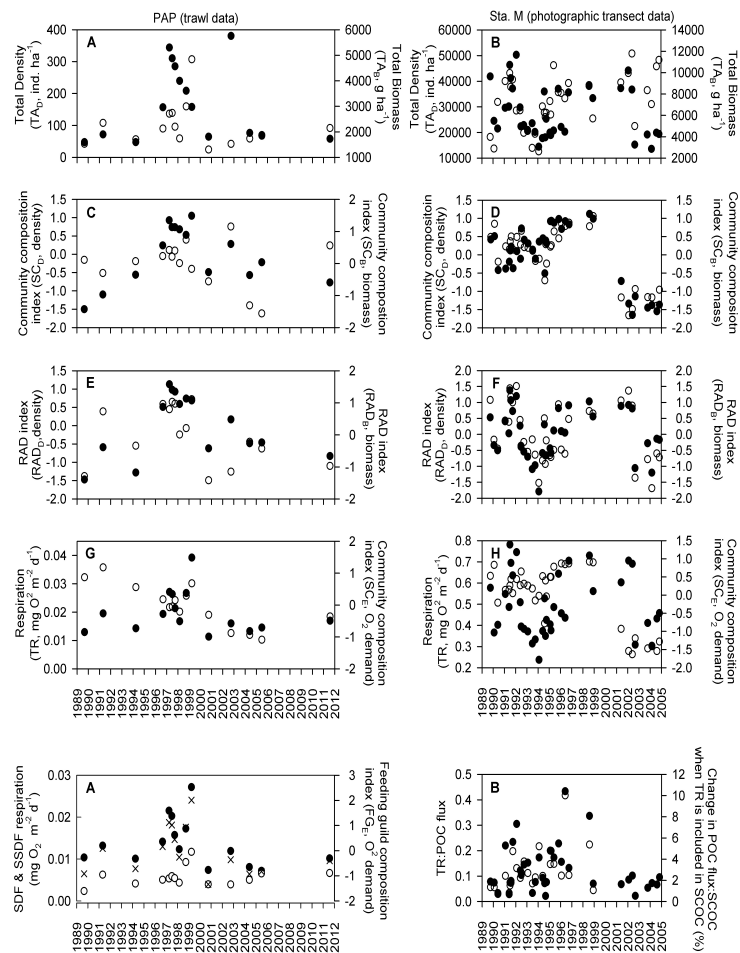
- A) density;
- B) biomass; and
- C) energetic demand (respiration),

Illustrates both changes in distribution shape, but also the ranks of the top ten most dominant fauna over time.

The SDF are open bars, SSDF diagonally hatched bars, SF have horizontally hatched bars, and P/S are cross hatched.



Time series of echinoderm megafauna community dynamics for the PAP and Sta. M research sites including A and B) density (\bullet , TA_D), biomass (\circ , TA_B); C and D) indices of species composition similarity based on density data (\bullet , SC_D) and biomass (\circ , SC_B); E and F) indices of rank abundance distribution similarity based on density (\bullet , RAD_D) and biomass (\circ , RAD_B); and G and H) total respiration of the studied megafauna (\bullet , TR) and an index of species composition similarity based on energetic demand (\circ , SC_E).



Community descriptor	Sta. M (n=37)		PAP (n=15)	
	TR	<i>p</i>	TR	<i>p</i>
univariate	r_s		r_s	
TA_D	0.63	<0.001	0.61	0.02
TA_B	0.87	<0.001	0.94	<0.001
multivariate	R	<i>p</i>	R	<i>p</i>
SC_D	0.09	0.053	0.34	0.007
SC_B	0.31	<0.001	0.20	0.07
SC_E	0.27	<0.001	0.42	0.02
RAD_D	0.25	<0.001	0.37	0.004
RAD_B	0.70	<0.001	0.49	<0.001
RAD_E	0.75	<0.001	0.78	<0.001

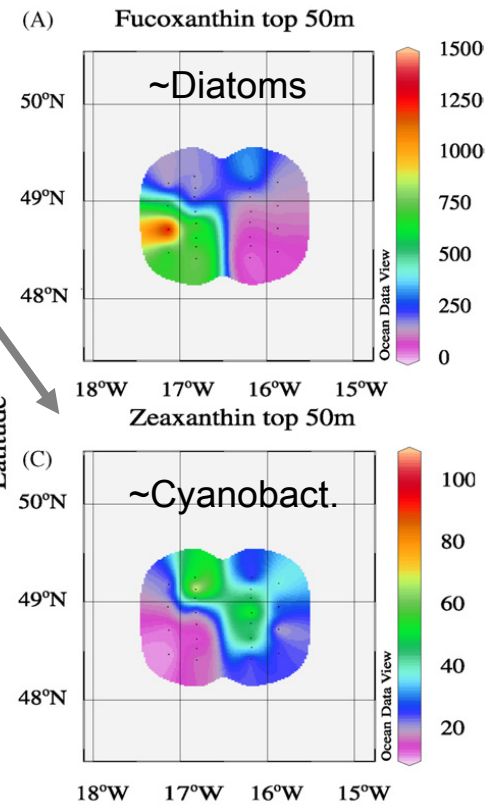
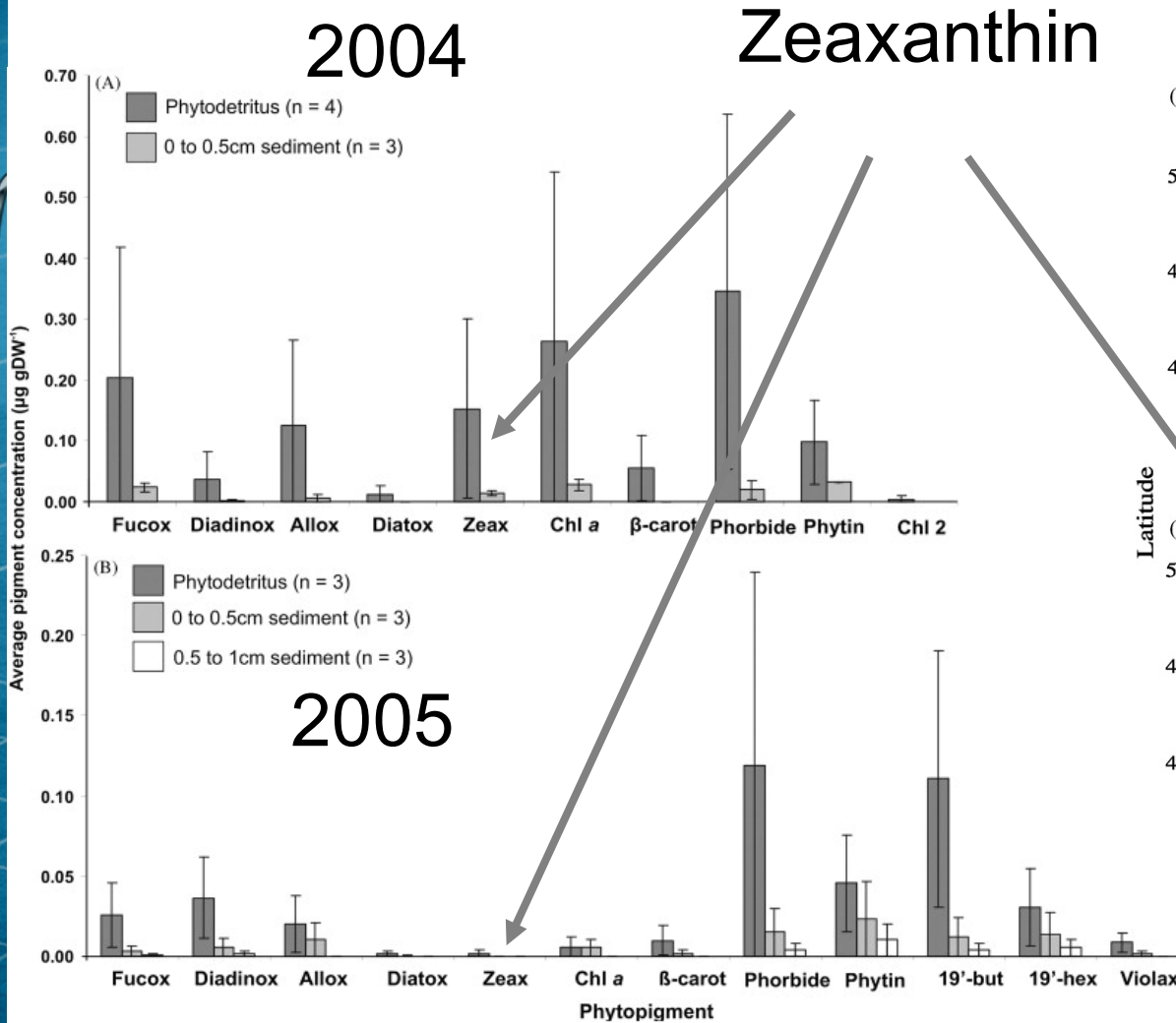
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Are pigments related to change?



Logonna-Daoulas (France)

17-22 sept. 2012



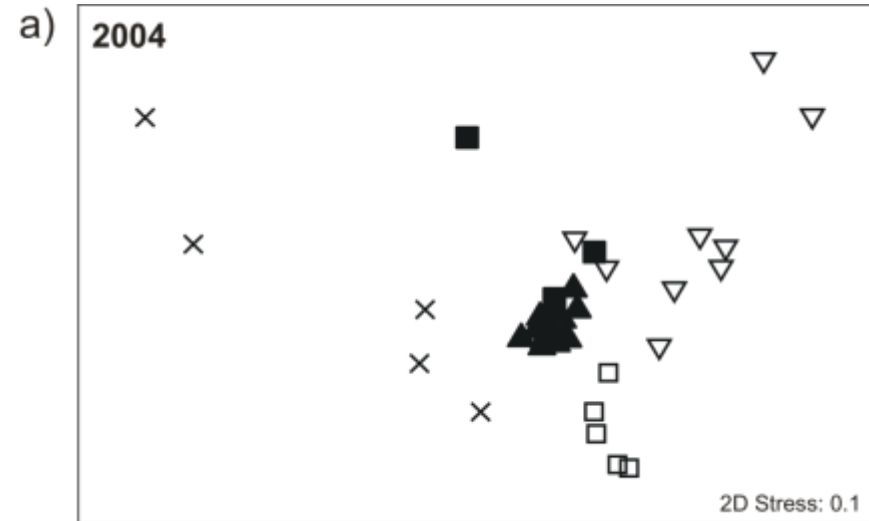
Fitz-George Balfour et al., 2010

Are pigments related to change?

MDS ordination of 37 individual holothurian ovary samples from PAP June 29 2004 (a) and July 2005 (b), based on $\sqrt{\cdot}$ -transformed pigment percentage contributions 30 and Bray-Curtis similarities.

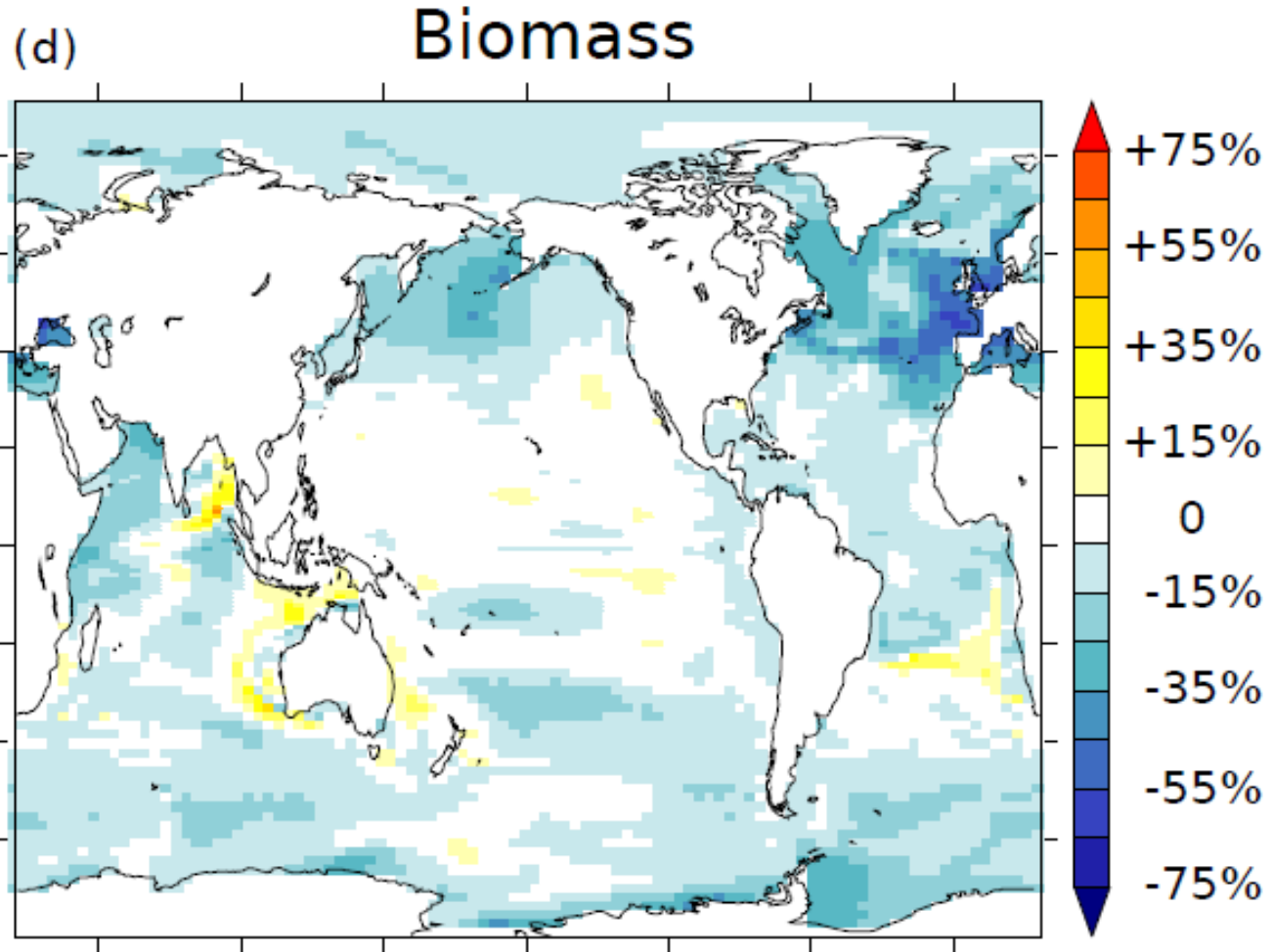
Key:

- ▲ = *Amperima rosea*;
- d = *Oneirophanta mutabilis*
- = *Peniagone diaphana*
- = *Psychropotes longicauda*
- X = *Paroriza prouhoi*



T. Smith, Billett, Wolff, Thompson,
and Tyler, *Deep-Sea Res.*, 2010

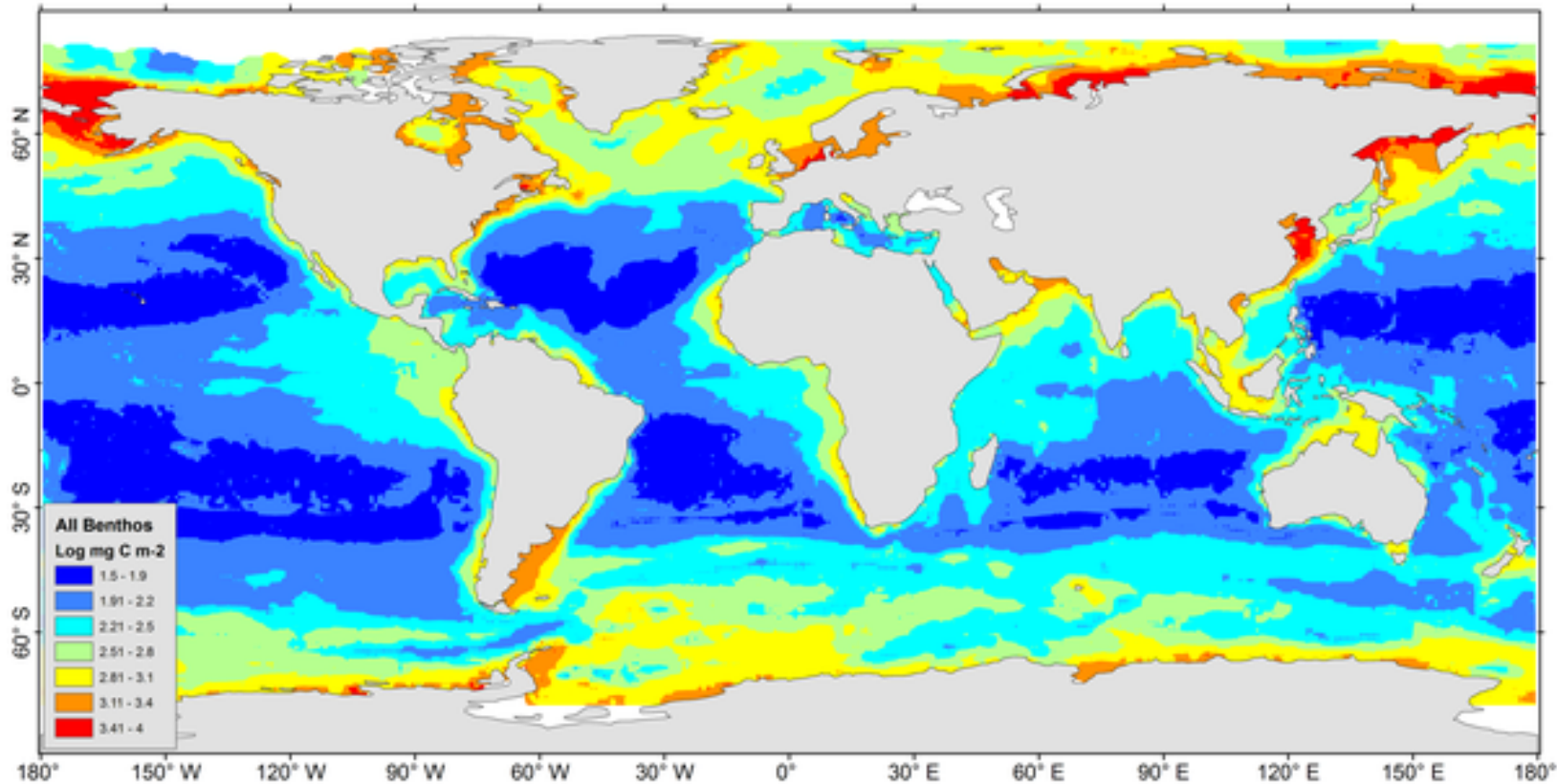
Biogeochemical Models Suggest Major Change Ahead



Change in biomass pool from preindustrial to ~2100
(IPCC SERS A2; function of nutrients; Steinacher et al. 2010)

Global Estimates of Seafloor Biomass

Time-series
analysis in
Marine
science and
applications
for industry



Logonna-
Daoulas
(France)

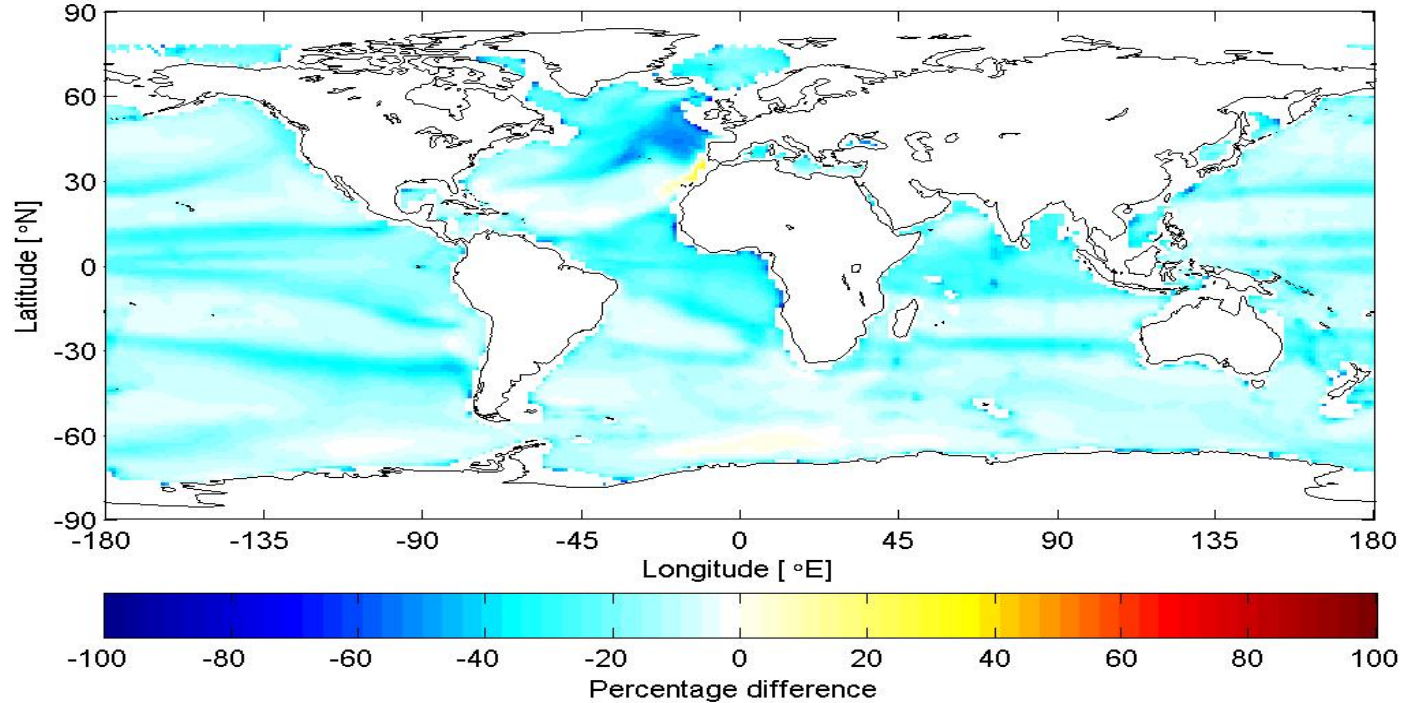
Established relationships between various parameters, including POC flux vs. Seafloor biomass
Wei et al., PLoS One, 2010

17-22 sept.
2012

Global Estimates of Seafloor Biomass into the Future

Time-series analysis in Marine science and applications for industry

% diff. in total biomass calc. from flux (Martin on EP100) to 500 m above seafloor between 1980-2009 and 2090-2099 IPSL-CM4 HISTA2



Logonna-Daoulas
(France)

17-22 sept.
2012

Jones, Ruhl et al., in prep

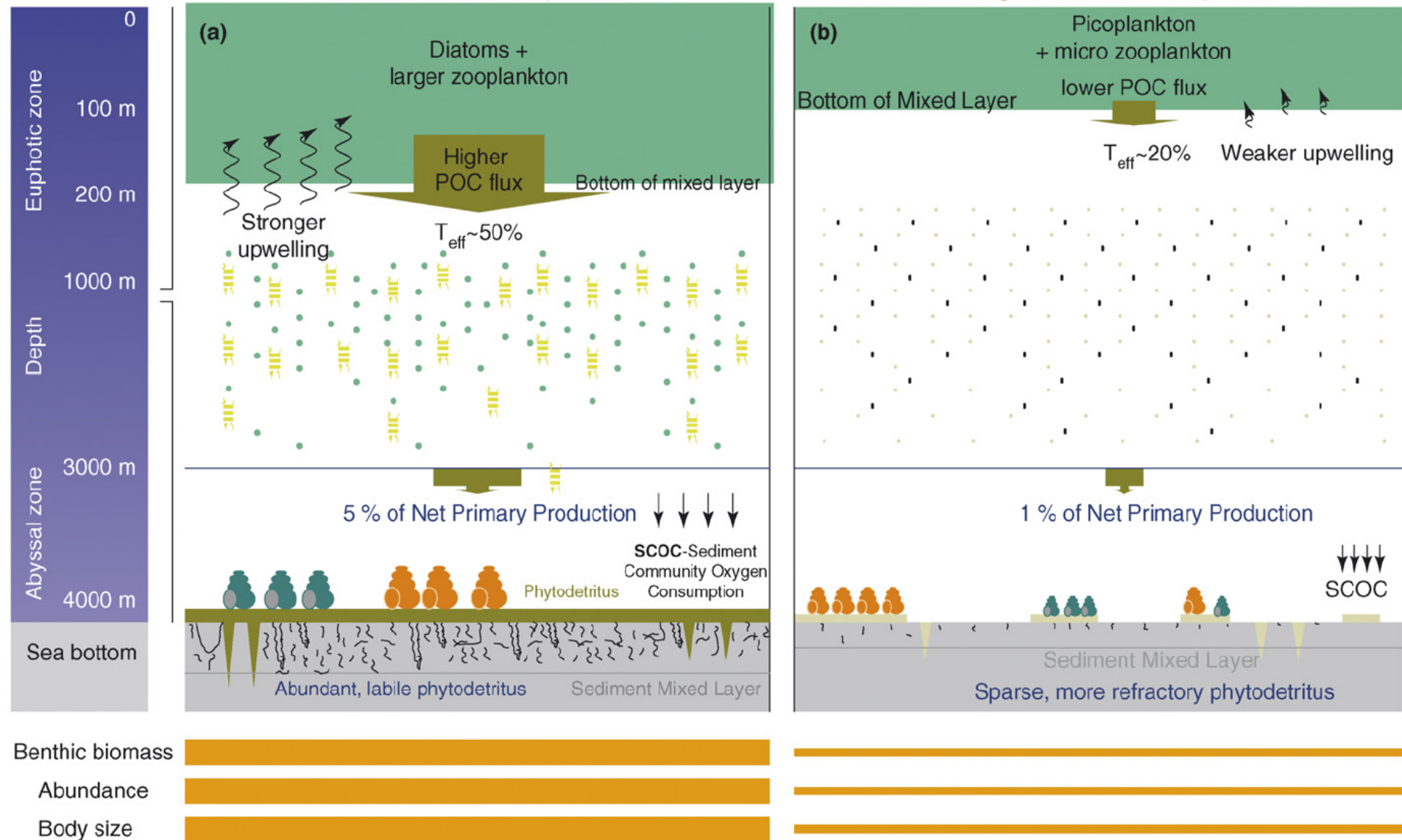
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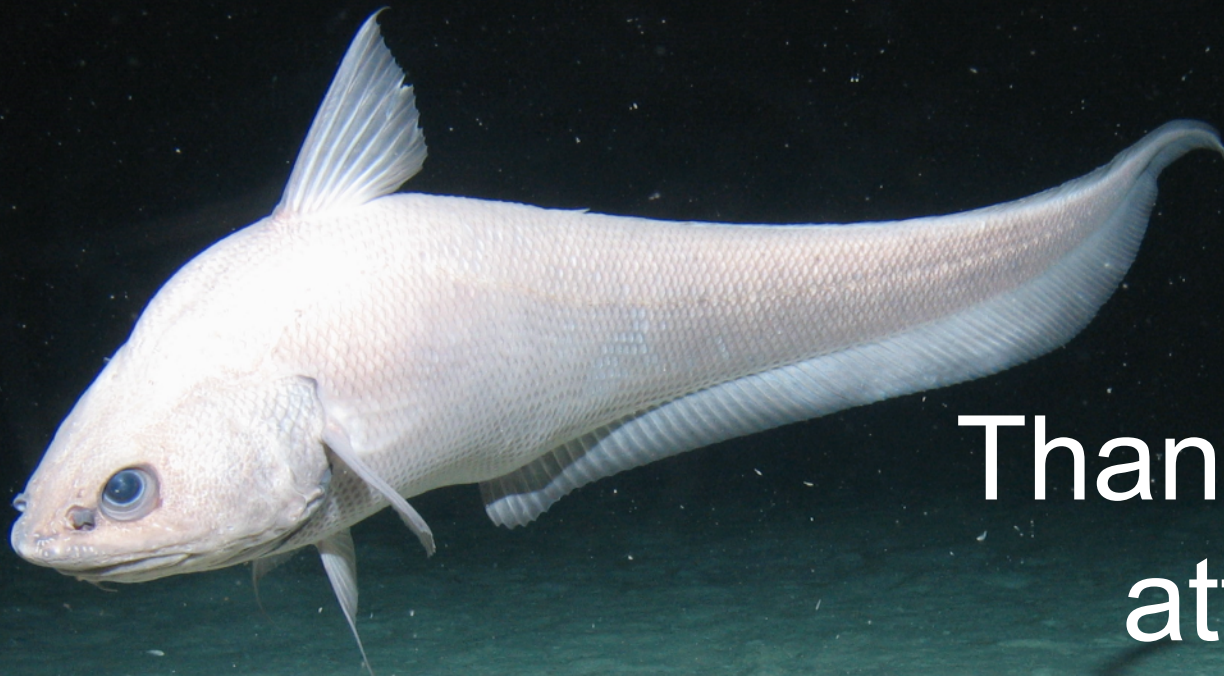
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Higher sea surface temperature





Thanks for your attention!

Photo: A. Jamieson, Oceanlab, University of Aberdeen

