



UKOA Annual Science Meeting
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Shipboard bioassays experiments

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Objectives

Quantitatively investigate the links between changes in the ocean carbonate system (Ocean Acidification; OA) and:

- natural microbial community physiology and morphometry (including CaCO_3 shells),
- plankton biodiversity and community structure,
- biogeochemical rates,
- food webs and
- climate-relevant processes.

Containerised lab facility

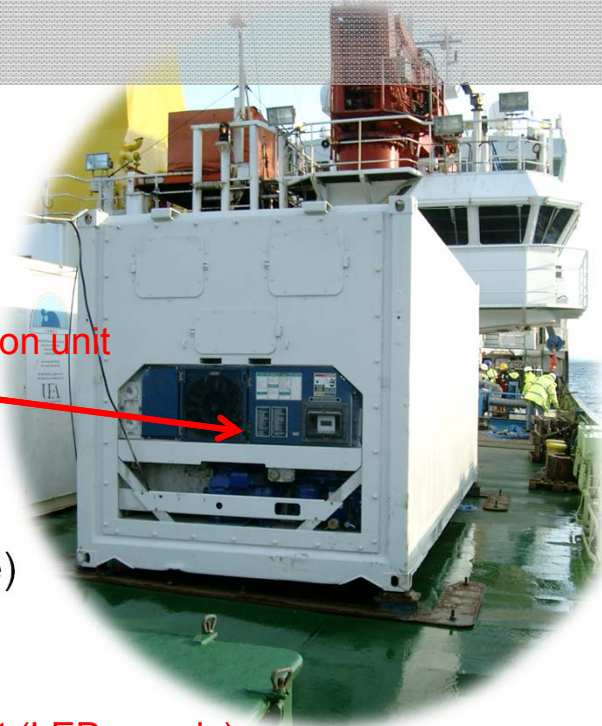
Controlled conditions in a refrigerated container:

Temperature adjusted to match with the *in situ*

Refrigeration unit

Artificial light sources: $100 \mu\text{E m}^{-2} \cdot \text{s}^{-1}$ (14/8 light/dark cycle)

White light (LED panels)

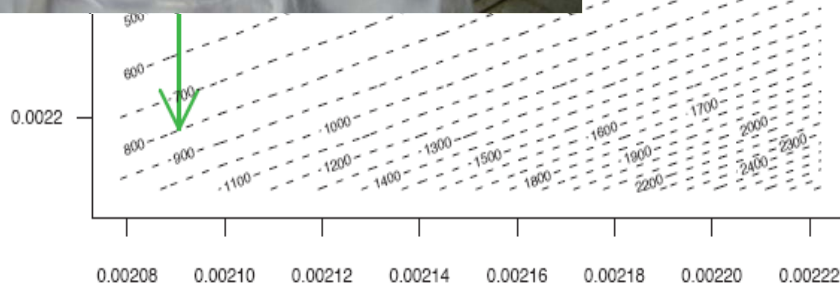


Set-up



TD of water, 24 x 20L = 480L,
 er into bottles on deck

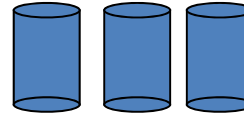
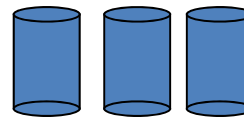
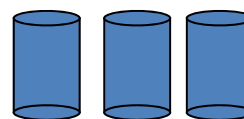
$\text{HCO}_3^- + \text{HCl}$ to achieve target pCO_2



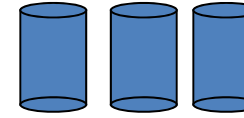
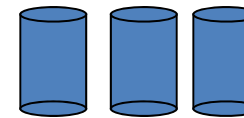
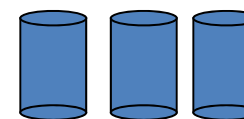
Dissolved inorganic carbon (mol kg^{-1})



m

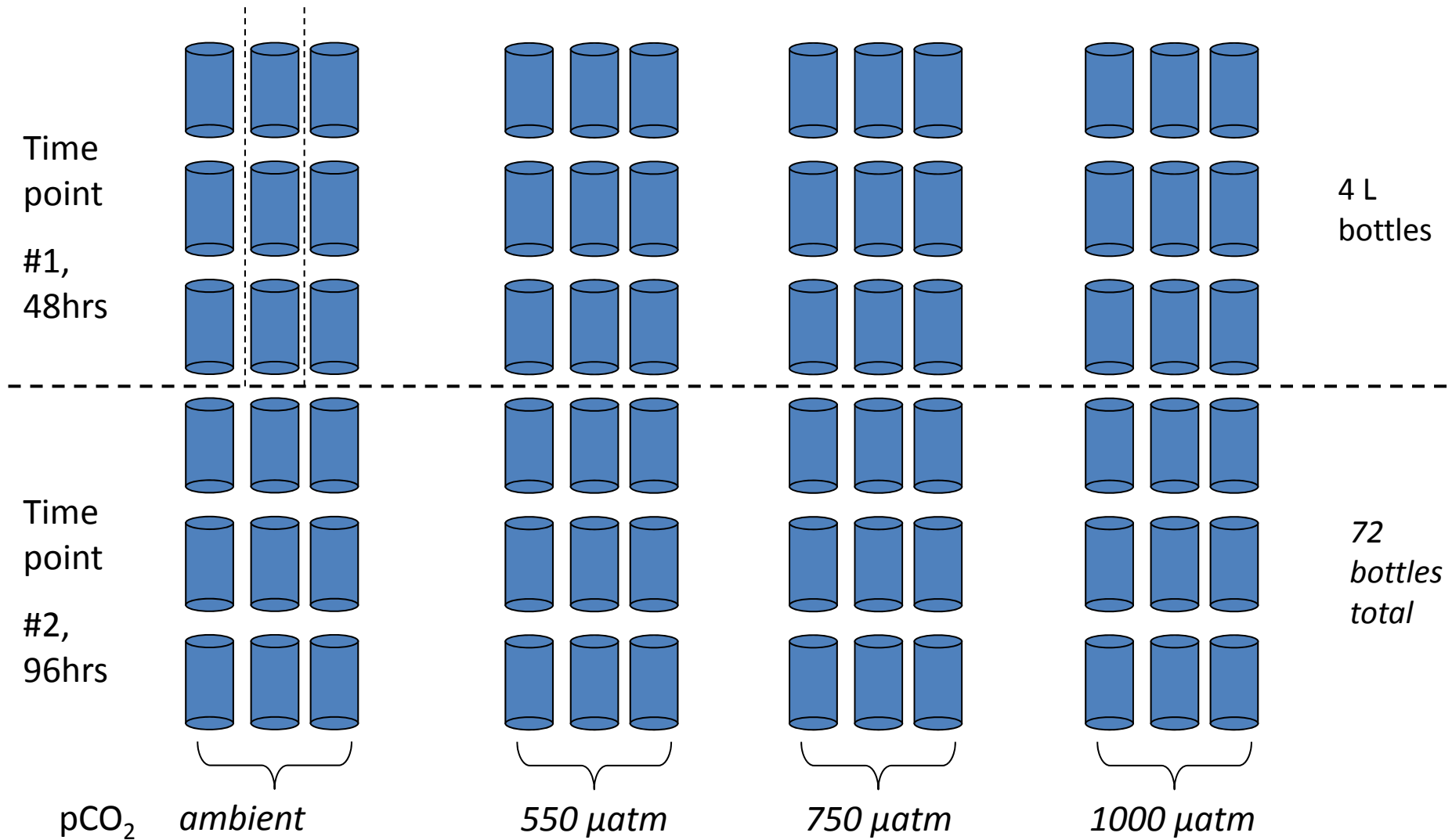


750 µatm



1000 µatm

Set-up



Cruise track (June-July 2011) and bioassay locations

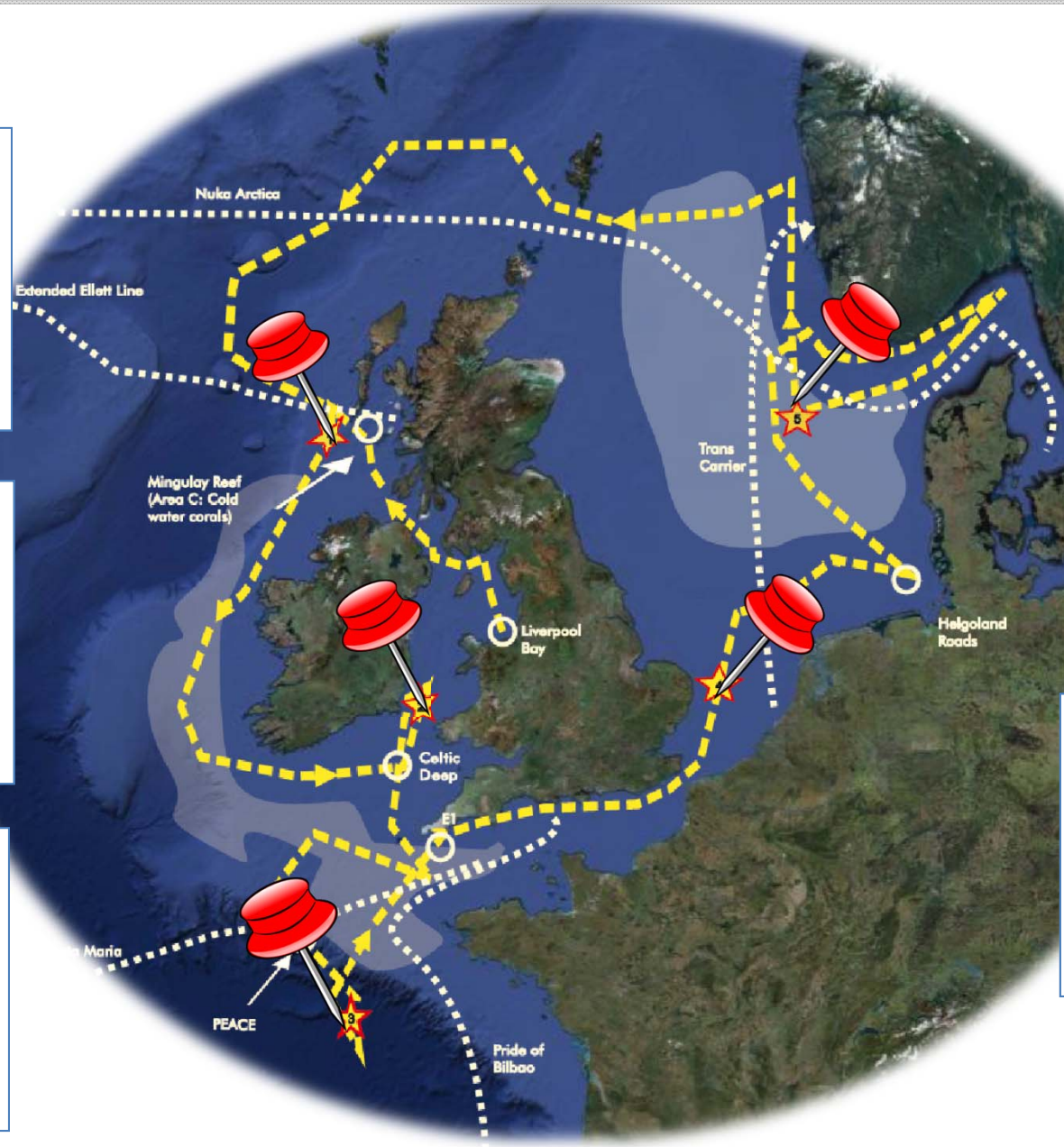
E01 Stratified
 T° C: 11.27
 Salinity: 34.8
 pCO₂: 338
 DIN: 1 μM
 Si: 2 μM
 P: 0.09 μM

E02 Mixed
 T° C: 11.77
 Salinity: 34.44
 pCO₂: 336
 DIN: 0.3 μM
 Si: 0.44 μM
 P: 0.14 μM

E03 Stratified
 T° C: 15.31
 Salinity: 35.77
 pCO₂: 342
 DIN: 0.56 μM
 Si: 0.6 μM
 P: 0.05 μM

E05 Stratified
 T° C: 13.96
 Salinity: 35
 pCO₂: 370
 DIN: 1.7 μM
 Si: 0.04 μM
 P: 0.06 μM

E04 Mixed
 T° C: 14.57
 Salinity: 34.05
 pCO₂: 407
 DIN: 0.8 μM
 Si: 0.8 μM
 P: 0.13 μM



Measured parameters

Carbonate chemistry (*UoSoton*)

Macro-nutrients (*NOCS*)

POC/PON/POP (*UoSoton*)

Total Chl_a/size fractionated chlorophyll_a (*UoSoton, UoEssex*)

Photophysiology (*Fv/Fm, σ, τ*) (*UoEssex*)

P/E (*UoEssex*)

Primary production (*NOCS*)

Calcite production (*NOCS*)

Coccolithophores assemblage (*NHM*)

PIC (*NHM*)

Microbial community composition (*NOCS, PML*)

Respiration (*UoSoton*)

Ammonium (*UoSoton*)

DOC/TEP (*UoSoton*)

DMSP production, DMSP-DMSP conversion rates (*PML*)

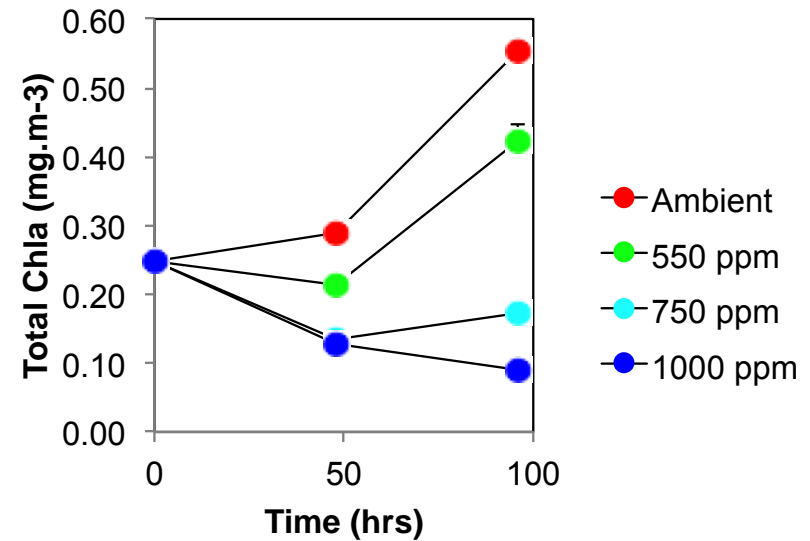
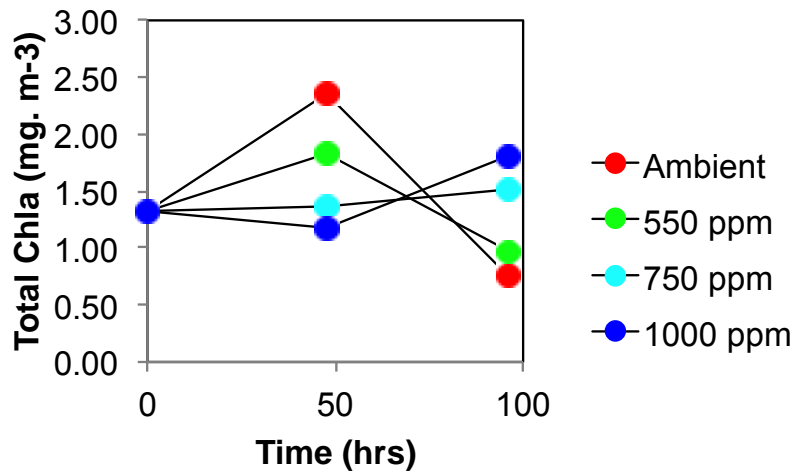
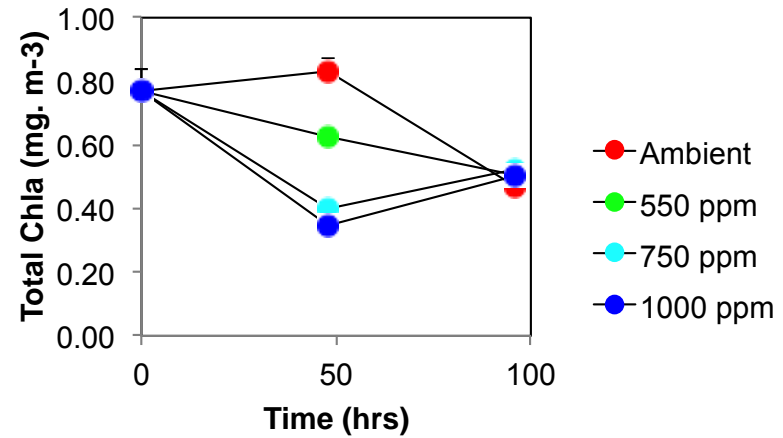
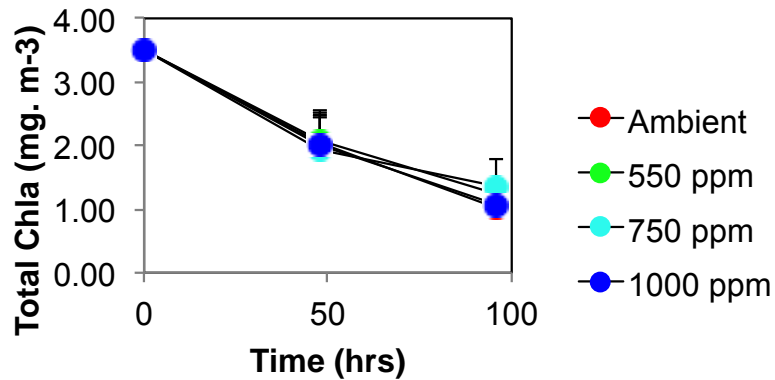
CH₄, N₂O (*PML*)

Oxidation rates of NH₄⁺ and NO₂⁻ (*PML*)

Genetic diversity (*MBA*)

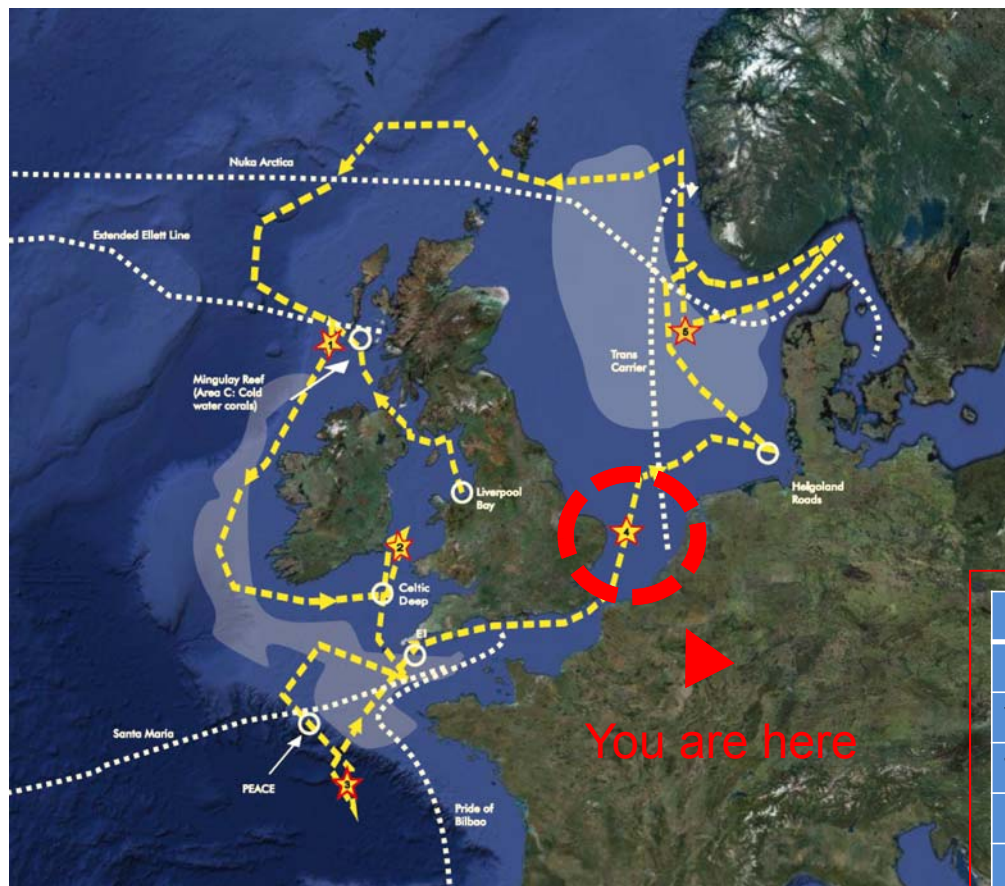
Proteins/RNA (*UoSoton*)

Diversity in responses to ocean acidification



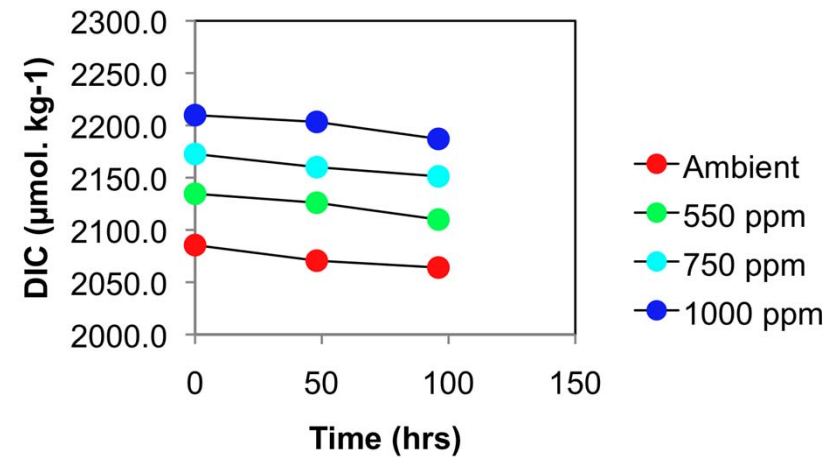
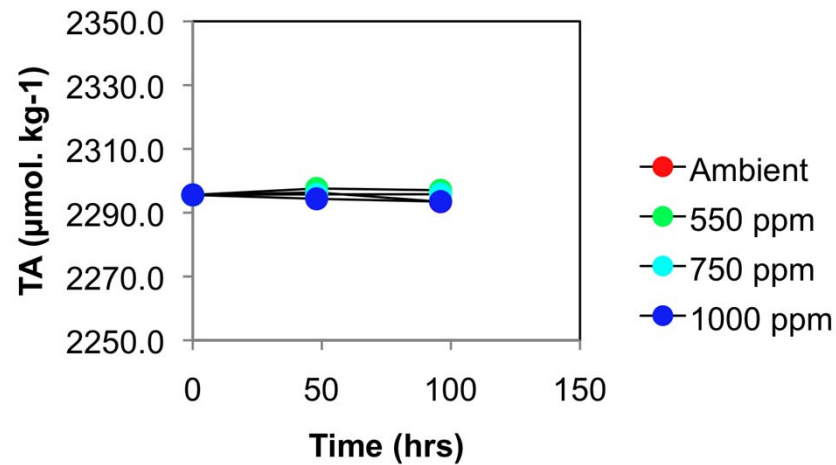
Discrepancies in Chl_a trends between bioassays illustrate the importance of initial conditions in response to OA.

Zoom on the South North sea bioassay (E04)



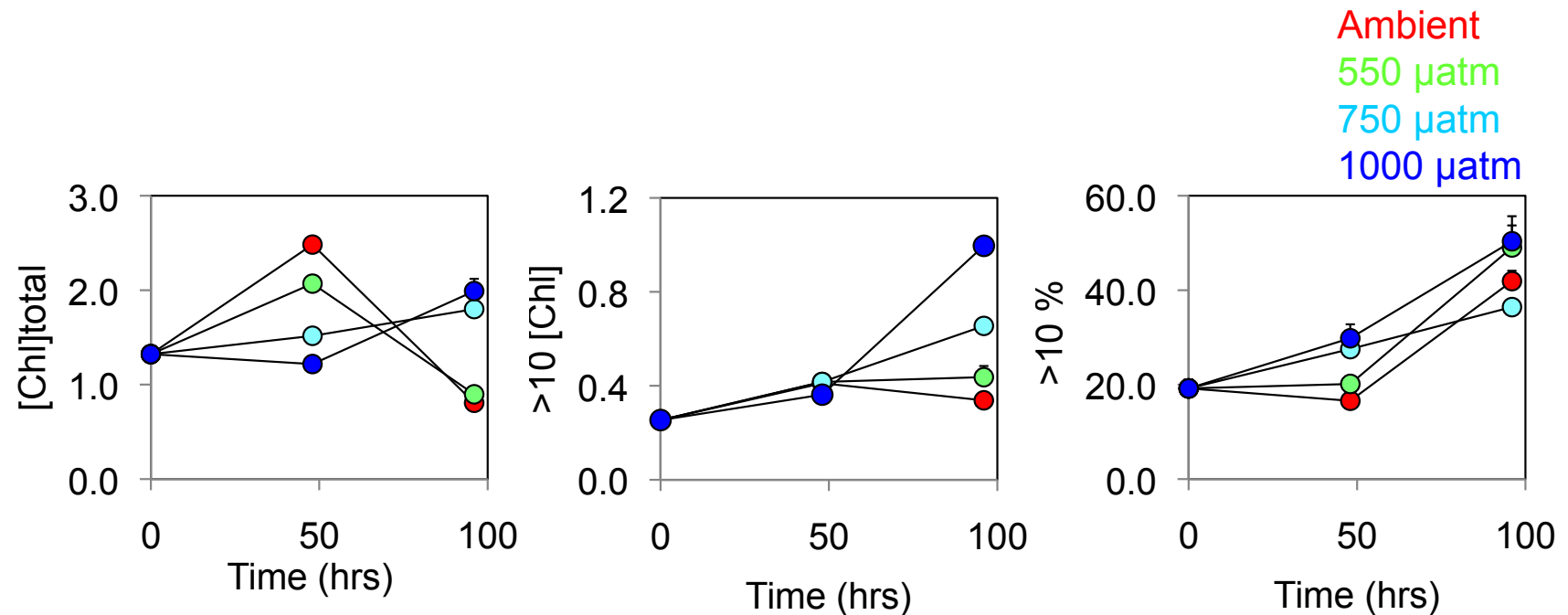
Latitude [deg+veN]	52.59
Longitude [deg+veW]	2.29
Water column	Mixed
Temperature (°C)	14.57
Salinity	34.05
pCO ₂ (ppm)	407
Nitrates/nitrites (µM)	0.8
Phosphates (µM)	0.13
Silicates (µM)	0.8

Carbonate chemistry in E04 bioassay



As expected, the TA remained unchanged with time and pCO₂ through the experiment while the DIC concentration is increasing with pCO₂ and decreasing with time.

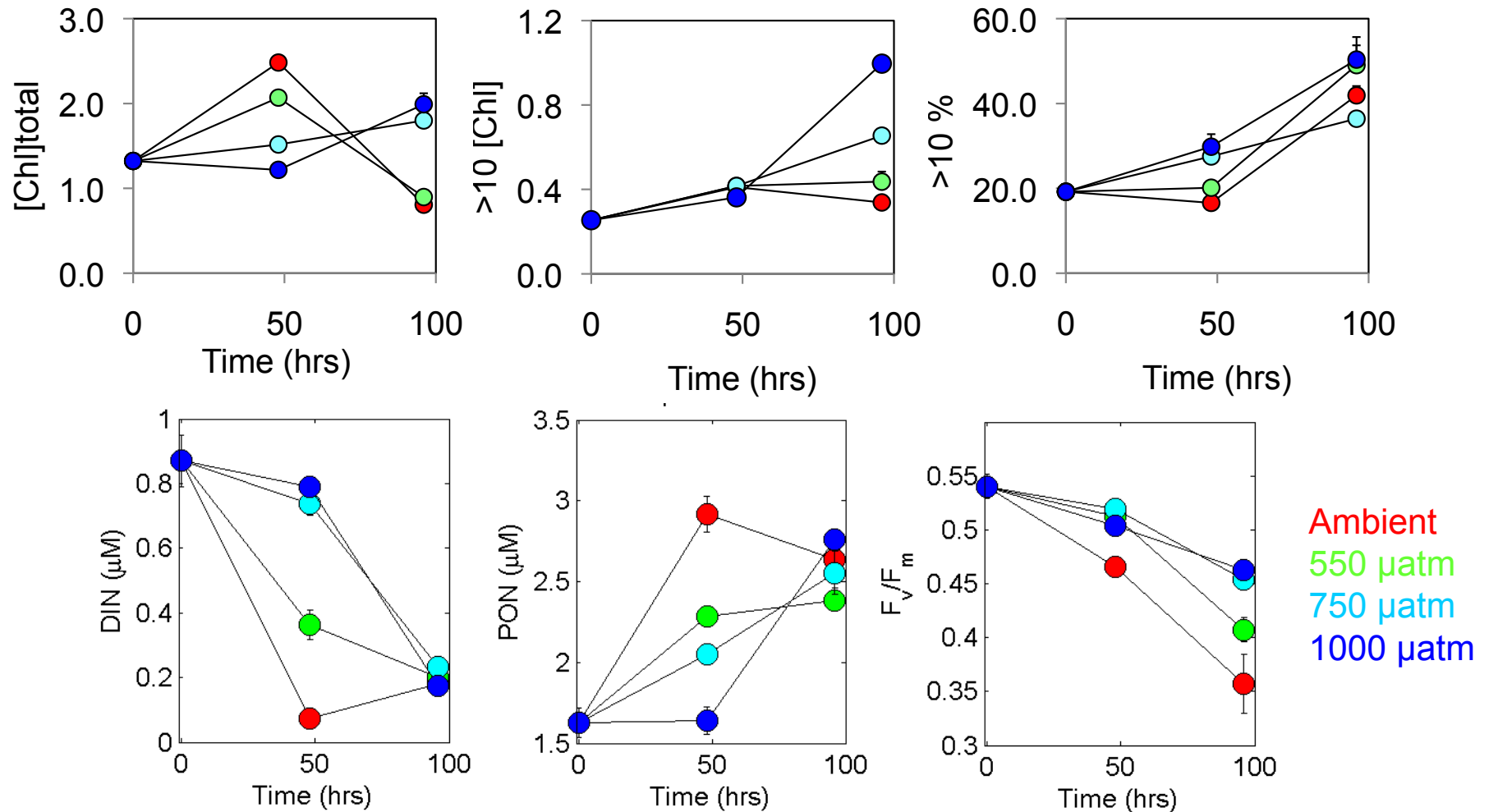
Interpretation of Time series experiments



Highest Chl_a values for the ambient condition after 2 days while the bloom is delayed with increasing pCO₂.

The small size phytoplankton dominates the microbial community (80% at T₀) and appears more sensitive to ocean acidification than > 10 μm fraction.

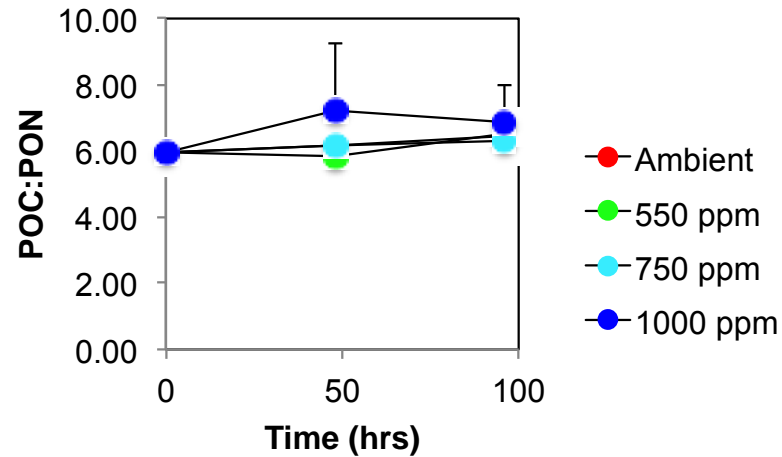
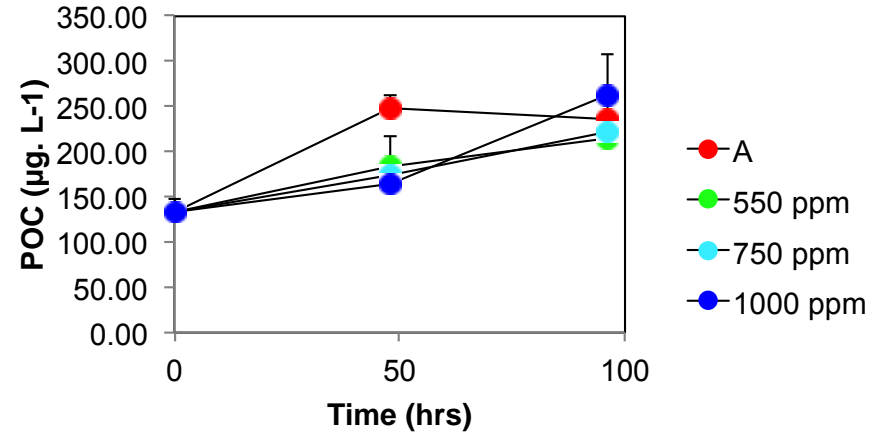
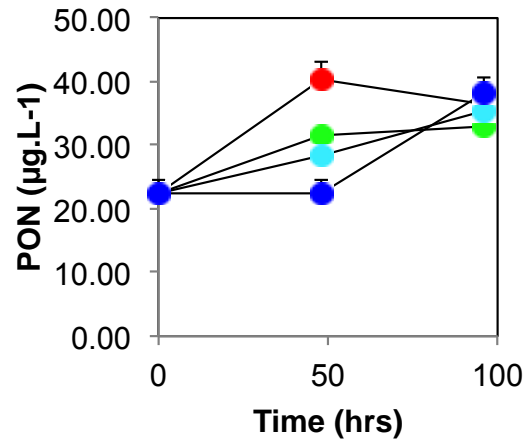
Interpretation of Time series experiments



Slow down in nutrients uptake and organic matter production with increasing pCO₂.

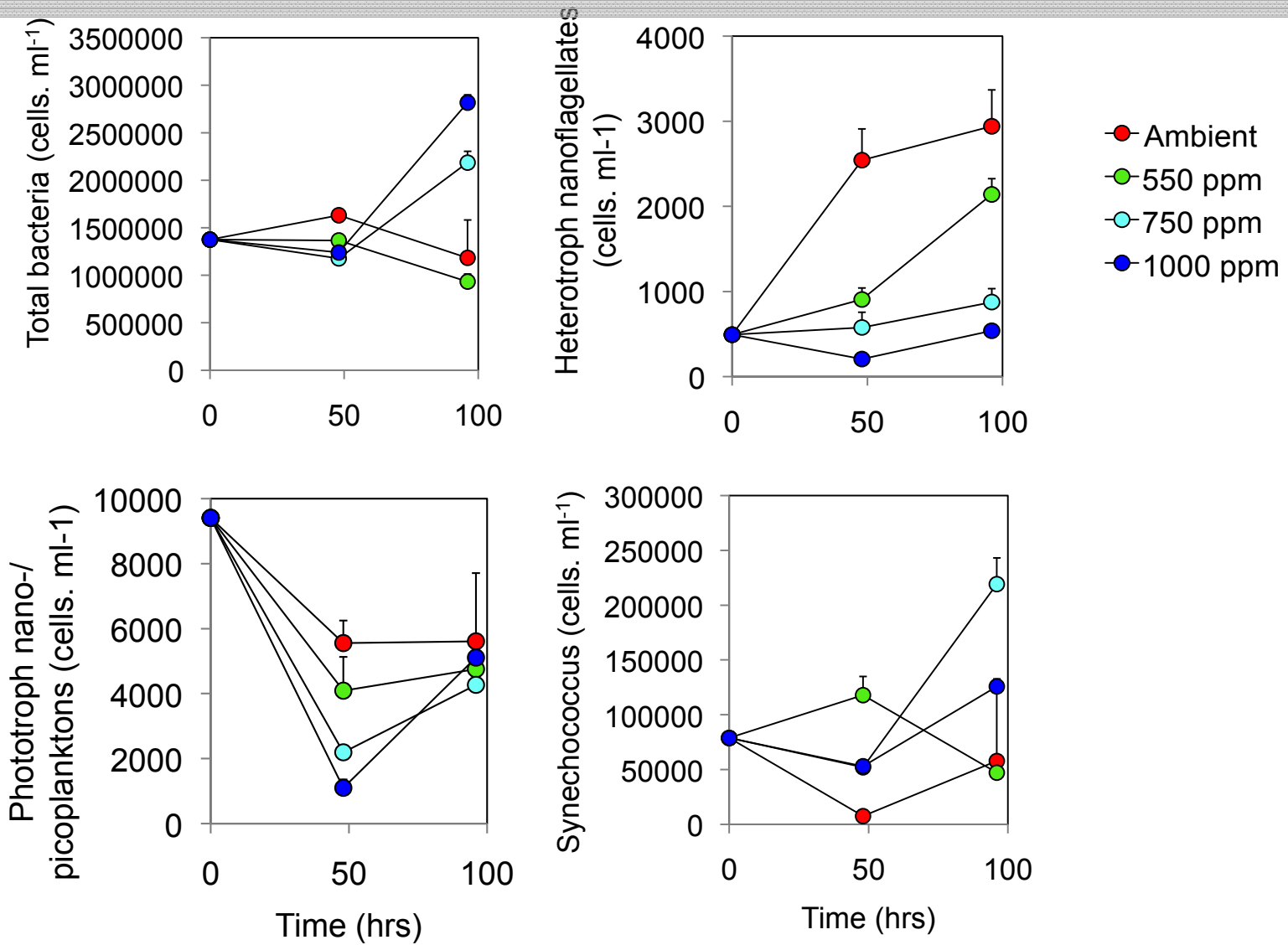
Increase in photo-efficiency with increasing pCO₂ (specific to this bioassay).

Interpretation of Time series experiments



The production of organic matter is affected by high pCO₂ while C:N ratio remains unchanged.

Interpretation of Time series experiments



Shift in community composition with increasing pCO₂.

Conclusions

The natural communities observed were impacted by ocean acidification.

After 2 days, the chlorophyll content is higher under low $p\text{CO}_2$. The peak of chlorophyll corresponds to the highest particulate organic matter concentration and the exhaustion of nutrient sources.

After 4 days, the trend is inverted. As the nutrients uptake is lower at high $p\text{CO}_2$, the bloom is delayed and occurs later.

C:N stoichiometry was not affected by higher $p\text{CO}_2$ and similar to the Redfield ratio of 6.6 in today's ocean.

Shift in microbial community structure (increase in bacteria and decrease in small size fraction and nanoflagellates) with changing $p\text{CO}_2$.

Take home messages

All the measurements performed during the cruise are required to interpret correctly the data from each bioassay,

A strong group-specific response to high $p\text{CO}_2$ was observed in the majority of bioassay experiments.

Such responses have important implications for trophic dynamics and biogeochemical cycling in both present day and future oceanic ecosystems

The diversity of initial environmental conditions as well as the multidisciplinary approach adopted as part of the project will potentially highlight new acclimation mechanisms of the planktonic community to future ocean acidification.

Acknowledgments



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