

Response of spring diatoms to CO₂ availability in the western North Pacific

Koji Suzuki¹, Hisashi Endo^{1,2}, Koji Sugie^{3,4}, and
Takeshi Yoshimura^{1,3}

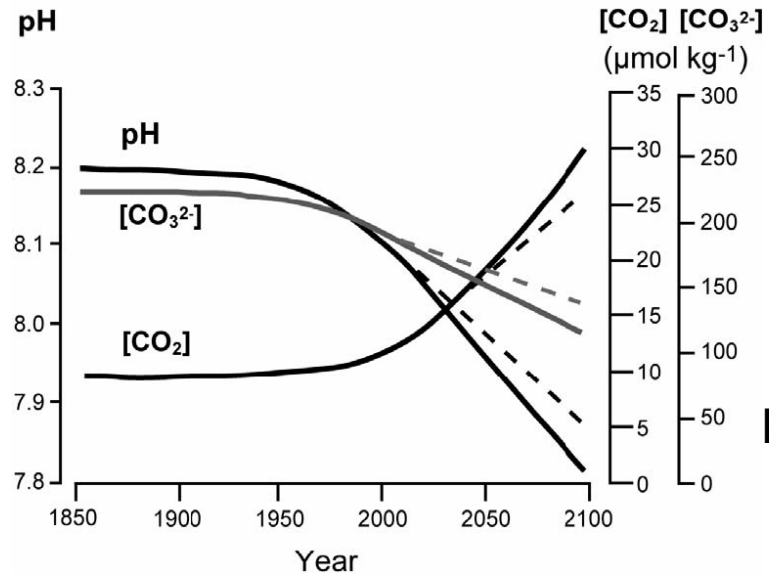
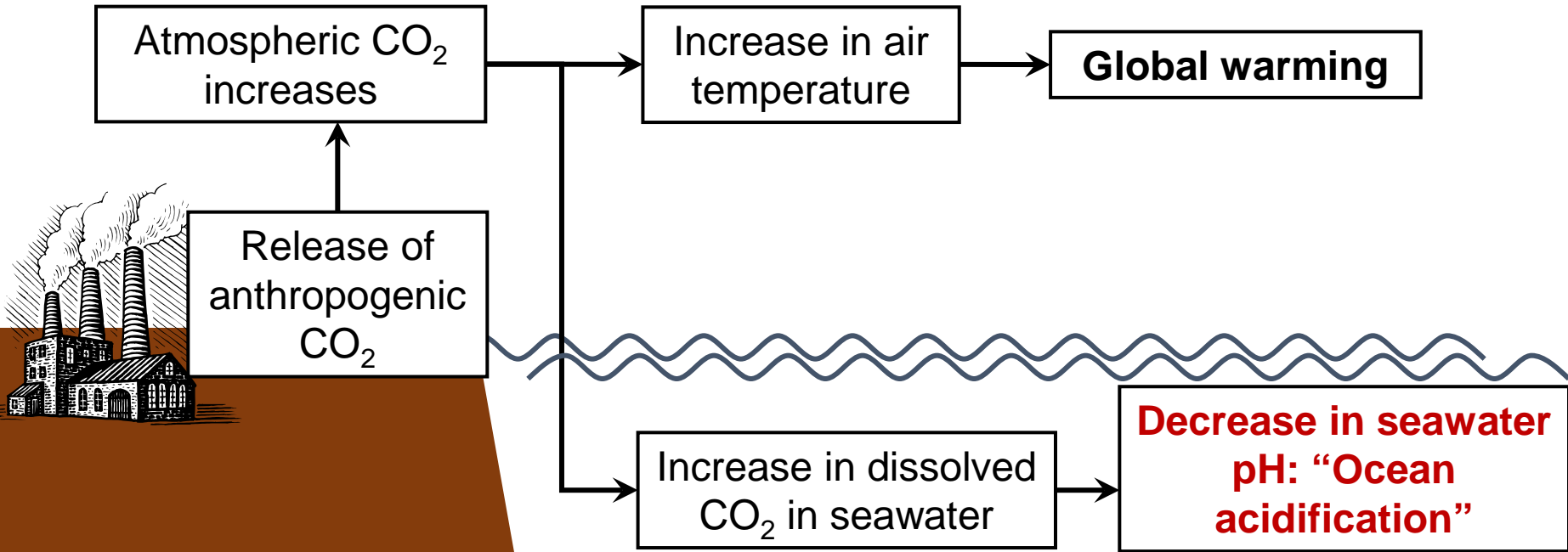
¹Hokkaido University, ²Kyoto University

³Central Research Institute of Electric Power Industry

⁴Japan Agency for Marine Earth-Science and Technology (JAMSTEC)



Ocean acidification

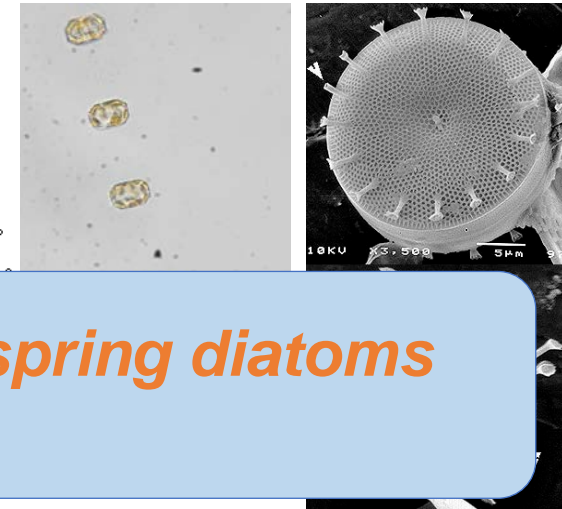
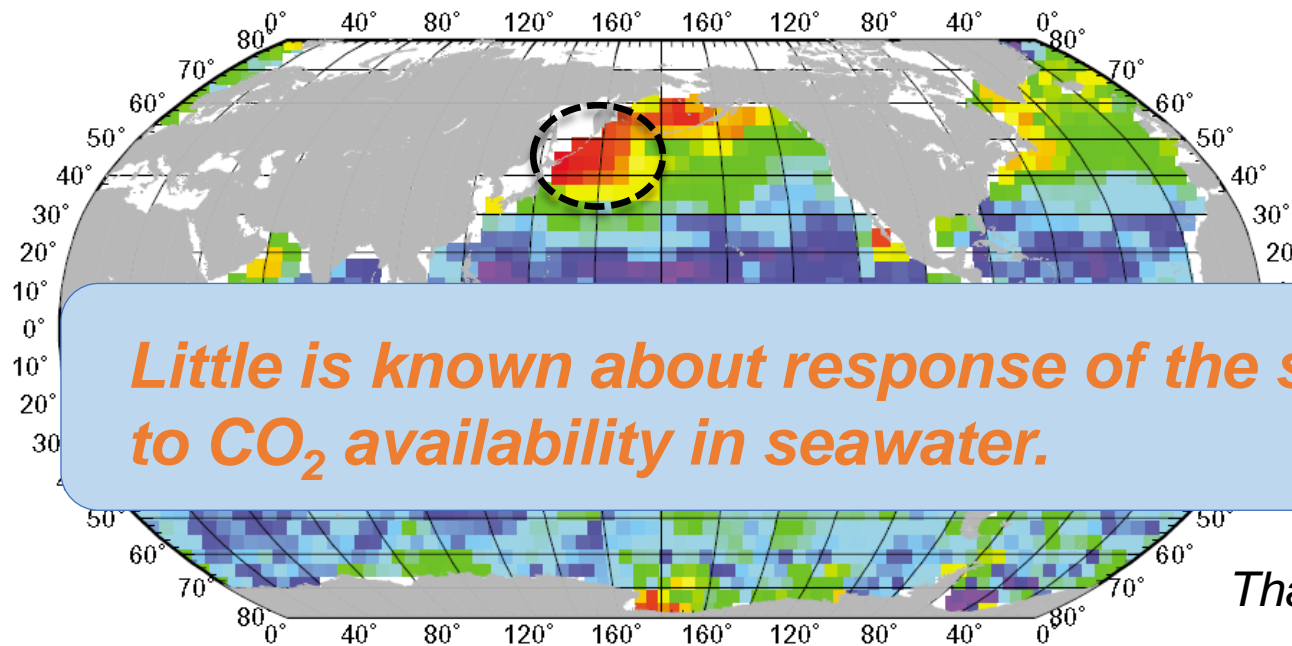


Riebesell
(2004)

In the western subarctic Pacific

- Highest biological drawdown of partial pressure of CO₂ ($p\text{CO}_2$) in surface waters among the world's oceans (Takahashi et al., 2002).
- The high $p\text{CO}_2$ drawdown is attributable to massive spring diatom bloom (Midorikawa et al., 2003; Ayers and Lozier, 2012).

Seasonal Biological Drawdown of Seawater $p\text{CO}_2$



Little is known about response of the spring diatoms to CO₂ availability in seawater.

Thalassiosira nordenskiöldii



$p\text{CO}_2$ Drawdown (μatm)

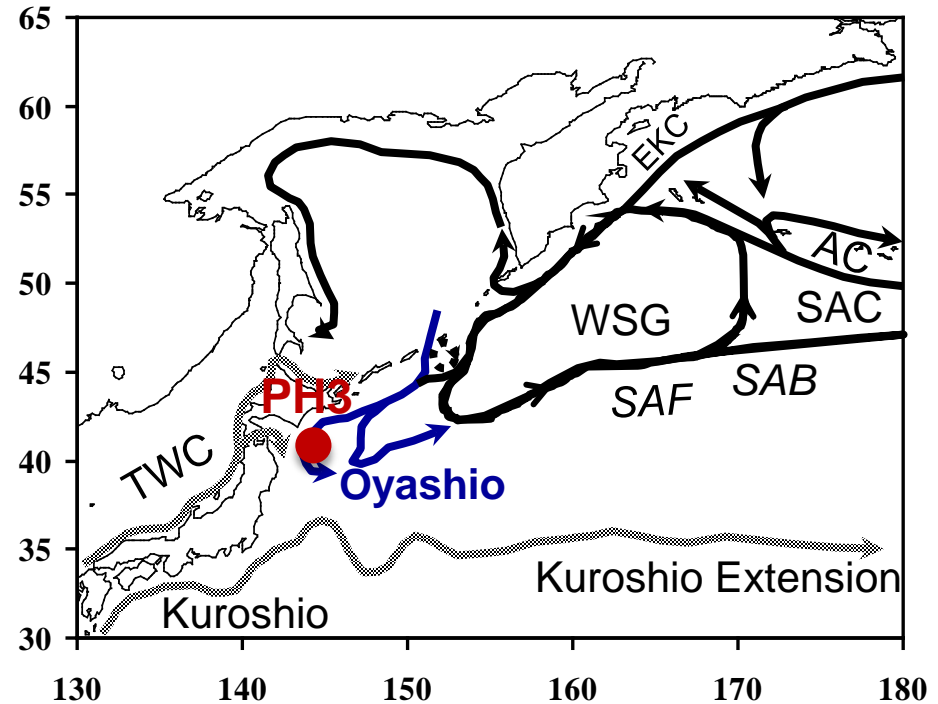
Takahashi et al. (2002)

On deck CO₂ bottle incubation experiment

- Sampling date: May 8, 2011
- Site: Stn PH3 (41°N, 144°E)
- Sampling depth: 10 m
- Incubation period: ca. 3 days

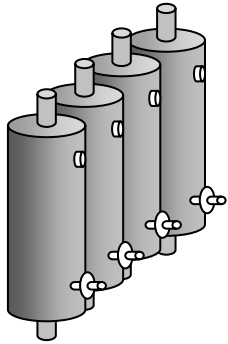


R/V *Tansei Maru* (JAMSTEC/ Univ. Tokyo)



Initial conditions at 10 m:

- Temperature: 5.0 °C
- Salinity: 33.1
- Chl *a*: 0.71 µg L⁻¹
- Nitrate: 14.0 µM
- Phosphate: 0.95 µM
- Silicate: 11.8 µM
- *p*CO₂: 342.8 µatm

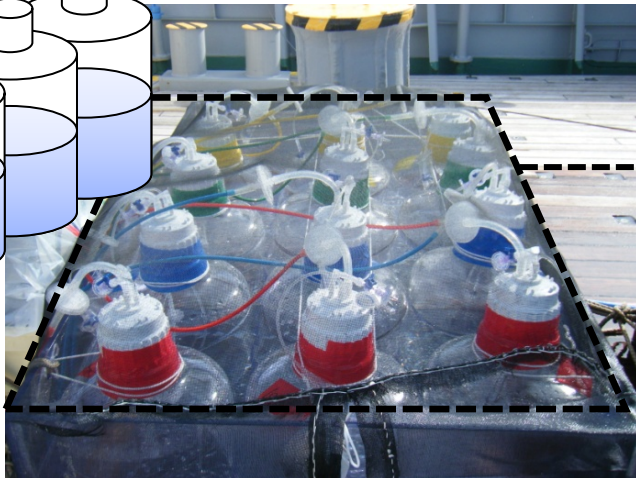
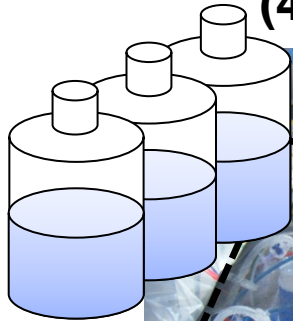


Clean Niskin bottle sampling

Pre-filtration with 197 μm teflon mesh

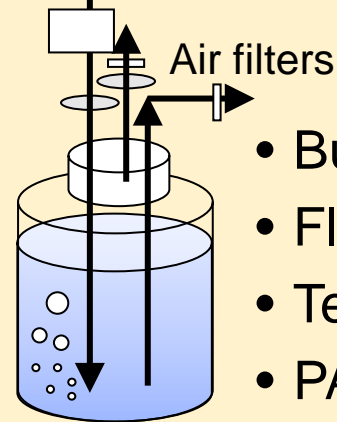


**12 L polycarbonate bottles
(4 CO_2 conditions \times triplicate = 12 bottles)**



Incubation conditions

Air + CO_2

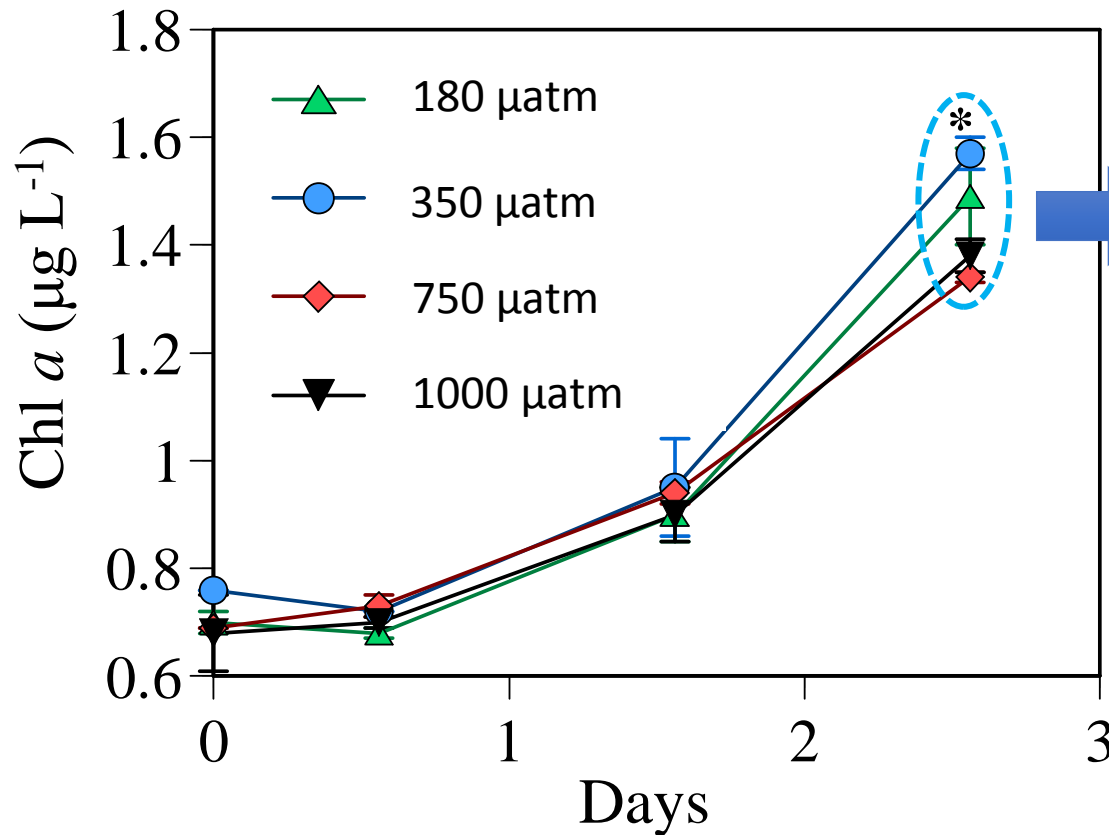


- Bubbling of CO_2 gases
- Flow rate: 50 mL min^{-1}
- Temperature: $5 \text{ }^\circ\text{C}$
- PAR: 50% light level

CO_2 levels:

- $180 \mu\text{atm CO}_2$
- $350 \mu\text{atm CO}_2$ (ambient)
- $750 \mu\text{atm CO}_2$
- $1,000 \mu\text{atm CO}_2$

Changes over time in chlorophyll (Chl) a concentration determined with high-performance liquid chromatography (HPLC)



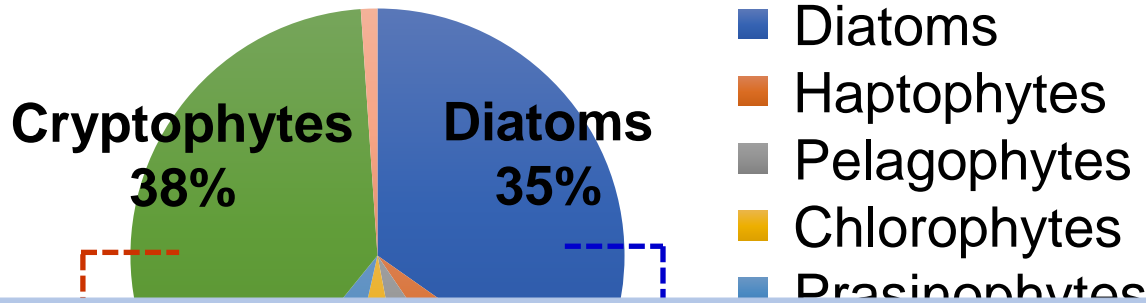
Chl a level
at 350 µatm pCO₂
v
Chl a level
at 750 and 1000 µatm
pCO₂

(Tukey's honest
significant test
(HSD), $p < 0.05$,
 $n = 3$)

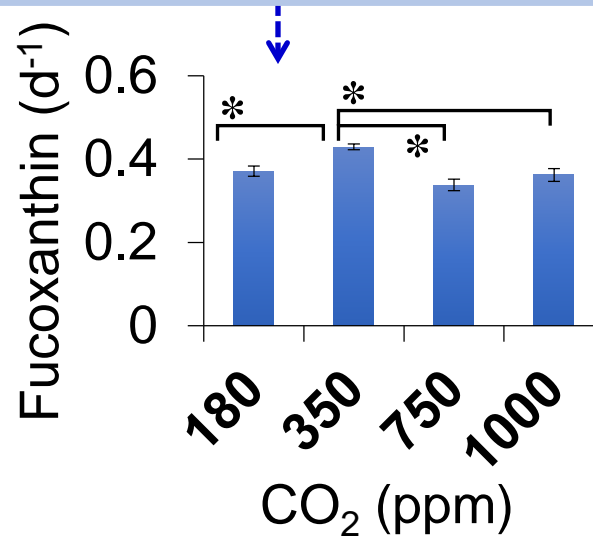
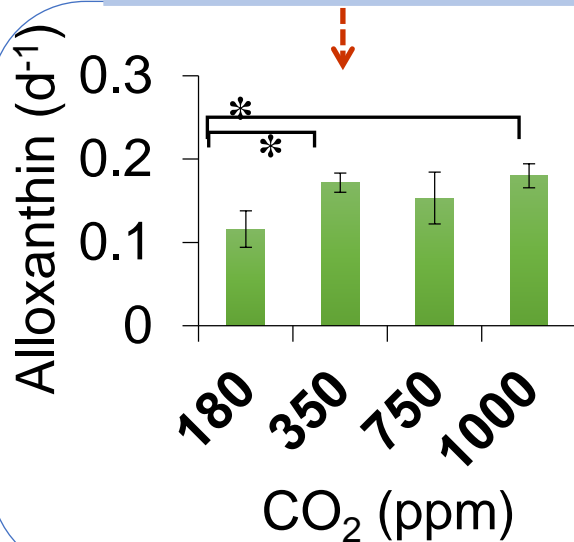
The net growth of phytoplankton assemblages was suppressed by an increase in CO₂ level.

Changes in algal community composition

Contributions of each algal taxa to Chl *a* biomass on **Day 0** as estimated with the program CHEMTAX (Mackey et al., 1996; Latasa, 2007)



The decreases in Chl *a* level at higher CO₂ levels after incubation were probably due to declines in diatom abundance.



On **Day 3**

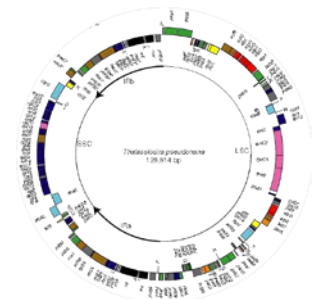
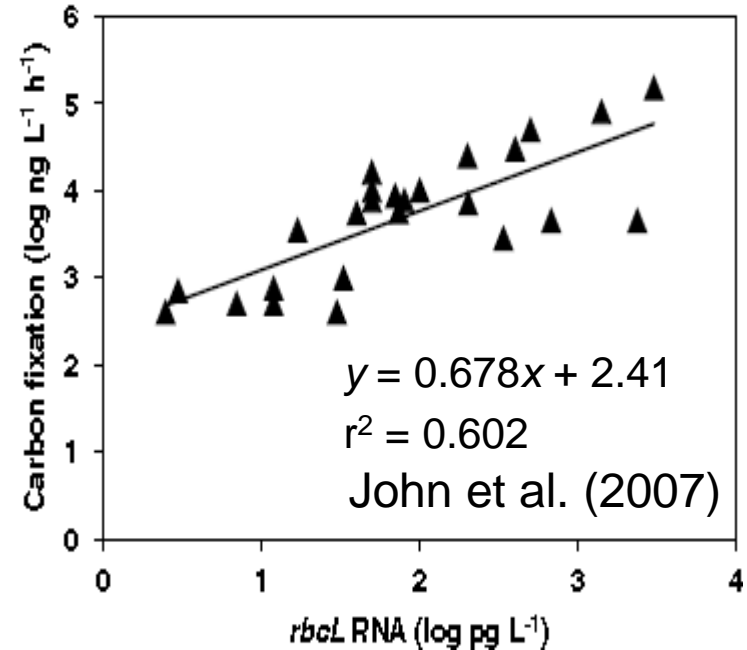
net growth rates of each taxonomic pigment marker

* Tukey's HSD, $p < 0.05$, $n = 3$

RubisCO

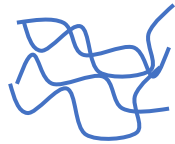
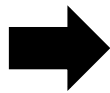
(Ribulose-1,5-bisphosphate carboxylase/oxygenase)

- In algae, CO₂ is fixed in the Calvin Benson cycle catalyzed by the enzyme RubisCO.
- The large subunit of RubisCO is encoded by *rbcL* gene, which can be regulated by environmental factors (John et al., 2007).
- Diatom abundance and photosynthetically active diatoms can be inferred from *rbcL* DNA or cDNA fragments (Endo et al., 2014; 2016)



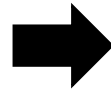
Plastid genome
(DNA)

Transcription



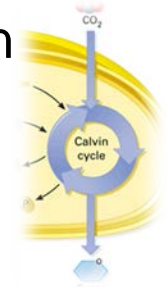
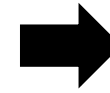
rbcL mRNA
(cDNA)

Translation



RubisCO

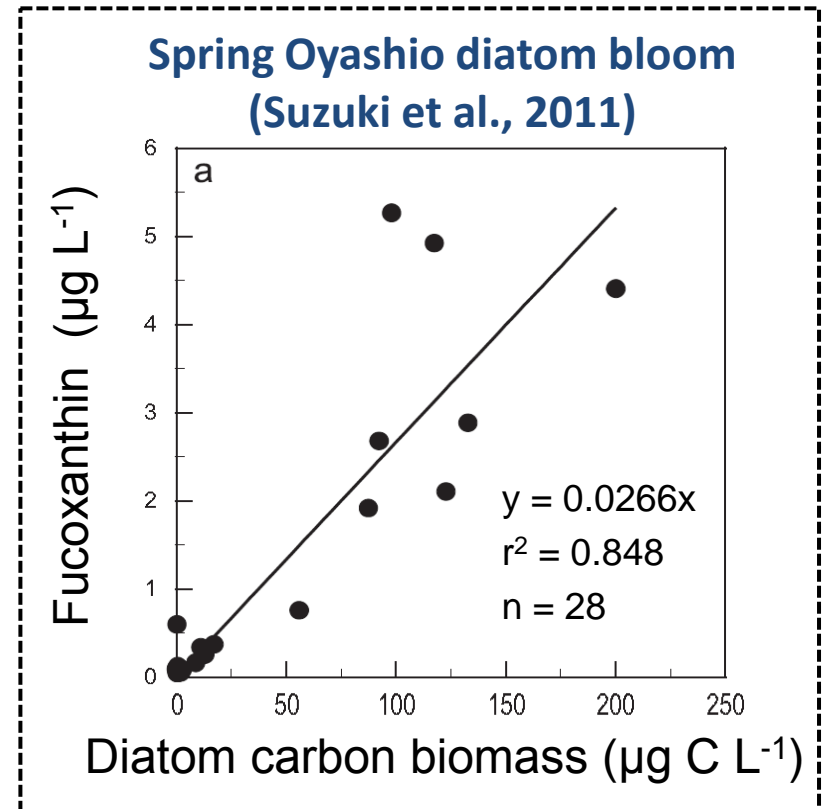
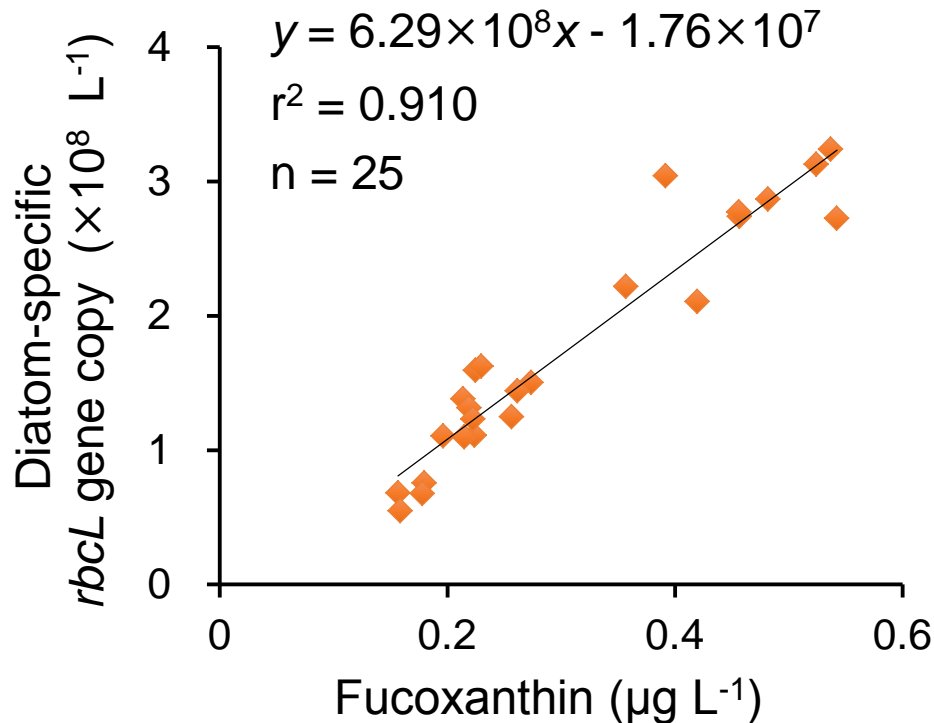
Functional
expression



CO₂ fixation

Copy numbers of diatom-specific *rbcL* gene as determined by quantitative PCR (qPCR)

Diatom-specific PCR primers: John et al. (2007)

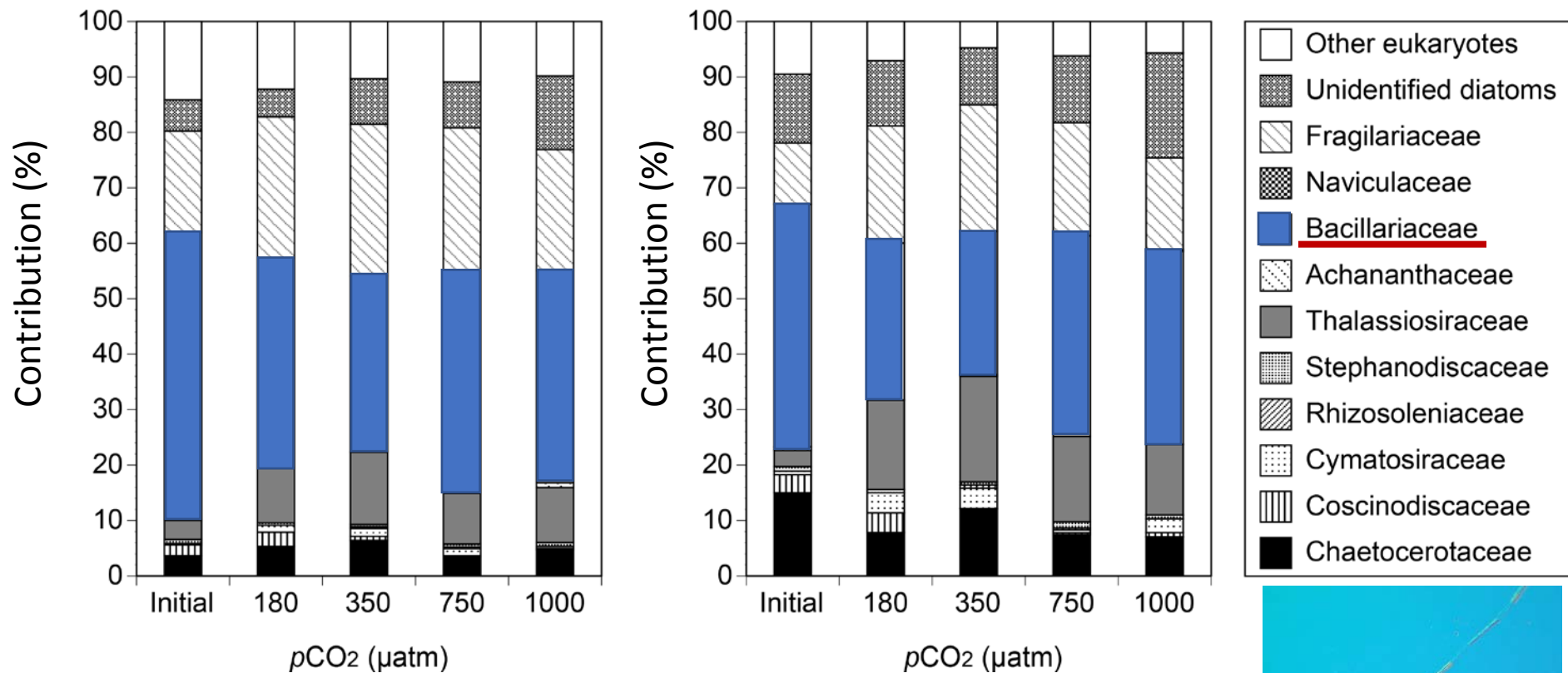


A significant correlation was found between fucoxanthin and diatom-specific *rbcL* gene levels → **Copy number of diatom-specific *rbcL* gene can become an indicator for diatom biomass.**

Relative contributions (%) of each diatom family to the *rbcL* DNA or cDNA libraries from the initial and each CO₂ treatment on day 3 as estimated with next-generation sequencing (Ion Torrent) technology

rbcL DNA library (abundance)

rbcL cDNA library (photosynthetic activity)



“**Bacillariaceae**” contains some pennate diatoms such as *Pseudo-nitzschia*, *Nitzschia*, and *Cylindrotheca* genera.

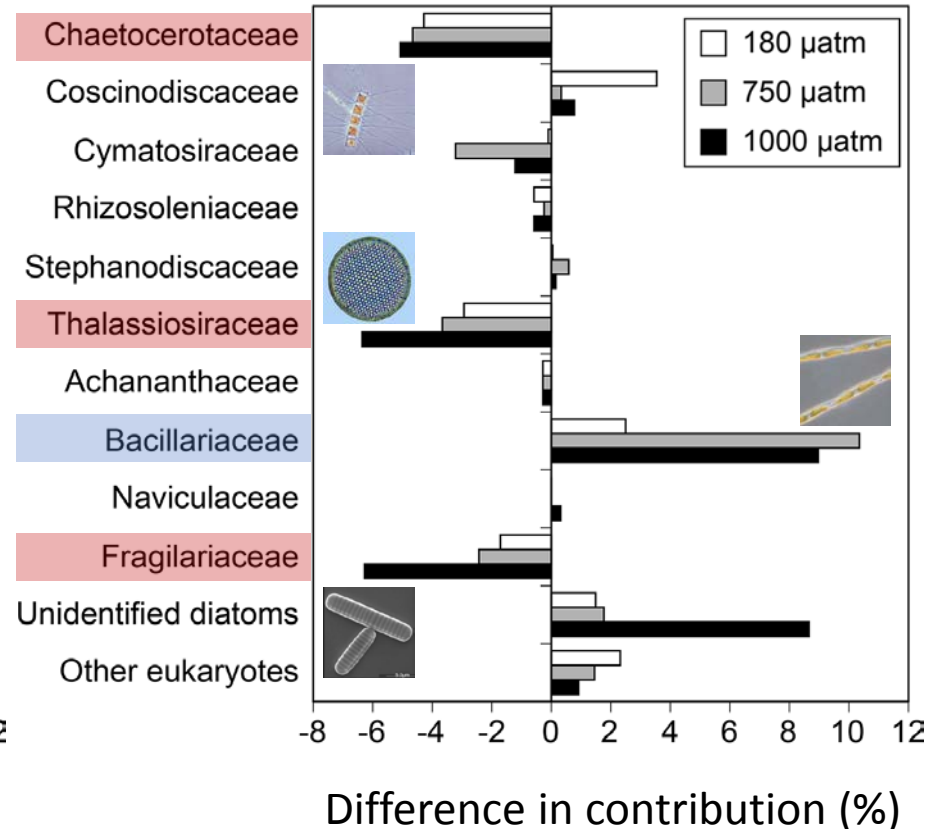
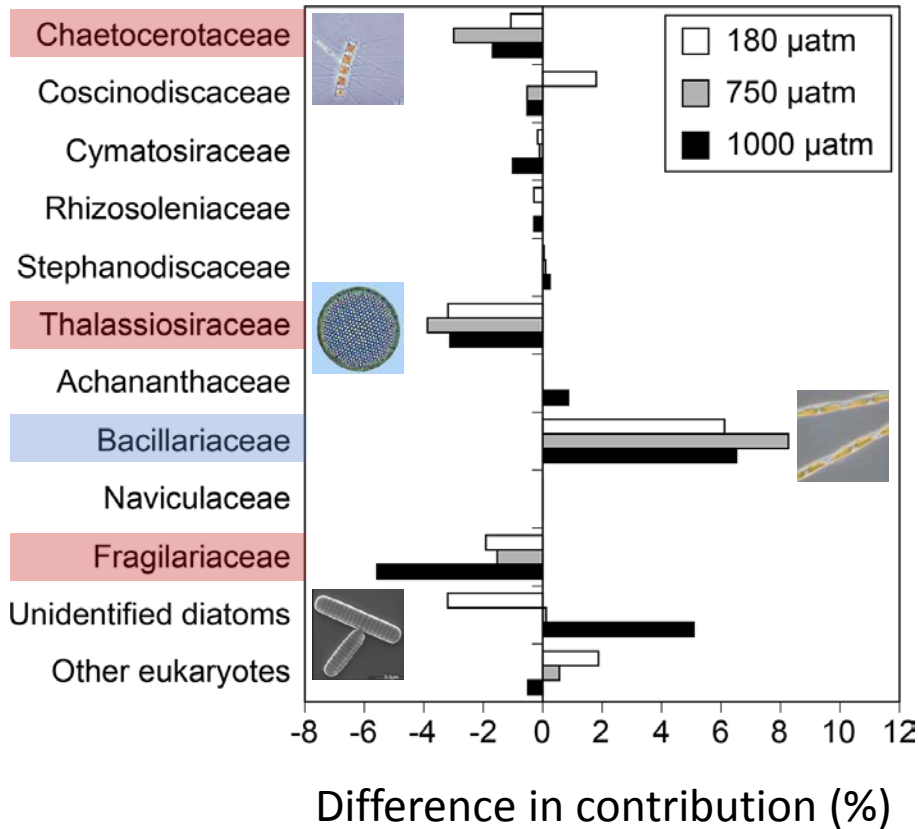


Pseudo-nitzschia seriata

Percent differences in *rbcL* contribution (%) between the control (350 μatm $p\text{CO}_2$) and other $p\text{CO}_2$ treatments in the diatom DNA or cDNA libraries

rbcL DNA library (abundance)

rbcL cDNA library (photosynthetic activity)

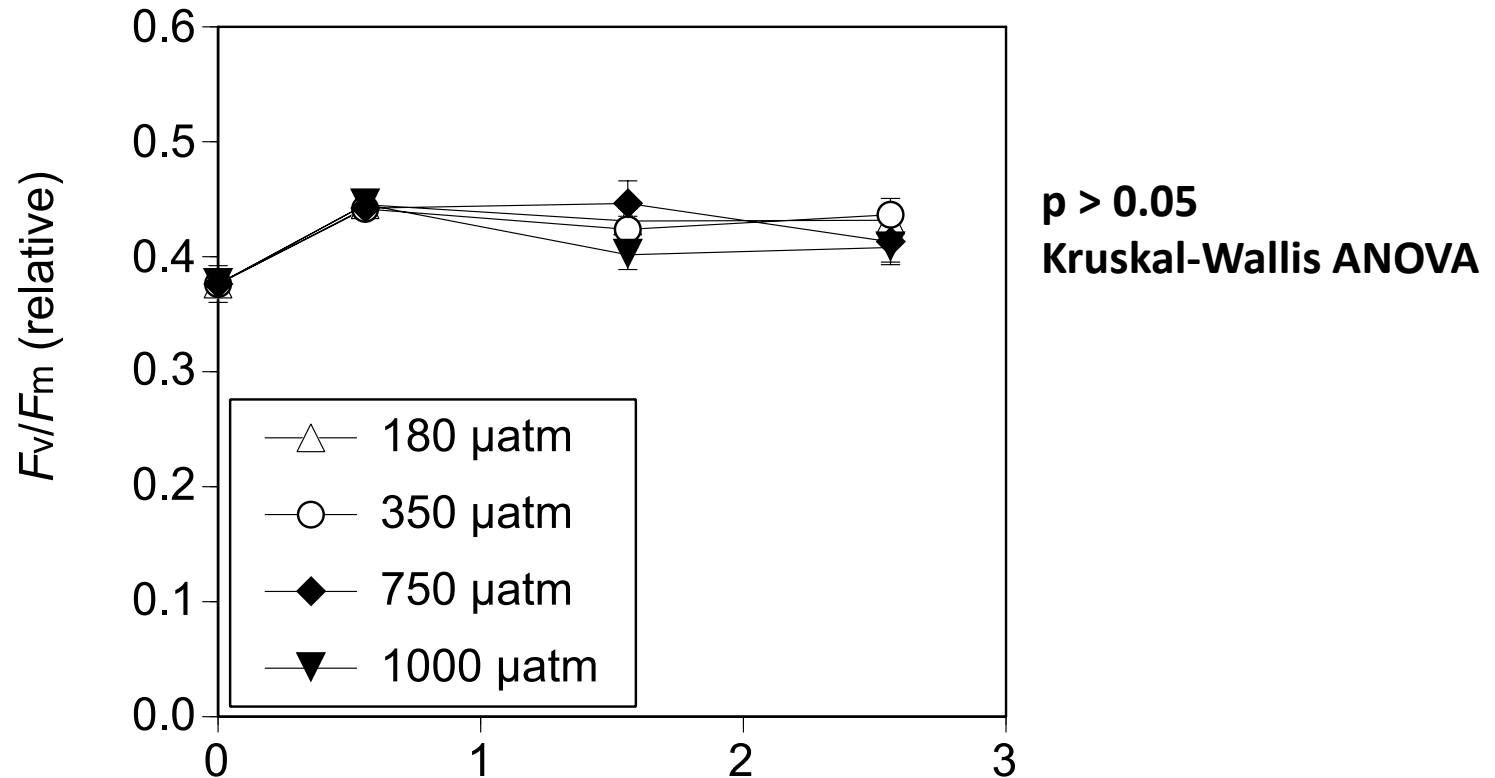


The vulnerable diatom families suggested



The predominant diatom family suggested

Changes over time in the photosynthetic competence (F_v/F_m) of phytoplankton among CO₂ treatments during incubation as determined by FRe fluorometry



The results indicate the photosystem II activity of the phytoplankton assemblages was little affected by CO₂ availability.

Summary

We investigated the impact of different CO₂ levels on spring diatoms in Oyashio waters of the western North Pacific.

- Net growth rates of fucoxanthin, a diatom marker, decreased at higher CO₂ levels during incubation.
- Diatom-specific *rbcL* DNA copies can also become an indicator of diatom biomass in the study area.
- Diatom-specific *rbcL* DNA and cDNA analyses revealed that Chaetocerataceae, Thalassiosiraceae and Fragilariaceae might be vulnerable in the high CO₂ world expected in the near future, whereas Bacillariaceae could be a strong group against CO₂ changes.

Thank you!

