



Primary productivity induced by iron and nitrogen in the Tasman Sea – An overview of the PINTS expedition

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Primary productivity induced by iron and nitrogen in the Tasman Sea – An overview of the PINTS expedition.

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Abstract:

The Tasman Sea and the adjacent Sub-Antarctic zone (SAZ) are economically important regions, where the parameters controlling the phytoplankton community composition and carbon fixation are not yet fully resolved. Contrasting nutrient distributions as well as phytoplankton biomass, biodiversity and productivity we observed between the North Tasman Sea and the SAZ. In-situ F_V/F_M , dissolved and particulate nutrients, iron biological uptake, and nitrogen and carbon fixation were used to determine the factor limiting phytoplankton growth and productivity in the North Tasman Sea and the SAZ. Highly productive cyanobacteria dominated the North Tasman Sea. High atmospheric nitrogen fixation and low nitrate dissolved concentrations indicated that non-diazotroph phytoplankton are nitrogen limited. Deck-board incubations also suggested that, at depth, iron could limit eukaryotes, but not cyanobacteria in that region. In the SAZ, the phytoplankton community was dominated by a bloom of haptophytes. The low productivity in the SAZ was mainly explained by light limitation, but nitrogen, silicic acid as well as iron were all depleted to the extent that they could become co-limiting. This study illustrates the challenge associated with identification of the limiting nutrient as it varied between phytoplankton groups, depths and sites.

Keywords: nutrients, limitation, subantarctic zone, phytoplankton, bioavailability

Running Head: Nutrient and light limitation in the Tasman Sea

1 **Introduction:**

2
3 The oceans absorb between 30 and 40% of annually emitted anthropogenic carbon dioxide
4 (CO₂; Battle et al. 2000), representing an important role in Earth's climate system.
5 Photosynthetic micro-organisms responsible for biological carbon fixation are also acting
6 as marine ecosystem engineers, regulating ocean chemistry and driving the
7 biogeochemical cycles of many key elements including, nitrogen, phosphorus, silicic acid,
8 iron, sulfur and carbon (Karl 2000; Chisholm 2000; Morel and Price 2003; LeBlanc et al.
9 2005; Moore et al. 2009; Boyd and Ellwood 2010). Phytoplankton therefore play a central
10 role in the metabolism of marine systems.

11
12 Nutrient limitation imposed by low concentrations of iron (Fe) and nitrogen as nitrate in
13 surface waters affects the growth of phytoplankton in up to 50% of the world's oceans,
14 including low-nutrient, low-chlorophyll (e.g., the North Atlantic and North Tasman Sea;
15 Moore et al. 2009; Law et al. 2011), and high-nutrient, low-chlorophyll regions (e.g., the
16 Southern Ocean; Martin et al. 1990; Boyd and Ellwood 2010), as well as coastal areas
17 (e.g., the California upwelling region; Hutchins and Bruland 1998). The widespread
18 significance of these two nutrients is due to their low concentrations in surface waters,
19 their biological importance and also their biogeochemical "coupling", with Fe being
20 required for atmospheric N₂ fixation and nitrate assimilation (Morel and Price 2003).
21 Indeed, Moore and Doney (2007) have suggested that "iron is the ultimate limiting nutrient
22 leading to nitrogen being the proximate limiting nutrient over wide regions today".
23 Vertical supply of Fe and nitrate in the mixed layer, although not the sole source, can be
24 important to sustain phytoplankton growth (e.g., Bowie et al. 2009; Law et al. 2011). With
25 predictions of shallower mixed layer depths in the future (IPCC 2007), the impact that Fe,
26 N and light availability have on phytoplankton growth and abundance needs to be
27 addressed in the context of present day oceans.

28
29 The Tasman Sea and the SAZ are both important regions for Australian biodiversity and
30 socio-economy. The Tasman Sea is a hot spot for climate change with one of the highest
31 predicted oceanic temperature increases due to changes in the strength of major boundary
32 currents (East Australian Current, EAC; Hobday et al. 2008). In addition, contrasting
33 concentrations of nutrients are reported, with macronutrient-(N, P and Si)-poor, warmer
34 subtropical waters in the North (e.g., Ellwood et al. 2013) grading to macronutrient-rich,

colder subantarctic waters in the South (e.g., Bowie et al. 2011). Previous studies also demonstrated a greater dissolved iron concentration in the North Tasman Sea compared to the SAZ (Boyd and Ellwood 2010; Lannuzel et al. 2011; Ellwood et al. 2013). The global ecosystem model developed by Moore and co-workers (Moore and Abbott 2000; Moore et al. 2002) predicts that nutrient distributions and their delivery to the mixed layer of this region would result in contrasting regional phytoplankton nutrient limitation leading to differences in their biomass, biodiversity, primary productivity, and carbon export during the austral summer. Recent studies demonstrated that in the oligotrophic North Tasman Sea, phytoplankton are mainly limited by the lack of nitrogen (Ellwood et al. 2013) with diazotrophs, able to bypass nitrate limitation by fixation of atmospheric N_2 , present in surface waters (Law et al. 2011). However, it has been suggested that the growth of diazotrophs in Northern Australia waters and the North Tasman Sea is mostly iron-limited (Kutska et al. 2003; Law et al. 2011). In the SAZ east of Australia, phytoplankton groups are predicted to be mainly limited here by Fe, with Si limitation possible for diatoms (Moore et al. 2002). The region spanning the North Tasman Sea to the SAZ is thus ideal for the study of the influence of iron and nitrogen limitation on phytoplankton communities.

This paper presents selected results from the PINTS ('Primary productivity induced by Iron and Nitrogen in the Tasman Sea') research voyage which took place during the austral summer of 2010. This project identified the distribution of Fe and N and other nutrients from the oligotrophic North Tasman Sea to the highly productive SAZ region in the South Tasman Sea. By comparing the distribution of nutrients with resident phytoplankton communities, and by relating these to key processes at play, such as N_2 fixation, Fe bioavailability, photophysiology and primary productivity, we investigated the control that nutrients and light impose on phytoplankton growth, biodiversity and carbon fixation. This study thus evaluates the distribution of Fe as a key essential trace element, including sink and source processes that affect both its distribution and biogeochemical cycling, and also contributes to the international GEOTRACES program in the Tasman Sea, a region identified as an area of interest for coarse-resolution global surveys of ocean surface waters in the GEOTRACES Science Plan (2006, http://www.geotraces.org/libraries/documents/Science_plan.pdf).

Methods:

Study region

During the PINTS voyage, 18 stations were sampled within the Tasman Sea during the austral summer 2010 (23rd Jan. – 15th Feb. 2010, *RV Southern Surveyor*, Table 1, Fig. 1). The voyage track consisted of three transects, with Transect 1 comprising five stations from Sydney to the oligotrophic North Tasman Sea (P1, 30°S). It crossed both the narrow abyssal basin (4500-m) off Sydney and the Lord Howe Rise (LHR) which stretches northward through the central Tasman. Transect 2 was a meridional transect from the Northern Tasman Sea (P1) to the SAZ (46.2°S, P3) recrossing the LHR at 35°S and extending again into abyssal waters in the south, with eight stations along its length. Transect 3 comprised seven stations extending zonally from the SAZ (P3) to the Tasman shelf (station [Stn] 19) crossing the abyssal plain.

Here, we place the PINTS voyage in the context of the regional circulation. Fig. 2a & b show the general structure of the mean EAC flow off eastern Australia as determined from multi-year ocean measurements (Ridgway et al. 2002). Transect 1 was aligned roughly along the core of the meandering eastward flow of the Tasman Front (TF; Godfrey et al. 1980). It is the portion of the EAC that separates from the western boundary at 31–34°S and extends across the Tasman Sea to the northern tip of New Zealand, forming the East Auckland Current (Fig. 2a, c). The transect also passed through a significant recirculation feature contained within the abyssal waters off Sydney (Fig. 2c). Fig. 2b indicates that from the coast to 160°E the region is very turbulent, dominated by mesoscale eddies, which have a strong signature in surface pressure and thus sea level. These eddies are 200–300 km in diameter and two or three eddies are generated annually and have lifetimes often exceeding a year (Nilsson and Cresswell 1981; Bowen et al. 2005). Fig. 2 shows that Transect 2 traversed the TF and a further series of relatively weak eastward components (Ridgway and Dunn 2003). It also passed through different water masses from subtropical waters north of the TF through to the northern edge of the Subtropical Convergence. The transect was just at the eastern edge of the region of high sea level variability (Fig. 2b), but there was still sufficient mixing to reduce any sharp frontal regions. Finally, Transect 3 encountered mostly quiescent conditions until it reached within 200-km of the Tasmanian east coast, where it met the southern extent of the EAC eddy field and the residual southward component of the EAC (EAC Extension). This latter feature turns both

eastwards southeast of Tasmania towards New Zealand (Bowie et al. 2011) and westward south of Tasmania – the Tasman Outflow – feeding the continental boundary current south of Australia and eventually is incorporated into the southern Indian Ocean gyre (Ridgway and Dunn 2007).

Water sampling

Salinity, temperature, fluorescence and oxygen were obtained from calibrated conductivity-temperature-depth (CTD, SeaBird SBE11) data using water collected from Niskin bottles (General Oceanics) (Reynolds and Navidad 2012). Macronutrients (reactive phosphorus, PO_4 ; silicic acid, Si; nitrate-plus-nitrite, NO_x) were obtained by the shipboard analysis of unfiltered water using an automated flow-injection analyser and colorimetric techniques (Reynolds and Navidad 2012). Analytical detection limits were 0.011, 0.013 and 0.016 μM for NO_x , PO_4 and Si, respectively (A. Navidad, pers. comm.). Samples were also collected from the CTD rosette for the determination of pigments concentration and composition, photophysiology, primary productivity, and N_2 fixation rate.

Since Fe and most of the other micronutrient elements are trace metals omnipresent in shipboard environments, samples for almost all the procedures below required the use of accepted and verified non-contaminating sampling procedures. In this case water was sampled directly after the deployment of the CTD (within 30–60 min of CTD recovery) using Teflon-coated Niskin X-1010 bottles (General Oceanics, USA) mounted on an autonomous rosette (Model 1018, General Oceanics, USA) and deployed using a Dynex hydroline (*Dynex Dyneema 75*, Hampidjan, Ltd, Nelson, New Zealand). Water samples for trace micronutrient analysis were collected and filtered (Pall, Acropack 200) in a clean van under a HEPA filter (ISO Class 5 conditions). Unfiltered seawater was also sampled for macronutrient analysis, and the determination of the iron biological uptake rate by the natural phytoplankton community. Sample acquisition and handling was as per GEOTRACES recommendations (Cutter et al. 2010; www.geotraces.org/libraries/documents/Intercalibration/Cookbook.pdf), using acid-washed non-contaminating material as detailed elsewhere (O'Sullivan et al. 1995; Bowie et al. 2009; Lannuzel et al. 2011). Samples were acidified using quartz-distilled HCl (qHCl, Seastar Baseline, Canada) to 0.02 M qHCl and kept for six months double-bagged in plastic boxes in clean storage, prior to analysis.

Synthetic Temperature-Salinity - synTS

To provide further regional context for the cruise sections, we obtained temperature and salinity fields generated from surface height (altimetry) and SST from satellite data (synTS; Ridgway et al. 2008). The system uses a multiple linear regression model obtained from historical in-situ observations.

Dissolved metal analysis

Dissolved iron (dFe) was determined by flow injection analysis with chemiluminescence detection (FI-CL) using in-line preconcentration on a TOYOpearl-8HQ chelating resin column adapted from Obata et al. (1993) and de Jong et al. (1998). Accuracy was checked using an in-house quality control (QC) standard and by analysing the surface standard (S1) collected during the SAFe cruise (Johnson et al. 2007) ($S1 = 0.109 \pm 0.007$ nM compared to the 'certified' value of 0.097 ± 0.043 nM). The mean limit of detection (LOD) was 0.04 ± 0.04 ($n=21$) for a 3 minute load. An automated method of flow injection – solid-phase extraction – inductively coupled plasma-mass spectrometry (FI-SPE-ICPMS) was used for determining dissolved Cd as detailed in O'Sullivan et al. (2013). Detection limit and precision of Cd analysis was of 5 pM and 3 %, respectively. Accuracy was checked using certified standard (SAFe D2) and reference seawater (NASS-5, National Research Council of Canada). Value of 904 ± 41 ($n=11$) and 190 ± 11 ($n=11$) pM of Cd were obtained for SAFe D2 and NASS-5, respectively. Values were within the certified range 1011 ± 28 pM and 205 ± 30 pM for SAFe D2 (<http://es.ucsc.edu/~kbruland/GeotracesSaFe/kwbGeotracesSaFe.html>, Nov. 2011) and NASS-5, respectively.

Particulate metal analysis

Suspended particulate trace metal samples were collected in situ using a 142 mm diameter stack consisting of a 54 μ m polyester pre-filter screen (Sefar Filtration, Inc.) followed by a Whatman QMA filter (nominal pore size 1 μ m), following methods outlined in Bowie et al. (2009). Filter stacks were housed in large volume McLane pumps (McLane Research Laboratories Inc., USA) suspended on plastic-sheathed steel mooring wire. Filters and supports were prepared and pre-cleaned as described in Bishop et al. (1985). Upon recovery of the pumps, filters were dried under a laminar flow bench and sub-samples taken using a circular plastic punch (15 mm diameter). Filters were stored frozen at -20 °C in acid-washed petri dishes until further analysis in the home laboratory. Blanks were true

1 'field' blanks that were deployed in situ (i.e., filter stacks that no seawater had passed
2 through on the McLane pumps).

3 Sampled particles were acid extracted (1 mL concentrated HNO₃ for 12 h at 95 °C) in
4 Teflon PFA vials and analysed by magnetic sector inductively coupled plasma mass
5 spectrometry (Finnigan ELEMENT 2) offering high spectral resolution capability,
6 following methods reported in Cullen and Sherrell (1999), Townsend (2000), and Bowie et
7 al. (2010). Recoveries for Fe were verified by analysing a certified reference material for
8 marine and plankton (Community Bureau of Reference BCR-414, with a mean recovery of
9 89% % for Fe ($n=3$) using HNO₃ only as the digestion solution.

11 ***POC/ PON analysis***

12 Water collected from the CTD was filtered through muffled quartz 1µm filters (450 °C for
13 five hours, Sartorius). The filters were then de-carbonated by fuming overnight in
14 hydrochloric acid and then drying for 48 h at 60 °C in a clean oven, to drive off excess
15 acid and ensure dryness. The active area of the filter (22.5 mm dia.) was subsampled, in a
16 class 1 laminar flow cabinet (Gelman CF43S), with a single 18 mm punch. The filter
17 punching was then encapsulated in silver 9 × 5mm capsules (Sercon SC0037) as
18 preparation for elemental analysis. Unused muffled filters were processed similarly to use
19 as an analytical blank. The capsules were stored over silica gel prior to analysis. McLane
20 filter subsamples (13.8 mm subsample of 133 mm active area) were transferred directly to
21 capsules and de-carbonated by acidification with two 20 µL aliquots of 2 M HCl (King et
22 al. 1998), followed by drying for 48 h before crimping and compressing the capsules.
23 Certified casein (OAS Elemental Microanalysis) was included with the samples as a
24 further analytical quality control.

26 The POC/ PON analysis was performed using a gas chromatography elemental analyser
27 (Thermal Finnigan Flash 1112 series catalytic combustion) run in standard conditions with
28 reference to acetanilide standards in the Central Sciences Laboratory of the University of
29 Tasmania. Precision of the analyses was 1% (Dr. Thomas Rodemann, pers. comm.).

31 ***Phytoplankton pigments***

32 Pigment samples were collected on GF/F filters (Whatman) using gentle filtration (< 5 mm
33 Hg) and stored in cryo-vials in liquid nitrogen prior to being analysed back on shore. The

1 pigments were extracted in 100% methanol in the dark and at 4 °C and analysed by HPLC
2 using a Waters – Alliance high performance liquid chromatography system, comprising a
3 2695XE separations module with column heater and refrigerated autosampler and a 2996
4 photo-diode array detector following a modified version of the Van Heukelem and Thomas
5 (2001) method (see Hassler et al. 2012a for details). Concentrations of chlorophyll-*a* (Chl-
6 *a*), chlorophyll-*b* (Chl-*b*) and β,β -carotene in sample chromatograms were determined
7 from standards from Sigma, while all other pigment concentrations were determined from
8 standards from DHI (Denmark). The presence or absence of pigments which relate
9 specifically to an algal class, termed biomarkers, was used to provide a simple guide to
10 the composition of a microalgal community, including identifying classes of small
11 flagellates that could not be determined by light microscopy techniques (Jeffrey and
12 Wright, 2006). Biomarker pigments analysed were Fucoxanthin (Fuco), 19-
13 Butanoyloxyfucoxanthin (But-Fuco), 19 Hexanoyloxyfucoxanthin (Hex-Fuco), Zeaxanthin
14 (Zea), Chlorophyll *b* (Chl-*b*), Peridinin (Peri) and Divinyl Chlorophyll-*a* (DVChl-*a*).
15

16 ***Flow cytometric enumeration of picoplankton***

17 Samples for enumeration of bacteria and cyanobacteria were fixed with 1% (v/v final
18 concentration) glutaraldehyde, frozen in liquid nitrogen and stored at -80 °C. Bacterial
19 enumeration was performed on samples stained with SYBR Green I (1:10,000)
20 (Invitrogen) (Marie et al. 1997), while cyanobacterial analysis was performed using
21 unstained samples. Samples were analysed using a Becton Dickinson LSR II flow
22 cytometer. Bacterial populations were discriminated according to SYBR Green
23 fluorescence and side-scatter (SSC) (Marie et al. 1997), while populations of
24 *Prochlorococcus* and *Synechococcus* were discriminated using SSC and red and orange
25 fluorescence. Data was analysed using Cell-Quest Pro (BD Biosciences) as in Seymour et
26 al. (2012).
27

28 ***Photophysiological community analysis***

29 To assess the photosynthetic efficiency (F_v/F_m) of phytoplankton community in-situ, a
30 prototype submersible Pulse Amplitude Modulated Water PAM (sub-Water PAM,
31 Gademan Instruments GmbH, Würzburg, Germany) fluorometer was deployed with the
32 CTD to a maximum depth of 100 m. The instrument had two channels (red and blue LED,
33 with wavelengths of 620 and 480 nm, respectively) to collect data on the photosynthetic
34 activity of the total phototrophic community and the total minus cyanobacteria community,

respectively. The sub-Water PAM was programmed to pump water for 10 s to flush the dark chamber every 5 min. Profiles were conducted by lowering the instrument at 10 m intervals and maintaining position for 10 min. At each depth, a saturating pulse of light (pulse duration = 0.6 s; pulse intensity > 3000 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$) illuminated the sample and maximum fluorescence (F_M) was determined. Maximum quantum yield of photosystem II (PSII), F_V/F_M , was then calculated according to the equation $(F_M - F_0)/F_M$ (Schreiber 2004).

¹⁴C productivity

Radiolabeled sodium bicarbonate ($\text{NaH}^{14}\text{CO}_3$, Perkin Elmer) was added to freshly sampled surface and subsurface chlorophyll-a maximum seawater to produce a working solution of $39.183 \times 10^3 \text{ Bq per mL}$ ($1.1 \mu\text{C mL}^{-1}$). Primary productivity incubations were conducted as described in Doblin et al. (2011), based on the small bottle ^{14}C technique of Lewis and Smith (1983). Briefly, a working solution was initially sampled to determine initial radioactivity, then triplicate 7 mL aliquots were each incubated in glass vials for 1 hour in the dark and at 2 light intensities at in-situ temperature ($\pm 0.5^\circ\text{C}$) controlled using a water-bath, or for surface seawater, using running seawater pumped from a depth of 5 m. Inorganic carbon was degassed by the addition of 250 μL of 6 M HCl (Technical grade, Crown Scientific) and agitated for 3 h prior to addition of 10 mL scintillation cocktail (Aquasure, Perkin Elmer) and counting using a TriCarb 2900 scintillation counter (Packard, Waltham MA, USA). Counts per minute were transformed into disintegration per minute using the built-in quench curve and radioactivity was expressed in carbon fixed per volume per hour using initial radioactivity and dissolved inorganic carbon (DIC) analysed using coulometric procedures (Johnson et al. 1998). Carbon uptake rates were standardised with in-situ Chl-*a* concentrations measured using HPLC.

Iron bioavailability

Phytoplankton iron uptake rates ($\text{amol Fe } \mu\text{g Chl-}a^{-1}$) were used to estimate iron bioavailability as per Hassler and Schoemann (2009a). One litre of unfiltered seawater, freshly collected from the depth of maximal fluorescence, was spiked with 2 nM $^{55}\text{FeCl}_3$ (Perkin Elmer, 20.82 mCi/mg Fe at the time of use). Solutions were sub-sampled (2 mL) to measure total initial radioactivity, then incubated for 20-24 h at in-situ temperature and light as determined from CTD deployments. Following incubation, the phytoplankton was gently sequentially filtered (< 5 mm Hg, 10 μm polycarbonate filter [Millipore], then 2 μm

1 and 0.7 μm glass fiber filters [MicroAnalytix]) and rinsed ($5 \times 3 \text{ mL}$, total contact time 10
2 min.) with an EDTA-oxalate solution to discard the adsorbed iron (Hassler and Schoemann
3 2009b). The filter was then rinsed with filtered seawater ($3 \times 3 \text{ mL}$, $< 0.7 \mu\text{m}$), collected in
4 scintillation vials, supplemented with 10 mL of scintillation cocktail (Ultima Gold, Perkin
5 Elmer) and then analysed using a liquid scintillation counter (Perkin Elmer, Tricarb 2900).
6 Intracellular disintegrations per minute were transformed into iron concentration using a
7 custom made quench curve, total initial radioactivity and dissolved iron concentration (see
8 Hassler and Schoemann 2009a for details). Pigment analyses were performed (see above)
9 in order to normalise iron uptake rate by phytoplankton biomass and investigate
10 differences in the iron uptake rate for the different dominant taxa in the micro- ($> 10 \mu\text{m}$),
11 nano- (2-10 μm), and pico-plankton (0.7-2 μm). Incubations were done in duplicate.

13 *Nitrogen fixation*

14 2.4-Litres were sub-sampled from the Niskin bottles into acid pre-washed polycarbonate
15 bottles (Nalgene). 4.5 mL of 99% $^{15}\text{N-N}_2$ was then injected through a butyl rubber septum
16 in the lid, followed by gentle rotation to encourage dissolution and subsequent incubation
17 for 24 hours in on-deck incubators (as per Law et al. 2011). The incubations were
18 terminated after 24 hours by filtration under low negative pressure onto pre-ashed 25 mm
19 0.7 μm Whatman GF/F filters, which were stored in a -20°C freezer. Background ^{15}N -
20 PON natural abundance was determined by filtration of 2.4 L surface seawater that was not
21 spiked with $^{15}\text{N-N}_2$. All filters were subsequently dried and analysed for ^{15}N incorporation
22 into PON by stable isotope mass spectrometry using an NA1500 elemental analyser (Fisons
23 Instruments) interfaced to a Delta^{plus} (Thermo-Finnigan) continuous flow isotope ratio mass
24 spectrometer (CF-IRMS), with nitrogen fixation rates determined using the calculations of
25 Montoya et al. (1996). The calculations were based on a dissolved concentration of 1.14
26 $\pm 0.09 \text{ mL/L}$, as reported in Law et al. (2011), following identification that dissolution of
27 $^{15}\text{N-N}_2$ was not instantaneous (Mohr et al. 2010).

29 **Results:**

31 *Oceanography – Syn TS*

32 The temperature along the PINTS track obtained from SynTS is compared with the
33 observed data from the CTD stations in Fig. 3. The section shows the temperature

1 anomaly (the mean field from a climatology (CARS; Ridgway et al. 2002) has been
2 removed). The SynTS result is in good agreement along the Transect 1, capturing the
3 main structure at the right magnitude. Along Transects 2 and 3, the comparison is not
4 quite as good. While the inferred SynTS temperature does show a similar pattern of warm
5 and cool anomalies along the track the magnitude of the features is far smaller. For
6 example, there was a strong negative feature ($< -4^{\circ}\text{C}$) in the upper 125 m at cast 8 in the
7 observed section, which is only weakly negative in the SynTS result. However, despite the
8 smaller amplitude of the SynTS features these comparisons suggest that the SynTS fields
9 will provide a very useful summary of the circulation at the time of the voyage.

10
11 The SynTS fields of T and S in Fig. 4 show the main features in the region during the
12 cruise. We see that the cruise track passed through the highly energetic EAC eddy field.
13 Steaming out of Sydney the track crossed a large cold-core cyclonic eddy and then passed
14 between the separated EAC flow (wrapped around an anti-cyclonic feature) to the north
15 and another warm-core anti-cyclonic eddy that had almost completed ‘pinching off’ from
16 the main EAC structure. The track then passed through the centre of another anti-cyclonic
17 eddy (158°E) although the nearest station (Stn 3) was located on the edge of the eddy. The
18 eastern transect passed through a train of less energetic anti-cyclonic and cyclonic features
19 from 30 to 38°S , although the stations tended to be on the periphery of the eddies. Moving
20 southwards, the transect passed from a higher to lower salinity environment with a distinct
21 salinity front from 43 – 45°S (Fig. 4). Even at 1000 m where salinity gradients were small
22 there was a frontal region representing the boundary between Antarctic Intermediate Water
23 (AAIW) of southern origin and the higher salinity northern variant which has been
24 modified as it is advected around the South Pacific Gyre (Sokolov and Rintoul 2000). On
25 the southern transect, the voyage moved into the SAZ and the track passed through a
26 region of meandering filaments from the north and south. Finally as it approached eastern
27 Tasmania it again crossed the EAC sourced waters associated with a seasonal warm-core
28 eddy at 148°E (Ridgway 2007).

30 ***Oceanography – Temperature, Salinity, Oxygen***

31 The salinity, temperature and oxygen from the CTD stations occupied along the PINTS
32 voyage track are presented in Fig. 5. In the following presentation of the voyage results
33 we recognize that the sampling interval of the CTDs was not able to completely resolve the
34 highly energetic EAC eddy field. However, the sections did clearly show that the track

sampled two distinct oceanic regions – a warm, salty, low oxygen water mass associated with the EAC that enters the Tasman Sea from the north and east of the South Pacific Gyre (Sokolov and Rintoul 2000) and a cool, fresh, high oxygen water mass that is sourced from the subantarctic region in the south. The transition between these two regions occurred fairly sharply at 44°S which represents the Sub-Tropical Front (STF). This tends to extend slightly northward between Tasmania and New Zealand (Orsi et al. 1995; Ridgway and Dunn 2007). The *T* and *S* sections also indicated that the track crossed another frontal zone between 32 and 35°S, the Tasman Front, which represents the separated EAC flow meandering eastward across the Tasman Sea (Figs. 4c-d; Godfrey et al. 1980). An anomaly section of *T* (not shown) indicated that the summer mixed layer penetrates to 20–30 m in the north and east and from 50–60 m along the southern transect (Transect 3).

The cold-core eddy off Sydney at the start of the track had a distinct subsurface expression, with low *T* and *S* from 60–900 m and also a smaller cell of low oxygen from 50–150 m. Station 3 showed the typical eddy induced deepening of the thermocline despite it being located to the west of the eddy centre. There was no corresponding salinity signature, because the salinity was well mixed down to about 150 m north of the Tasman Front (Fig. 4). In fact, Fig. 5 showed a uniform high salinity feature in this region with the maximum in salinity occurring in the northeastern corner of the cruise track (Stns 4–6). The Tasman Front was shown by a subsurface frontal region in both *T* and *S* between 50–250 m (Fig. 5). There was also a strong expression in both *T* and *S* at the STF in the upper 400 m (44°S) which extended down to about 400 m. Along the southern Transect 3 both *T* and *S* showed an alternating series of high and low features extending to 400 m. They features identified the cross-frontal exchange of properties, northward intrusions of cool, fresh, subantarctic water and warm, salty water from the north. East of Tasmania, at the end of the voyage track, the high *T* and *S* associated with the anti-cyclonic eddy projected down to more than 1000 m.

The dissolved oxygen section (DO, Fig. 5c) showed a high degree of structure along the path. The magnitude and vertical extent of DO increased uniformly – the depth of the 220 μ M contour increased from 50 m east of Sydney to 800 m southeast of Tasmania. In fact the advection of high DO water into the southern Tasman Sea was associated with cross-frontal processes indicated by the Subantarctic intrusions at 152°E (Figure 5). The DO signature of these intrusions extended deeper than that of *T* and *S* (> 800 m). A thin

1 filament (40 m) of high DO water at 60 m depth was sampled along the section from 43°S
2 to the Sydney endpoint. It was apparently formed in the south Tasman (at the surface) and
3 spreads northwestwards along the 25.4 isopycnal.

4 5 ***Macronutrients***

6 Although the distance between CTD deployments was too great to efficiently resolve the
7 oceanography in Transect 1 and 2 (see above), the effect of mesoscale features can be seen
8 on macronutrient distributions, with doming of isolines along Transect 1 (Fig. 6 and Suppl.
9 Fig. 2). In general, for a given depth along Transect 1, the concentrations of NO_x , PO_4 and
10 Si decreased eastwards (P1, Stn 5). At P1, the surface water was strongly depleted in
11 macronutrients ($\text{NO}_x < 0.011 \mu\text{M}$, PO_4 0.02–0.12 μM , $\text{Si} < 0.7\text{--}1.0 \mu\text{M}$). Indeed
12 concentrations of NO_x using a nanomolar analytical method were on average 8.5, 9.2, and
13 9.8 nM at 15, 30 and 50 m depth respectively (M. Woodward, pers. comm.). Along
14 Transect 2, macronutrient concentrations generally increased from North (P1) to South
15 (P3, Fig. 6, Suppl. Fig. 2). A doming of all macronutrients in surface water north and in
16 the STF was observed from Stn 7 to 11 with depleted concentrations at P3 (Suppl. Fig. 2).
17 As previously observed (Ellwood et al., 2013), despite crossing the Tasman Front, no
18 frontal footprint in macronutrient concentrations was observed (down to 1000 m depth).
19 Along transect 3, macronutrient concentrations increased towards Tasmania, but remained
20 lower at depth (1000 m) than measured along Transects 1 and 2 (Fig. 6). As for Transect 1,
21 mesoscale features affected macronutrient distribution, resulting in doming isoclines.
22 Close to Tasmania (Stns. 17 and 18), surface water was depleted in NO_x and PO_4 , but not
23 Si, possibly resulting in a heterogeneous diatom distribution along Transect 3 (Suppl. Fig.
24 2). Considering all data, the dissolved NO_x : PO_4 disappearance ratio was 14.7 ± 0.1 ($r^2 =$
25 0.98 , $n = 386$).

26 27 ***Organic particles***

28 Total Chl-*a* concentration was used as a surrogate of phytoplankton biomass. Patchiness of
29 Chl-*a* was measured along all transects (Fig. 7a). Overall, the total Chl-*a* in surface water
30 (0–125 m) was well correlated with the fluorescence data from CTD deployments (Suppl.
31 Fig. 3) and increased along the cruise track, with maximum values observed in the SAZ
32 region (P3 and Stn 14). Along Transect 1, the depth of the maximum Chl-*a* decreased from
33 40 m to 110 m with very low Chl-*a* concentrations for the top 50 m ($0.2 \mu\text{g L}^{-1}$ isocline) at

P1. Close to the LHR (Stn 3), two distinct Chl-*a* maxima were measured. Travelling southwards from P1 to P3, the total Chl-*a* in surface water increased with a shallower depth of Chl-*a* maximum. At P3, the maximum Chl-*a* was found in surface waters (top 25 m). Moving west along Transect 3, the total Chl-*a* in surface water decreased toward Tasmania and the Chl-*a* maximum deepened from 25 m (P3) to 40–50 m (Stn 18).

POC and PON were measured in surface water (25 m) and at the depth of the maximal fluorescence (Suppl. Fig. 3) along the cruise track. As expected, the ratio Chl-*a*: POC and Chl-*a*: PON was statistically lower ($p < 0.001$) in surface waters (0.30 ± 0.09 ; $0.03 \pm 0.01 \mu\text{g } \mu\text{mol}^{-1}$, $n = 17$, respectively) than at the depth of maximum fluorescence (0.73 ± 0.41 ; $0.10 \pm 0.04 \mu\text{g } \mu\text{mol}^{-1}$, $n = 18$, respectively). At the depth of maximum fluorescence, the Chl-*a*: POC ratio was greater ($> 0.65 \mu\text{g } \mu\text{mol}^{-1}$) at Stns 2, 5–8, 9–10 and 16–18, coinciding with high Chl-*a*: PON ($> 0.09 \mu\text{g } \mu\text{mol}^{-1}$). Particulate material at the surface showed a different pattern (data not shown). The greatest ratios for Chl-*a*: POC and Chl-*a*: PON at the depth of fluorescence maximum were observed at Stn 5 ($1.99 \mu\text{g } \mu\text{mol}^{-1}$), and Stns 2, 16 and 18 ($0.12 \mu\text{g } \mu\text{mol}^{-1}$), respectively. The lowest ratios for Chl-*a*: POC and Chl-*a*: PON at the depth of fluorescence maximum were observed at Stns 4 and 13 (0.37 – $0.38 \mu\text{g } \mu\text{mol}^{-1}$) and Stn 4 ($0.04 \mu\text{g } \mu\text{mol}^{-1}$).

The POC: PON ratio (or C/N, Fig. 7b) was used to indicate a potential nitrogen (N) limitation. A ratio of C/N above the optimal Redfield ratio (6.6; Redfield et al. 1963) indicates a deficit of nitrogen relative to carbon in organic matter, possibly related to N limitation. In the study region, the ratios C/N were in general greater in surface water. The C/N ratios were close to the Redfield ratio from Stns 12 to 15 in surface water (25 m), and at Stn 4 and Stns 10 to 17 at the depth of maximal fluorescence. An extremely high value (16.5) was measured in surface waters at Stn 7. C/N was weakly correlated to Chl-*a* ($r^2 = 0.42$, $n = 29$) for C/N value < 10.0 (4 points were excluded, all of which had Chl-*a* $< 0.2 \mu\text{g/L}$). The negative slope (-0.19 ± 0.04), suggests that greater phytoplankton growth were observed at lower C/N ratio.

Comparison between North Tasman Sea (P1) and SAZ (P3)

Overall for the study region, the greatest differences in fluorescence (and Chl-*a*), macronutrient profiles, as well as C/N were observed in Transect 2 between P1, the North

1 Tasman Sea, and P3, the Sub-Antarctic Zone. Below, we present more detailed results to
2 investigate differences in phytoplankton abundance, biodiversity and productivity in
3 relation to both macro- and micro-nutrients at these two sites.

4
5 *Nutrient distribution*- The concentrations of the macro- (NO_x , PO_4 , Si) and micro-nutrient
6 Cd were greater in surface water and at the depth of maximum fluorescence for P3
7 compared to P1 (Fig. 8, and Suppl. Fig 4). Only for Fe, the concentrations in surface
8 waters and at depths of maximum fluorescence were smaller at P3. For depths greater than
9 2000 m, nutrient concentrations were greater at P1, suggesting differences in the source,
10 sink and transport of nutrients for these two sites.

11
12 The Fe^* was calculated to discuss potential dFe limitation for phytoplankton growth. The
13 Fe^* at a given site and depth compares the concentrations of dFe and PO_4 concentrations
14 with respect to Fe and P phytoplankton cellular quota ($R_{\text{Fe:P}}$). Fe^* equals $[\text{dFe}] - R_{\text{Fe:P}} \times$
15 $[\text{PO}_4]$, and an average phytoplankton $R_{\text{Fe:P}}$ was considered (0.47 mmol: mol; Parekh et al.
16 2005). A negative Fe^* is indicative of Fe limitation (Parekh et al. 2005; Bowie et al. 2009;
17 Lannuzel et al. 2011). Here, Fe^* decreased with depth from 0.29 at 15 m to -0.01 at 95 m
18 for P1 and from 0.09 at 15 m to -0.06 at 80 m at P3. Negative Fe^* values for the top 100 m
19 were only observed at 95 m and 80 m at P1 and P3, respectively.

20
21 The disappearance ratios Cd: PO_4 , PO_4 : NO_x , Fe: PO_4 were similar considering all data
22 from both sites (Fig. 9a – c). A “kink” was observed in the Cd: PO_4 relationship for
23 dissolved concentrations of $\text{PO}_4 < 0.8 \mu\text{M}$, comprising data from 0–125 m at P1 and P3.
24 Fe: NO_x and Fe: Si showed a typical “scavenging” relationship for Fe (Fig. 9e and 9g).
25 When considering only data from 0–125 m, the euphotic zone inhabited by phytoplankton,
26 differences in disappearance ratios for Fe: PO_4 , Fe: Si and Fe: NO_x were observed between
27 stations (Fig. 9 d, f, and h, respectively). Dissolved Fe: PO_4 and Fe: Si ratios were greater
28 at P1, suggesting a lower relative biological demand for Fe at this site, possibly related to a
29 limitation Fe utilisation by the low concentrations of PO_4 or NO_x . Using the NO_x
30 concentrations measured at P1 (M. Woodward, pers. comm., 15-50 m) Fe: NO_x ratio was
31 negative (-122 nmol: μmol , $r^2 = 0.98$, $n=3$).

1 *Phytoplankton biology* – The vertical distribution of phytoplankton biomass (Fig. 10d), the
 2 cyanobacteria and nanoplankton distribution profiles (Fig. 10b, f) and the dominant
 3 phytoplankton groups inferred from biomarker pigments (Fig. 10c, g) were different at P1
 4 and P3. Despite similar total biomass inferred by total Chl *a*, the vertical distribution of
 5 phytoplankton is quite different (Fig. 10d). At P1, a deep Chl *a* maximum was observed
 6 (90-110 m), whereas high biomass was present in the top 30 m at P3, suggesting difference
 7 in nutrients and light availability amongst sites. Order of magnitude greater N₂ fixation
 8 rate were measured at P1 relative to P3 (Fig. 10h); as expected from the warmer
 9 temperature and the oligotrophic status of P1 (Law et al. 2011). However P1 has an
 10 unusual surface mixed layer profile with a maximum N₂ fixation rate at 50m. At P1,
 11 cyanobacteria of the type *Prochlorococcus* and *Synechococcus* were present with
 12 *Prochlorococcus* dominating at depth of maximal fluorescence, whereas *Synechococcus*
 13 dominated at P3 (Fig. 10b and DV Chl-*a* from Fig. 10c). Biomarker pigments
 14 demonstrated that haptophytes (Hex-Fucoxanthin) were also present at P1 and diatoms
 15 (Fucoxanthin) were only marginally abundant. At P3, cyanobacteria contribution to
 16 biomass was less important, and haptophytes were dominating the phytoplankton biomass
 17 as suggested by the MODIS calcite map of the study region during the time at sea and
 18 coccolithophorids abundances (Mark Baird and Cliff Law, pers. comm.). At P3, diatoms
 19 (Fucoxanthin) were more abundant than at P1 and dinoflagellates (Peridinin) were also
 20 present (Fig. 10g). This contrast in phytoplankton communities could result in differences
 21 in vertical patterns of nutrient and light limitation, primary productivity, generation of
 22 detritus and export.

23
 24 F_V/F_M is commonly used as an indicator for nutrient or light limitation (e.g., Behrenfeld et
 25 al. 2006). Here, we measured the in-situ F_V/F_M for the entire phytoplankton community
 26 including cyanobacteria (Red excitation) in contrast to the phytoplankton community
 27 excluding *Synechococcus* (Blue excitation; Fig. 10a and e). At P1, the F_V/F_M for both the
 28 cyanobacteria and the overall community was lower in surface and increased with depth to
 29 reach a maximum value at the depth of maximal fluorescence (95 m). This suggests that
 30 light was not limiting at that depth, but nutrient limitation or photo-inhibition processes
 31 occurred at shallower depths. Similar F_V/F_M profiles were measured at P3; however, the
 32 total phototrophic community showed a more constant and an overall greater F_V/F_M (Red)
 33 compared to P1. Sub-optimal F_V/F_M (< 0.6) at the maximum fluorescence P3 suggested
 34 that phytoplankton growth was limited. The greater difference in red and blue F_V/F_M at P1

1 indicated a greater contribution of *Synechococcus* to the total phototrophic signal. The
2 dominance of coccolithophorids at P3 provided a strong haptophyte signal that masked any
3 minor fluorescence signature from the cyanobacteria; this is why both red and blue F_V/F_M
4 values were similar.

5
6 As for the dFe concentrations, the particulate iron (pFe) concentrations were lower at the
7 depth of maximal fluorescence at P1 (Table 2). The ratios pFe: POC were similar in
8 surface water at P1 and P3, and lower at depth for P1. In contrast, the ratio pFe: Chl-*a*
9 increased at depth at P1, and was greatest at P3. The dFe concentrations normalised by
10 POC were smaller than the pFe concentrations normalised by POC for both sites, whereas
11 the dFe concentrations normalised by Chl-*a* were greater than pFe: Chl-*a* at site P1 and
12 lower at P3.

13
14 The phytoplankton iron uptake rate (Fig. 11a) is a measure of their ability to take up Fe
15 which can be related to Fe biological requirement and Fe limitation (Hassler et al. 2012b).
16 Here, Fe uptake rates were measured for three phytoplankton size classes to elucidate
17 whether Fe differentially controls phytoplankton within the natural community present at
18 the depth of maximal fluorescence. To account for the differences in phytoplankton
19 biomass between sites and size class (Tot Chl-*a*, Fig. 11b), Fe uptake rate was normalised
20 to Chl-*a* (Fig. 11c). At P1, the phytoplankton community composition was dominated by
21 pico- and nano-plankton. From the composition of biomarker pigments (Fig. 11d), the
22 picoplankton were mainly cyanobacteria of the type *Prochlorococcus*, whereas the
23 nanoplankton were mainly haptophytes. The Fe uptake rate per unit of biomass was lower
24 for the picoplankton than for the nanoplankton, suggesting different abilities to take up
25 iron or different iron requirement between prokaryotic and eukaryotic phytoplankton at
26 this site. At P3, iron uptake rates per unit of biomass were similar for pico- and nano-
27 plankton despite greater relative abundance of nano-plankton, suggesting that parameters
28 other than Fe controlled the abundance of phytoplankton at this site. At P3, the size classes
29 did not resolve the different phytoplankton functional groups well with haptophytes
30 dominating and diatoms present in the micro- and nano-plankton size class.
31 Dinoflagellates and cyanobacteria of the *Synechococcus* type were mainly present in the
32 micro- and pico-plankton, respectively.

1 Despite a lower F_V/F_M (Red) and biomass of the total phototrophic community in surface
2 water at the site P1 compared to P3, Chl-*a*-specific carbon fixation from the phytoplankton
3 community was greater at P1 (Fig. 12b). It is to be noted that the F_V/F_M (Blue) was greater
4 at P1 relative to P3. The small difference between the biological carbon fixation at low and
5 high light in surface waters, suggest that these communities were well adapted to their in-
6 situ light levels (382 and 56 $\mu\text{mol photons m}^{-2} \text{ s}^{-1}$ at P1 and P3, respectively). At the depth
7 of maximum fluorescence, the biological carbon fixation was greater at P1 than P3 (Fig.
8 12c-d). The largest difference in biological carbon fixation between these two sites was
9 observed at low light, suggesting that the phytoplankton community of P1 is well adapted
10 to low light prevailing at depth. At P1, an in-situ light intensity of 25 $\mu\text{mol photons m}^{-2} \text{ s}^{-1}$
11 was measured at the depth of 74 m (Suppl. Fig. 3). The large difference in carbon fixation
12 between the low and high light treatment at the site P3 demonstrated that the in-situ
13 biological carbon fixation was not optimal, and limited by the low light intensity found at
14 the depth of maximal fluorescence (13 $\mu\text{mol photons m}^{-2} \text{ s}^{-1}$ at 25 m).

16 **Discussion:**

17 During the time of the cruise, the oceanography in the region was representative for austral
18 summer (Ridgway and Dunn 2003). A comparison of the PINTS sections with
19 climatological fields did not show any large-scale offsets in temperature, salinity or any
20 other properties; rather the differences were at relatively small spatial scales and likely to
21 be associated with the presence of meso-scale eddies. Despite CTD deployments being
22 too widely spaced to accurately represent the detailed oceanography in the study region,
23 distinctive patterns in the distribution of micro- and macro-nutrients, and total Chl-*a* were
24 observed. Marked differences in most of the parameters studied were observed between
25 the North Tasman Sea and the SAZ southeast of Australia, as also predicted by the model
26 from Moore and co-workers (Moore and Abbott 2000; Moore et al. 2002) and from recent
27 studies (Bowie et al. 2009; Law et al. 2011; Ellwood et al. 2013). Despite observed
28 concentrations for dFe and Chl-*a* being greater than predicted by the mixed-layer
29 ecosystem model (Moore et al. 2002), comparative trends between P1 and P3 were
30 accurate for dFe, Chl-*a*, NO_x , atmospheric N_2 and C biological fixation. As predicted by
31 their model, the greater primary productivity at P1 is not associated with greater carbon
32 export as the phytoplankton community was dominated by small cyanobacteria. The model
33 predicted greater CaCO_3 export at P3, which was corroborated by the dominance of

1 haptophytes, including *Emiliania huxleyii* and the MODIS calcite composite map (this
2 study; C. Law and M. Baird, pers. comm.).

3
4 The in-situ F_V/F_M values at P1 and P3 also suggested a growth limitation or a photo-
5 inhibition which significantly differs for the cyanobacteria and the whole phytoplankton
6 community at P1. Despite a sub-optimal F_V/F_M being a good indicator for Fe limitation in
7 high-nutrient-low-chlorophyll regions, previous studies have also reported high F_V/F_M
8 under N and Fe co-limitation (Behrenfeld and Miligan 2013), preventing the use F_V/F_M as
9 a sole indicator of nutrient limitation in our study region.

10
11 A ‘kink’ in the disappearance ratio trend for Cd was observed for the surface waters of P1
12 and P3, suggesting a decrease in phytoplankton growth rate (Sunda and Huntsman 2000;
13 Cullen et al. 2003). Previous studies in the south Tasman Sea have observed such a trend
14 for $PO_4 < 0.6 \mu M$, with a similar disappearance ratio at greater PO_4 concentrations (323
15 and 460 for austral summer and winter, respectively, at 15–100m; Ellwood 2008a; Butler
16 et al. 2012; Hassler et al. 2012a). The parameters usually discussed as a control of
17 phytoplankton growth rate, with an effect on disappearance ratios are: (i) iron limitation
18 (e.g., Martin et al. 1990; deBaar et al. 1997; Cullen et al. 2003), (ii) combined effect of
19 micronutrients (Sunda and Huntsman 2000; Twining et al. 2004; Cullen and Sherrell 2005;
20 Gault-Ringold et al. 2012), and (iii) community structure and differing nutrient
21 requirement (e.g., deBaar et al. 1997; Arrigo et al. 1999; Sunda and Huntsman 2000; Ho et
22 al. 2003; Quigg et al. 2003; Twining et al. 2004; Finkel et al. 2007). In surface waters, a
23 greater disappearance ratio is often associated with a variation in the relative biological
24 uptake but could also be related to different recycling and remineralisation rates.
25 Unfortunately, neither the bacterial loop nor the grazing rates were measured in this study.
26 Owing to previously high grazing rates and remineralisation processes reported in the
27 region (Pearce et al. 2011; Jacquet et al. 2011), further studies should address their impact
28 on disappearance ratio and the biogeochemical cycle of nutrients. Biological requirement
29 can also be modulated by the bioavailability of other nutrients, light, temperature and
30 change of pCO_2 that occur under bloom situations, with impact on disappearance ratios
31 (see Finkel et al. 2010 for a review).

32
33 Comparison of the carbon fixation measured under $250 \mu mol \text{ photons } m^{-2} s^{-1}$ with the
34 optimal carbon fixation predicted from in-situ temperature (Behrenfeld and Falkowski

1997; Fig. 12d) suggests that maximal biological carbon fixation was only measured in surface water at P1. It is noted that this maximal carbon fixation at that site was remarkable given the low dissolved NO_x and PO_4 concentrations; the ability to fix atmospheric N_2 is likely essential for an efficient carbon fixation (Ellwood et al. 2013). Sub-optimal carbon fixation has important impacts for carbon export and climate, and could be due to saturating light greater than the $250 \mu\text{mol photons m}^{-2} \text{ s}^{-1}$ used in this study (e.g., Westwood et al. 2011) or due to light or nutrient limitation.

Within the study region, nitrogen and iron were the two nutrients that most likely limited phytoplankton growth. Overall the good correlation between C/N to Chl-*a* observed here, suggests that N is a key nutrient for most of the study region, as recently demonstrated in the North Tasman Sea (Ellwood et al. 2013). Despite this, C/N ratios for eukaryotic phytoplankton can significantly differ from the Redfield ratio (6–11; Quigg et al. 2003), with C/N ratios reported for *Trichodesmium* and marine *Synechococcus* and *Prochlorococcus* being close to the Redfield ratio (5–6.2; Bertilsson et al. 2003; Kutska et al. 2003; Nuester et al. 2012). At P1, several parameters indicate that nitrogen limitation was an important control of the phytoplankton community in the surface water of P1: these are low NO_x concentrations ($< 0.011 \mu\text{M}$), a C/N ratio greater than the Redfield ratio (6.6; Redfield et al. 1963), and the presence of diazotrophs fixing atmospheric nitrogen. As NO_x concentrations and F_v/F_m increased with depth, coincident with a decreasing C/N ratio, this suggested that nitrogen limitation was somewhat relieved at the depth of the maximal fluorescence, possibly allowing for the transition between a *Synechococcus*-dominated cyanobacterial community to a *Prochlorococcus*-dominated community at depth. Such trend in cyanobacteria distribution in relation to NO_x concentration has been previously observed in the North Tasman Sea (Ellwood et al. 2013).

The Australian continent is a significant source of iron-rich dust delivered to the North Tasman Sea, the SAZ and the Southern Ocean (Hesse 1994; Hesse 1997; Gabric et al. 2002; Hesse and McTainsh 2003; Boyd et al. 2004; Bowie et al. 2009; Mongin et al. 2011, Cropp et al. 2013). Periodic inputs of dust laden with iron is thought to stimulate phytoplankton growth (Bowie et al. 2009; Moore et al. 2009), resulting in a shift of the dominant phytoplankton species composition and an increase in carbon fixation (Lenes 2001), thus providing a vital link in the planet's climate control system and being part of a key climate feedback loop linking the lithosphere, atmosphere and ocean (Jickells et al.

2005). In the study region, Moore et al. (2002) model runs, in the absence of, and with, iron-saturating dust deposition scenarios, showed that dust input can increase primary productivity, phytoplankton biomass and induce a shift in the composition of the phytoplankton community. Conflicting results were obtained from recent simulations of wet and dry dust deposition in the North Tasman Sea, with enhanced N_2 fixation by unicellular cyanobacteria (Law et al. 2011) and no effect on phytoplankton growth (Ellwood et al. 2013). During the PINTS study, experiments simulating dry Australian dust deposition, showed that such an event can induce phytoplankton growth and a clear shift in the composition of the phytoplankton community both at P1 and P3 (Hassler et al. unpublished data), suggesting that iron released from dust could affect the structure of the phytoplankton community in our study. This highlights, as observed for artificial large scale Fe fertilisation, the importance of the oceanography and biological conditions at the time of dust deposition (Boyd et al. 2007; Cropp et al. 2013).

Here, differences in dFe: POC, pFe: POC, dFe: Chl *a*, and pFe: Chl *a* between sites and depths, suggested differences in the iron biological requirement or limitation for the in-situ phytoplankton community. The fact that ratios with dFe are smaller than ratios with pFe at P3, suggests that the pool of dFe is not enough to sustain further phytoplankton growth at that site. However, the link of these parameters to iron limitation have to be interpreted with care as they can also indicate (1) a phytoplankton species-dependent or light-dependent cellular Chl-*a* content (see Hassler and Schoemann 2009a; Morrissey and Bowler 2012 for discussion), (2) difference in the lithogenic pFe contribution or (3) the detritus contribution to POC and PON. In addition, luxurious iron uptake, in case of episodic iron enrichment (e.g. subsequent to dust deposition) can increase pFe to POC ratio. Indeed surface iron enrichment was observed at P1. Such dFe profile in this region was previously discussed as the result of dust deposition event (Law et al. 2011).

The N_2 fixation process results in a greater iron biological requirement by diazotrophs than most other phytoplankton (Kutska et al. 2003). A 15-fold increase in Fe uptake was calculated due to an increase in N_2 fixation rate by unicellular cyanobacteria in the North Tasman Sea (Law et al. 2011). The N_2 fixation rate observed in the top 50 m at P1 were similar to the rates previously observed in the study region (Law et al. 2011). In such a case, despite high Fe^* values (up to 0.29 at 15 m), potential Fe co-limitation could occur in the surface water of P1. Diazotrophs and cyanobacteria are able to efficiently modulate

1 their iron biological requirement in response to their environment and during the diel cycle
2 through the control of biological processes (e.g., photosynthesis, N₂ fixation) and gene
3 expression (Saito et al. 2011; Morrissey and Bowler 2012). At the depth of maximal
4 fluorescence, dFe and pFe were extremely low, at a level usually limiting the growth of
5 most eukaryotic phytoplankton (e.g., Sunda and Huntsman 2000, Sarthou et al. 2005;
6 Sedwick et al. 2007) and Fe* was negative indicating probable Fe limitation (Bowie et al.
7 2009). In addition, dFe: POC were smaller than pFe: POC. Finally, iron bioaccumulation
8 data, suggested a low iron requirement for cyanobacteria relative to eukaryotic
9 phytoplankton at that depth. Indeed iron internalisation constants (data not shown)
10 demonstrated that cyanobacteria (picoplankton) were not iron limited in this case. Primary
11 productivity data demonstrated that light was not limiting the in-situ plankton community
12 at P1, even at the depth of maximal fluorescence. Therefore, our data have demonstrated a
13 vertical difference in nutrient limitation, where in surface waters Fe could limit
14 diazotrophs and nitrate was mostly limiting the growth of other phytoplankton; but at
15 depth iron was the main limiting nutrient for the in-situ eukaryotic phytoplankton. Law et
16 al. (2011) demonstrated that the vertical diffusive supply of nitrate was able to maintain
17 the deep chlorophyll maximum in this region, and that nitrate and possibly low
18 temperature were limiting diazotrophy at these depths. Si concentrations were lower than
19 the average diatom requirement for growth ($3.9 \pm 5.0 \mu\text{M}$; Sarthou et al. 2005), so could
20 also limit diatom abundances and growth at depth.

21
22 In the SAZ, the situation is quite different than from the North Tasman Sea. SeaWiFS
23 satellite imagery, model and field observations reveal a persistent phytoplankton bloom
24 present in the surface waters from October to April (Moore et al. 2002; Bowie et al. 2011;
25 de Salas et al. 2011). The earlier SAZ-Sense study performed during the 2007 austral
26 summer demonstrated that this bloom was likely fuelled by episodic, iron-laden dust
27 deposition which may have been transported south in the EAC, and lateral transport from
28 the shelf sediments in EAC eddies (Bowie et al. 2009; Mongin et al. 2011) and the
29 composition of its phytoplankton community was likely influenced by iron availability
30 (Hassler et al. 2012a). However, other in-situ indicators of iron limitation gathered during
31 the SAZ-Sense study such as Fe* (Lannuzel et al. 2011), F_v/F_m (Petrou et al. 2011) and
32 biological uptake of Fe (Schoemann et al., unpublished) suggest that iron is not the main
33 limiting nutrient. Here, as was observed in winter (Ellwood et al. 2008), the strong
34 correlation between Fe and macronutrients concentration demonstrated its role as a

1 nutrient. The absence of Fe:PO₄ correlation during the SAZ-Sense study (Hassler et al.
2 2012a), could have been due to dust enrichment in surface waters.

3
4 The model developed by Moore et al. (2002) predicts that phytoplankton are mainly
5 limited by Fe with Si co-limitation for diatoms. Despite being in accordance with the
6 potential Fe and Si limitations in the SAZ, our data demonstrated a critical limiting role for
7 light. Primary productivity and in-situ light measurements demonstrated that the primary
8 productivity was strongly limited by the light prevailing at depth of the maximum
9 fluorescence. The plankton biomass in the SAZ region is so high ($> 1 \mu\text{g L}^{-1}$ Chl-*a* in some
10 places; de Salas et al. 2011; this study) that significant shading occurs (Westwood et al.
11 2011; this study). In the top 30 m, where most of the biomass was observed, most nutrients
12 were strongly depleted due to biological consumption. In this case, Si (0.7 μM), Fe (0.2
13 nM), NO_x (1.6 – 3.0 μM) were low enough to potentially limit or co-limit further growth
14 of phytoplankton (e.g., Sarthou et al. 2005; Veldhuis et al. 2005; Schoemann et al. 2005;
15 Sedwick et al. 2007). Given that different phytoplankton have different biological
16 requirements for growth (e.g., Ho et al. 2003), a range of co-limiting scenarios might
17 simultaneously occur in the region (Hassler et al. 2012a).

18
19 Here F_V/F_M values were below 0.6 at the depth of maximum fluorescence, and pFe: POC
20 were greater than dFe: POC, suggested that dFe is potentially not able to sustain
21 phytoplankton growth. As iron biological requirement of phytoplankton is greater under
22 low light (e.g., Raven et al. 1999; Timmermans et al. 2001), light limitation increase the
23 likelihood of iron co-limitation. In our study, as opposed to the austral summer of 2007
24 (Bowie et al. 2009; Mongin et al. 2011), no dFe surface enrichment from dust deposition
25 was observed, thus limiting iron inputs needed to sustain phytoplankton growth, as attested
26 by a lower Fe* in surface waters (Bowie et al. 2009). Iron bioaccumulation data suggest
27 that iron limitation will affect the micro, nano and pico-plankton communities differently
28 (this study; Hassler et al. 2012a). For these reasons, it is hard to unequivocally identify the
29 control that nutrient(s) exert on phytoplankton growth in this SAZ region. This study
30 suggests that, in the SAZ southeast of Australia, nutrient limitation is likely to be complex,
31 heterogeneous between phytoplankton groups and size classes, and variable in time due to
32 the evolution of the bloom, as well as nutrient inputs from dust and the mesoscale
33 processes at play in the region (Ellwood et al. 2008b; Bowie et al. 2009; Bowie et al.
34 2011). In addition, the potential for iron to limit phytoplankton growth is a function of its

complex chemistry and biological requirements, as well as the availability of other nutrients (e.g., N, Si, Cu; see Hassler et al. 2012b; Shaked and Lis 2012; Sunda 2012 for recent reviews). In the SAZ region, light limitation of phytoplankton growth needs to be taken into account during the austral summer.

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Figure captions:

Figure 1: The Tasman Sea region showing the bottom topography (depth, [m]). The PINTS stations along the cruise track are shown by the coloured symbols. Transect stations are shown in yellow, process stations in red (P1=Stn 5, P2=Stn 9, P3=Stn 12) and reoccupation of the process station 3 from the SAZ-Sense expedition (*Aurora Australis*, Jan-Feb 2007) in green (SAZ=Stn14). Thicker solid lines indicate the East Australian Current (EAC), Tasman Front (TF), and EAC Extension. The dashed line represents the path of the Subtropical Front (STF).

Figure 2: (a) Surface circulation of the Tasman Sea region as shown by the surface steric height field ($h_{0/200}$, m) obtained from the CARS climatology (Ridgway and Dunn, 2003). (b) The surface height variability (m) obtained from 18 years of satellite altimetry observations. (c) The full-depth transport shown by the depth-averaged steric height from CARS.

Figure 3: Temperature anomaly data (with respect to the annual mean from CARS) obtained from synTS along the PINTS cruise track in the upper 1000 m (a). Temperature anomaly from CTD stations (b). Note the increased resolution in the vertical axis from 0-125 m. The location of the CTD stations is indicated by the dashed lines and the end of each transect is shown by the full lines.

Figure 4: Temperature (a) and salinity (b) at 60-m inferred from SynTS. Temperature (c) and salinity (d) at 250 m inferred from SynTS. The contour intervals are 0.25°C (a, c); 0.025 (b) and 0.02 (d) g kg⁻¹. The upper panels show the pattern for the first portion of the voyage (January 2010) and the lower panels represent the latter half of the voyage (February 2010).

Figure 5: CTD data collected, (a) temperature in °C, (b) salinity in g kg⁻¹ and (c) oxygen in µM down to 1000m along the PINTS voyage track. The voyage track is shown on the right panel overlaid on bathymetry.

Figure 6: Phosphate (PO_4 , a), nitrate + nitrite (NO_x , b) and silicic acid (Si, c) distribution along the PINTS voyage track down to 1000 m depth. The voyage track is shown on the right panel overlaid on bathymetry.

Figure 7: Total Chlorophyll-*a* (Tot Chl-*a*, a) and particulate organic carbon to particulate nitrogen ratio (C/N, b) along the PINTS voyage track. Tot Chl-*a* distribution is shown down to 125 m, whereas C/N are shown for surface (25m) and the depth of maximum fluorescence. The depth of maximum fluorescence coincides with the depth of maximum Tot Chl-*a*. The voyage track is shown in the right panel, overlaid on bathymetry.

Figure 8: Profile of dissolved trace elements and nutrients at sites P1 (Stn5, full circles) and P3 (Stn 12, empty circles). Dissolved iron (dFe, a), cadmium (dCd, b), silicic acid (Si, c), nitrate + nitrite (NO_x , d), and phosphate (PO_4 , e).

Figure 9: Disappearance ratios for Cd (panel a), nitrate + nitrite (NO_x , b), and Fe (panels c to h) at sites P1 (Stn 5, full circles) and P3 (Stn 12, empty circles). Disappearance ratio are shown against phosphate (PO_4 , panels a-d), silicic acid (Si, panels e-f), and nitrate + nitrite (NO_x , panels g-h). Disappearance ratios are calculated by linear regression (slope with 95 % confidence interval and coefficient of correlation) using data down to 4000 m (panels a-c, e, g) and down to 125m (panels d, f, h). NA = not applicable.

Figure 10: Biological parameters from sites P1 and P3. Maximum quantum yields from the whole phytoplankton community (F_v/F_m Blue, full circles) and from cyanobacteria (F_v/F_m Red, empty circles) are shown in panels a and e ($n=3$, error bars represents the standard deviation); cell abundance are shown in panels b and f; total chlorophyll-*a* (Tot Chl-*a*) is shown in panel d; biomarker pigments are shown in panels c and g; and N_2 fixation arte are shown in panel h ($n=3$, error bars represents the standard deviation).

Figure 11: Iron bioavailability of size fractionated phytoplankton community from the depth of maximum fluorescence at stations P1 and P3. Average iron uptake rates are presented with half interval ($n= 2$, panel a and c). Size fractionated total Chlorophyll-*a* (Tot Chl-*a*, b) and biomarker pigments relative concentration (c) is presented for microplankton ($> 10 \mu\text{m}$), nanoplankton (2-10 μm) and picoplankton (0.7-2 μm).

1 Figure 12: Biological carbon fixation by phytoplankton at sites P1 and P3 for two depths
2 (surface, 15m and maximum of fluorescence (P1= 125m, P3 = 25m). Carbon fixation is
3 shown at two light intensities (30 ± 3 and $271 \pm 35 \mu\text{mol photons m}^{-2} \text{s}^{-1}$, $n=6$, represented
4 in dark and light bars respectively). Optimal carbon fixation was calculated from in-situ
5 temperature (open stars) using the equation from Behrenfeld and Falkowski (1997). Error
6 bars represent half interval ($n=2$).
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Table 1: Stations coordinates and dates for the PINTS expedition. Activities are physical oceanography (O), nutrients (N), Biology (B), Trace metals (TM), Process study (P). Full water column (0-4500m) sampling is indicated by an asterisk, otherwise 0-1000m were sampled when bottom depth was > 1000 m. A single time of sampling was not appropriate at process stations (P1, P2, P3 and reoccupation from the SAZ-Sense voyage (2007)) as 3 to 17 CTD casts and 2 to 7 trace metal rosette casts were performed.

Stns #	Activity	Latitude (°E)	Longitude (°S)	Date (UTC, 2010)	Time (UTC)
1	O, N, B	153 5	33 2	23-Jan	0:49
2	O, N, B, TM	155 5	32 5	24-Jan	17:12
3	O, N, B, TM	158 4	32 0	26-Jan	1:02
4	O, N, B, TM	161 5	31 0	26-Jan	19:49
5 - P1	O, N, B, TM, P*	165 0	30 0	27 to 31-Jan	NA
6	O, N, B, TM	164 3	31 4	31-Jan	22:55
7	O, N, B	163 3	35 3	2-Feb	2:26
8	O, N, B, TM	162 4	38 3	3-Feb	1:13
9 - P2	O, N, B, TM, P*	162 1	39 6	3 to 4-Feb	NA
10	O, N, B, TM	161 1	42.3	6-Feb	18:19
11	O, N, B	160 5	43 5	7-Feb	7:41
12 - P3	O, N, B, TM, P*	159 5	46 2	7 to 9-Feb	NA
13	O, N, B, TM	156 5	46 0	10-Feb	22:54
14 - SAZ	O, N, B, TM*	153 2	45 6	11 to 12-Feb	NA
15	O, N, B, TM	150 5	45 5	13-Feb	16:05
16	O, N, B, TM	149 4	44 6	14-Feb	1:57
17	O, N, B, TM	148 4	44 1	14-Feb	13:12
18	O, N, B, TM	148 0	43 4	14-Feb	23:10

1 Table 2: Particulate data collected by Mc Lane pumps on a 1 μm quartz filter. Particulate iron (pFe), organic carbon (POC) and particulate
 2 nitrogen (PON) from surface water and the depth of maximal fluorescence (F max) at P1 and P3. For comparison, similar data considering the
 3 dissolved iron (dFe, $< 0.2 \mu\text{m}$) and total chlorophyll-*a* (Chl-*a*) are shown.

4

	pFe (nM)	C/N	pFe/POC (nmol: μmol)	pFe/Chl- <i>a</i> (nmol: μg)	Chl- <i>a</i> /POC (μg : μmol)	dFe/POC (nmol: μmol)	dFe/Chl- <i>a</i> (nmol: μg)
P1							
Surface (30 m)	0.23	8.95	0.12	0.11	0.06	0.12	1.90
F max (95m)	0.12	6.52	0.08	0.39	0.24	0.03	0.61
P3							
F max (30 m)	0.77	6.61	0.15	1.11	0.14	0.04	0.30

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Primary productivity induced by iron and nitrogen in the Tasman Sea – An overview of the PINTS expedition.

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Supplementary Material:

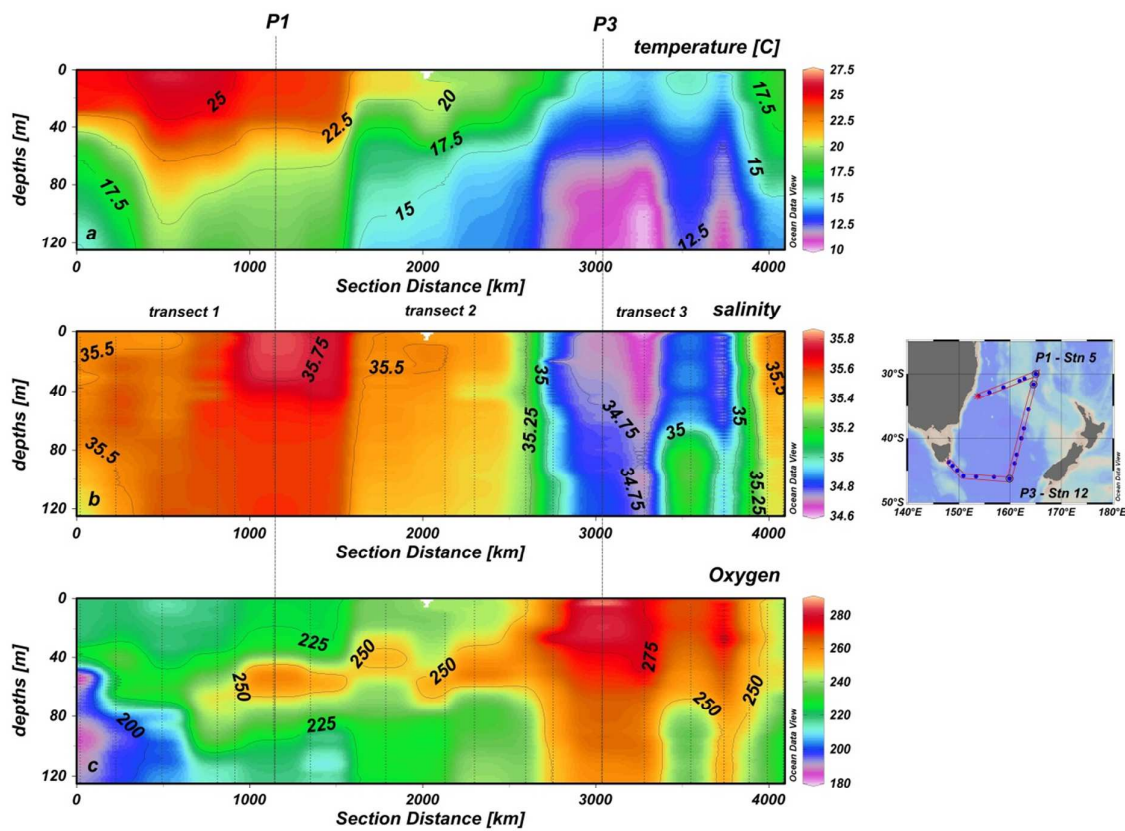


Figure 1: Data collected from the CTD deployment along the PINTS cruise track. Temperature (a), salinity (b) and oxygen (c) are shown down to 125m. This figure focuses on the area where phytoplankton have been measured.

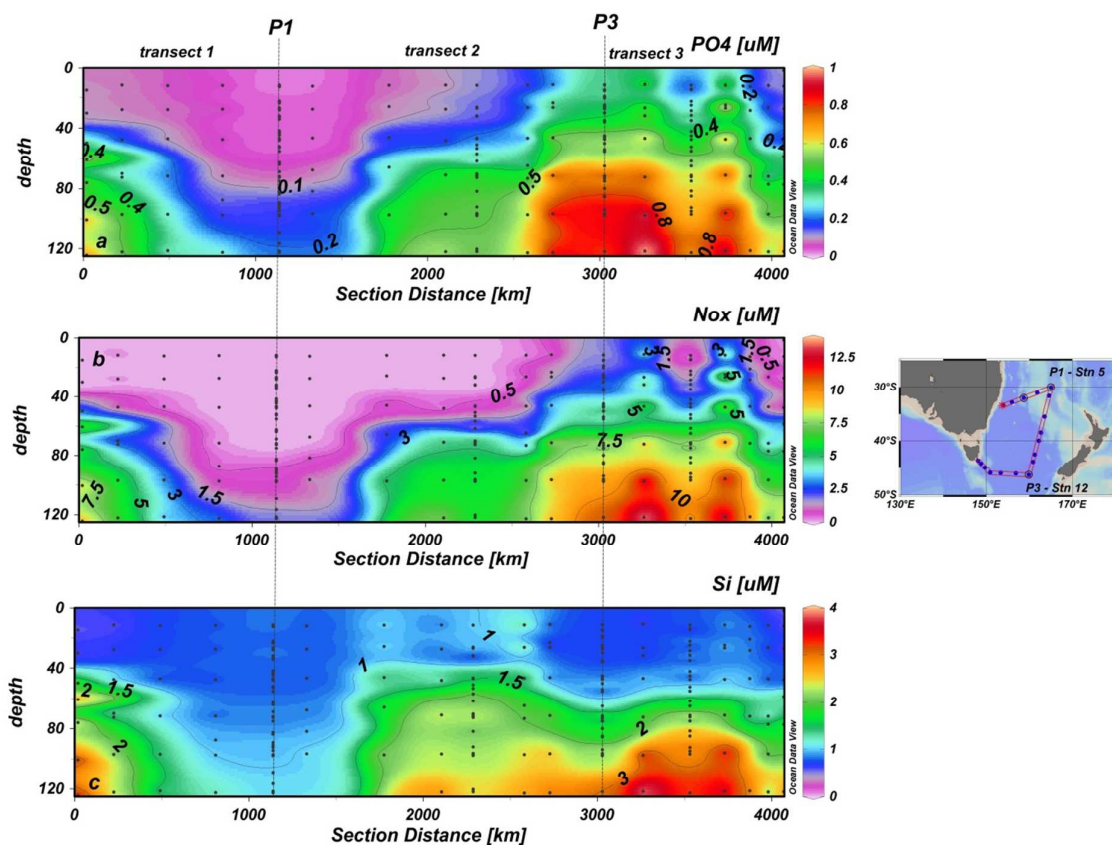


Figure 2: Data collected from the CTD deployment along the PINTS cruise track. Phosphate (PO_4 , a), nitrate + nitrite (NO_x , b) and silicic acid (Si , c) are shown down to 125m. This figure focuses on the area where phytoplankton have been measured.

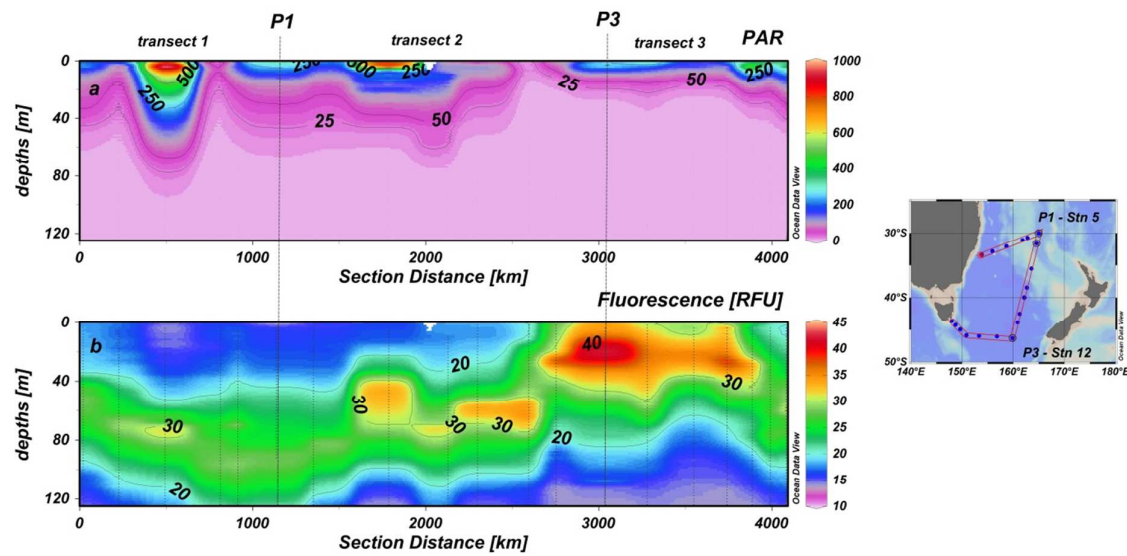


Figure 3: Data collected from the CTD deployment along the PINTS cruise track. Photosynthetic active radiation (PAR, a), and fluorescence (b) are shown down to 125m. This figure focuses on the area where phytoplankton have been measured.

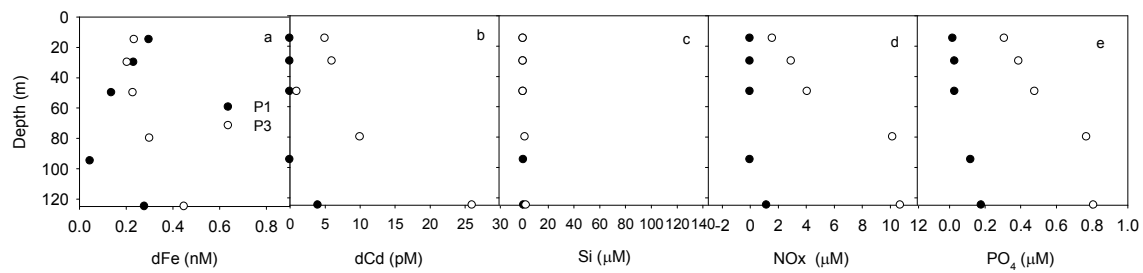
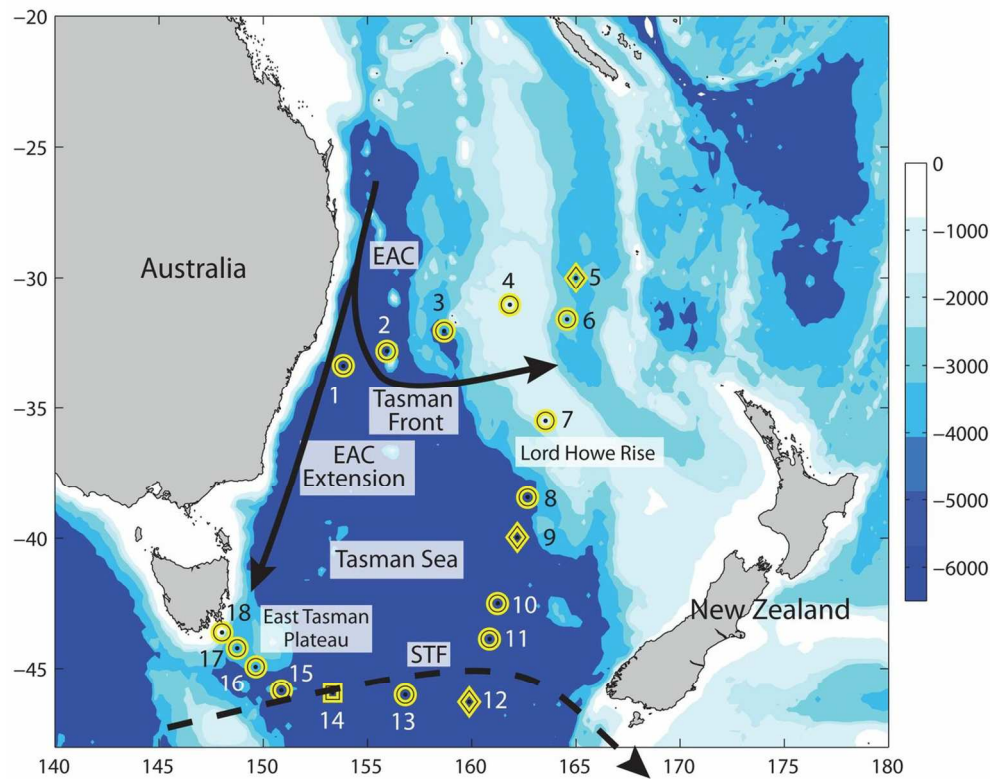
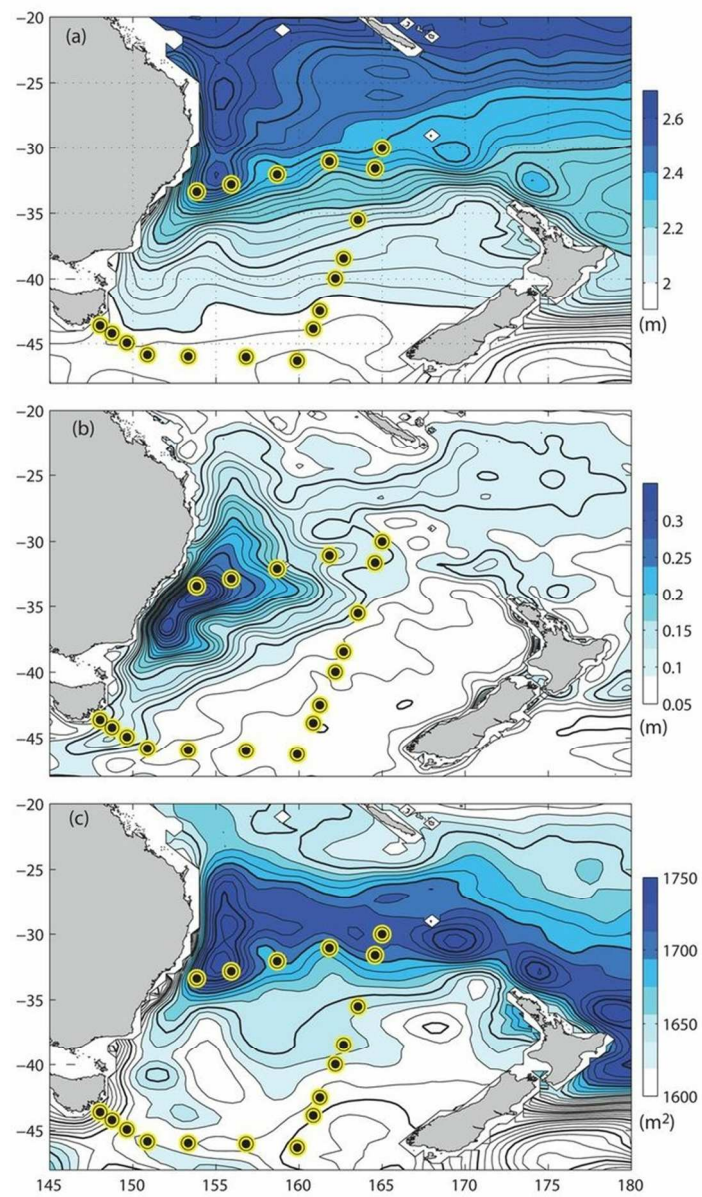


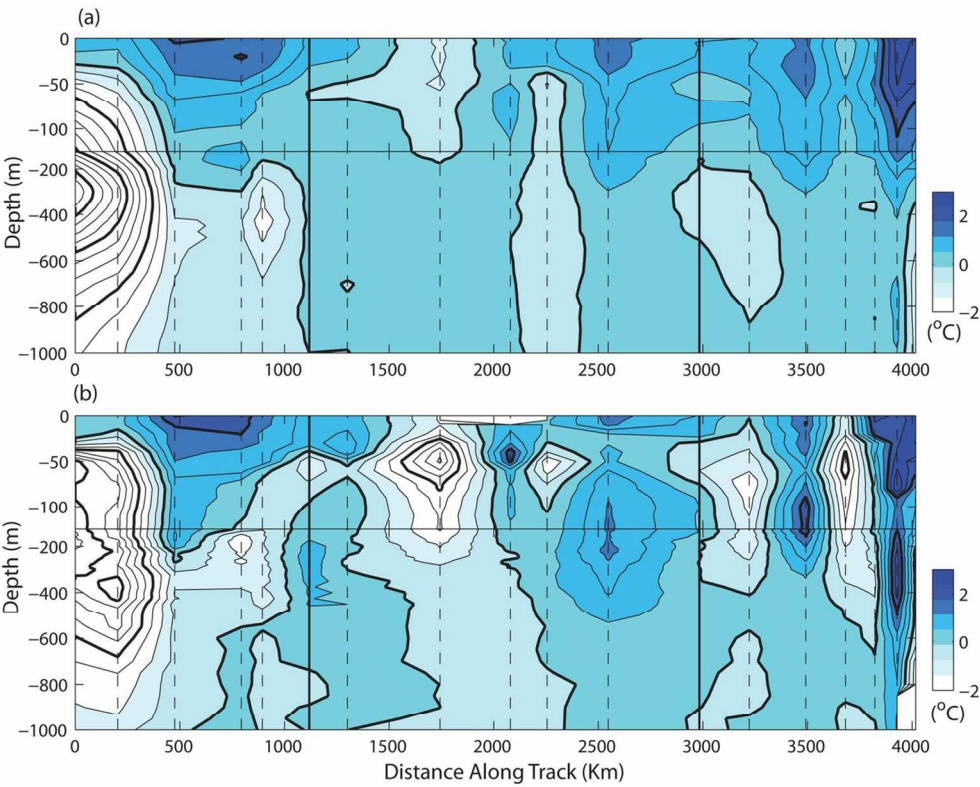
Figure 4: Profile of dissolved trace elements and nutrients in surface waters at sites P1 (full circles) and P3 (empty circles). Dissolved iron (dFe, a), cadmium (dCd, b), silicic acid (Si, c), nitrate + nitrite (NO_x, d), and phosphate (PO₄, e).



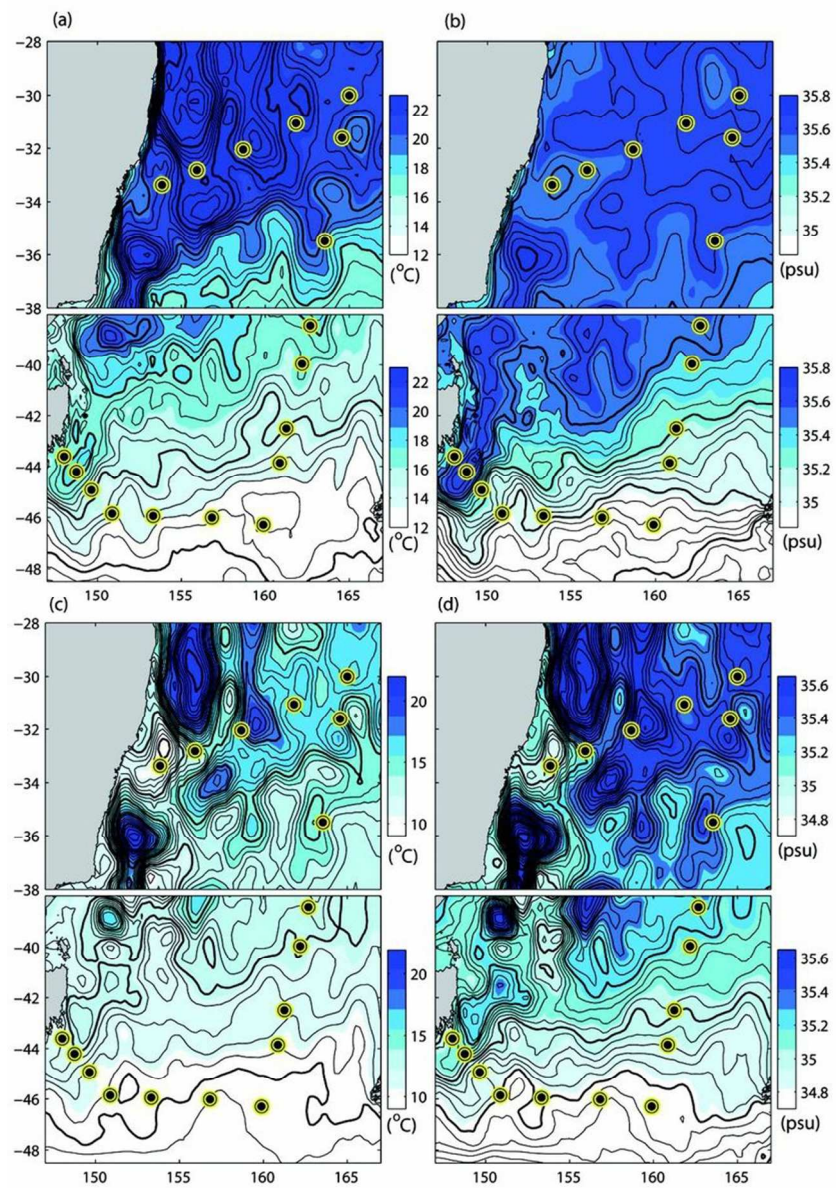
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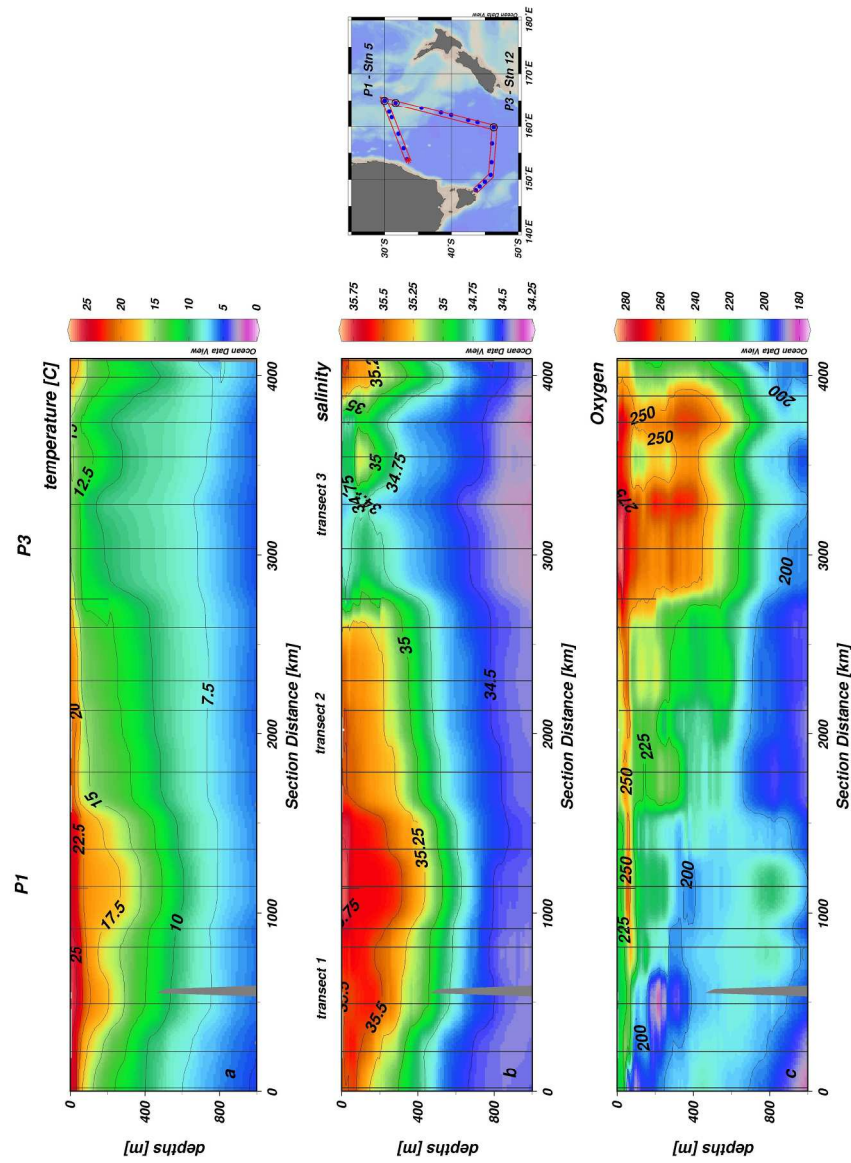
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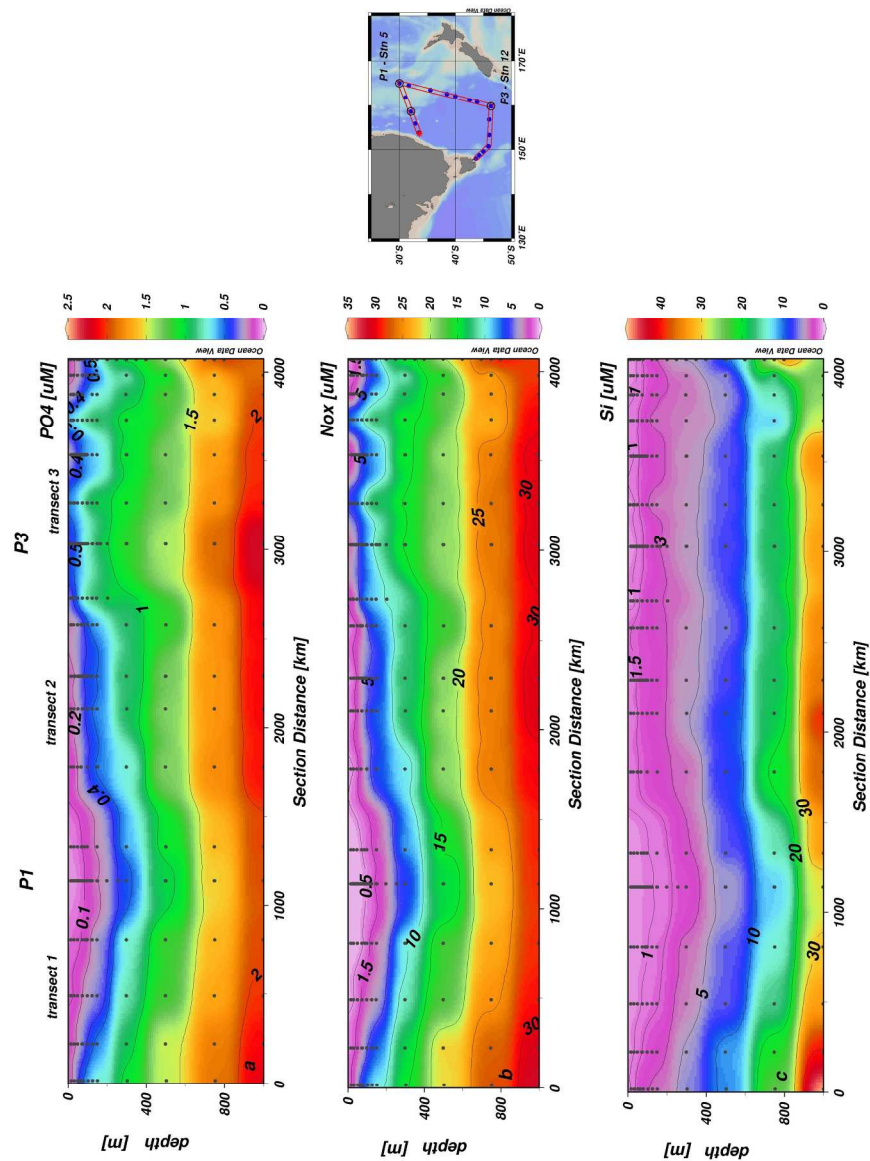
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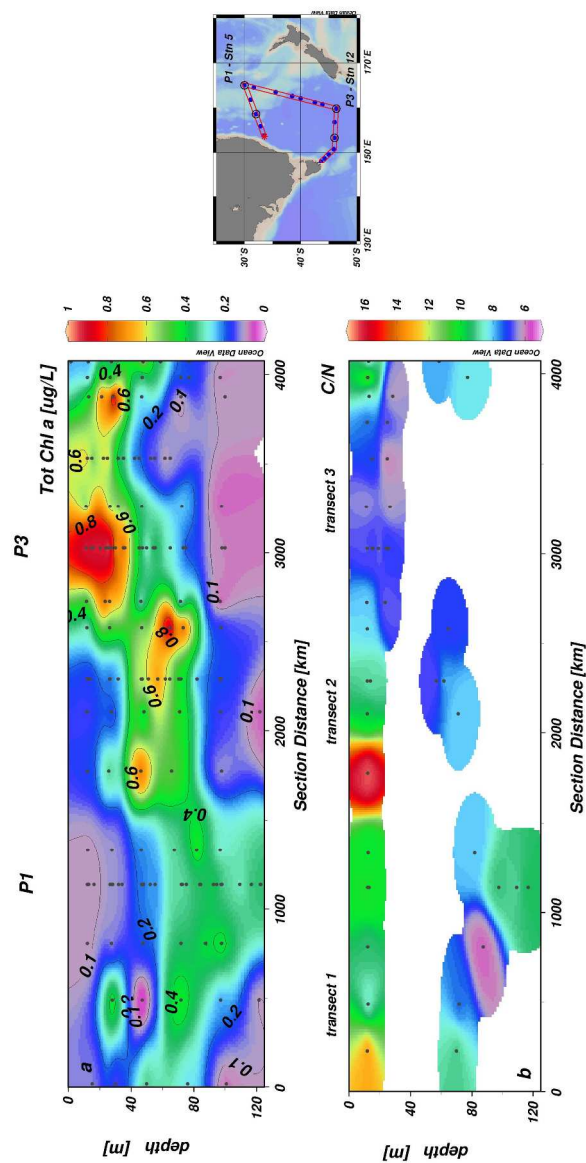
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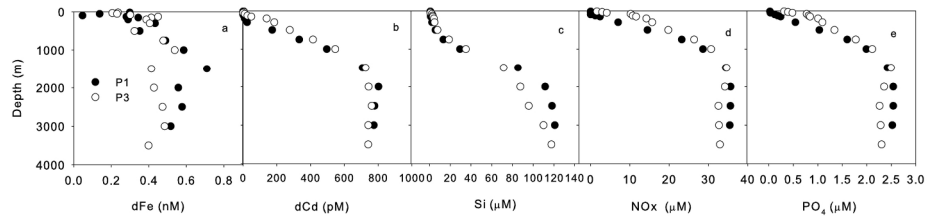
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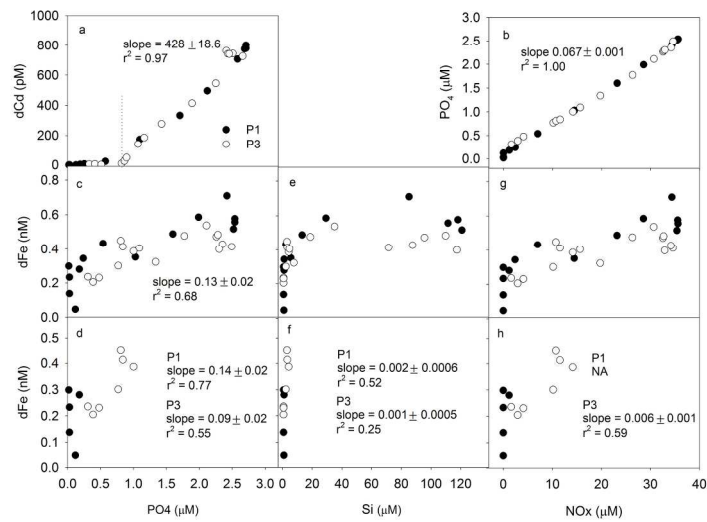
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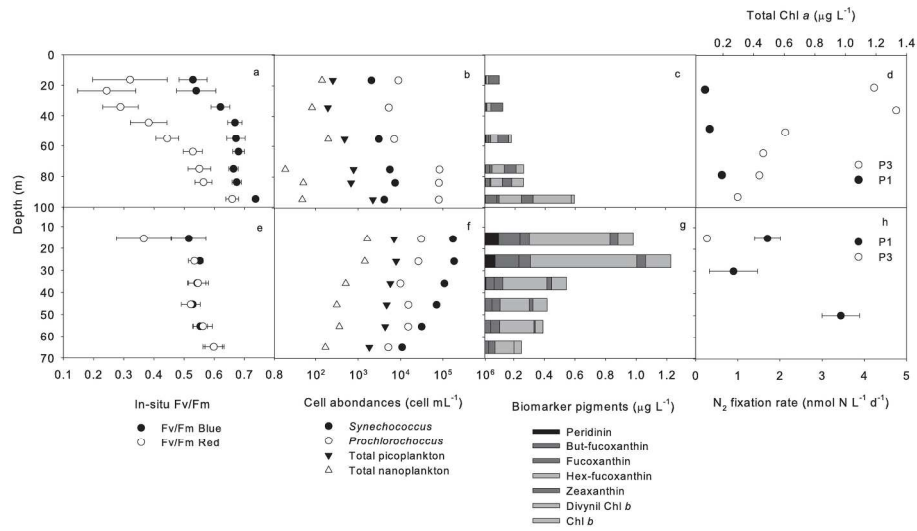
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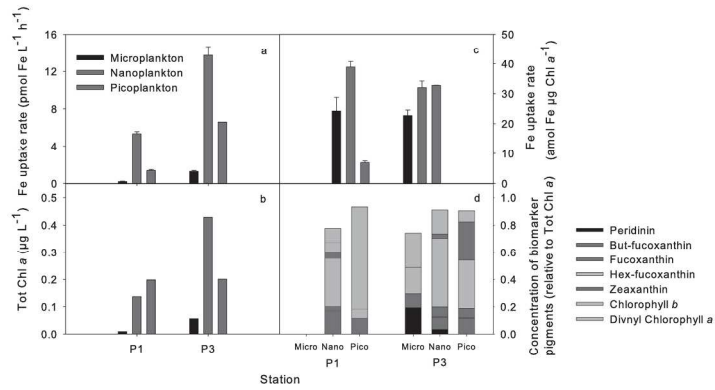
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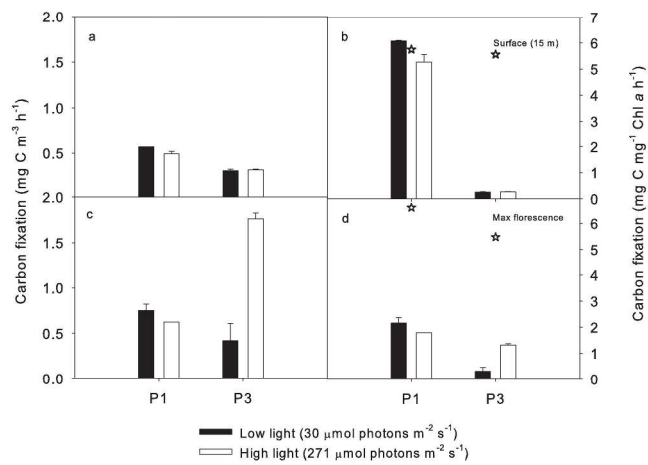
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209x147mm (300 x 300 DPI)



209x148mm (300 x 300 DPI)



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