



Title	Dynamics of particulate and dissolved organic and inorganic phosphorus during the peak and declining phase of an iron-induced phytoplankton bloom in the eastern subarctic Pacific
Author(s)	Yoshimura, Takeshi; Nishioka, Jun; Ogawa, Hiroshi; Tsuda, Atsushi
Citation	Journal of marine systems, 177, 1-7 https://doi.org/10.1016/j.jmarsys.2017.09.004
Issue Date	2018-01
Doc URL	http://hdl.handle.net/2115/76473
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Type	article (author version)
File Information	Yoshimura.pdf



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1 Dynamics of particulate and dissolved organic and inorganic phosphorus during the peak and
2 declining phase of an iron-induced phytoplankton bloom in the eastern subarctic Pacific

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4 Takeshi Yoshimura^{1,*}, Jun Nishioka², Hiroshi Ogawa³, Atsushi Tsuda³

5

6 ¹Central Research Institute of Electric Power Industry, 1646 Abiko, Abiko, Chiba 270-1194,
7 Japan

8 ²Institute of Low Temperature Science, Hokkaido University, Sapporo, Hokkaido 060-0819,
9 Japan

10 ³Atmosphere and Ocean Research Institute, The University of Tokyo, 5-1-5 Kashiwanoha,
11 Kashiwa, Chiba 277-8564, Japan

12

13 *Corresponding author (present address): T. Yoshimura (yoshimura-t@fish.hokudai.ac.jp)

14 Faculty of Fisheries Sciences, Hokkaido University

15 Graduate School of Environmental Science, Hokkaido University

16 Kita 10, Nishi 5, Kita-ku, Sapporo, Hokkaido 060-0810 JAPAN

17 Tel: +81-11-706-2324

18

19 April 26, 2017 (submitted)

20 August 4, 2017 (resubmitted)

21 Submitted as a Research Paper to Journal of Marine Systems

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24 ABSTRACT

25 Phosphorus (P) is an essential element for all organisms and thus the P cycle plays a
26 key role in determining the dynamics of lower trophic levels in marine ecosystems. P in
27 seawater occurs conceptually in particulate and dissolved organic and inorganic (POP, PIP,
28 DOP, and DIP, respectively) pools and clarification of the dynamics in these P pools is the
29 basis to assess the biogeochemical cycle of P. Despite its importance, behaviors of each P pool
30 with phytoplankton dynamics have not been fully examined. We measured the four
31 operationally defined P pools (POP_{op}, PIP_{op}, DOP_{op}, and SRP) during an iron-induced
32 phytoplankton bloom (as part of the subarctic ecosystem response to iron enrichment study
33 (SERIES)) in the eastern subarctic Pacific in summer 2002. During our observations of the
34 iron-enriched patch from day 15 to day 26 after the iron infusion, chlorophyll-*a* concentration
35 in the surface layer decreased from 6.3 to 1.2 $\mu\text{g L}^{-1}$, indicating the peak through decline
36 phase of the phytoplankton bloom. At the bloom peak, P was partitioned into POP_{op}, PIP_{op},
37 and DOP_{op} in proportions of 60, 27, and 13 %, respectively. While chlorophyll-*a* and POP_{op}
38 showed similar temporal variations during the declining phase, PIP_{op} showed a different peak
39 timing with a 2 day delay compared to POP_{op}, resulting in a rapid change in the relative
40 proportion of PIP_{op} to total particulate P (TPP = POP_{op} + PIP_{op}) at the peak (25 %) and during
41 the declining phase of the bloom (50 %). A part of POP_{op} was replaced by PIP_{op} just after
42 slowing down of phytoplankton growth. This process may have a significant role in the
43 subsequent regeneration of P. We conclude that measurement of TPP alone is insufficient to
44 show the interaction between P and phytoplankton dynamics and fractionation of TPP into
45 POP_{op} and PIP_{op} provides useful insights to clarify the biogeochemical cycle of P.

46

47 Keywords:

48 Phosphorus

49 Organic matter

50 Inorganic matter

51 Phytoplankton bloom

52 **1. Introduction**

53 Phosphorus (P) is a factor regulating phytoplankton productivity in the ocean. Better
54 understanding of the biogeochemical cycle of P is required to evaluate the role of P in
55 controlling marine ecosystems and thus to determine the link with other bioactive elements
56 such as carbon (C) and nitrogen (N). However, comprehensive studies on P are few compared
57 with C and N (Karl, 2014). P in seawater occurs in both particulate and dissolved pools, and
58 each of which contain organic and inorganic forms (POP, PIP, DOP, and DIP). Although the
59 most favorable form of P for phytoplankton growth is orthophosphate (PO_4), recent studies
60 have shown that other forms of P also play roles in the P cycle (Dyhrman, 2016; Karl, 2014).
61 Since these different P pools would have different regeneration pathways and thus have
62 different turnover times (Björkman and Karl, 2003), information on the size and dynamics of
63 each P pool is necessary to characterize the P cycle. However, fractionated analyses of total
64 particulate P (TPP) and total dissolved P (TDP) pools are scarce, as pointed out in Labry et al.
65 (2013) and Karl and Björkman (2015), respectively.

66 Past studies have shown that the fractionated measurement of P is a strategy for a
67 better understanding of the P cycle (Loh and Bauer, 2000). TPP pool can be operationally
68 differentiated into HCl-extractable P and non-extractable P pools, and TDP pool can be
69 differentiated into soluble reactive P (SRP) and soluble non-reactive P (SNP) pools; P pool
70 can be fractionated into four operationally defined fractions using filtration and chemical
71 fractionation methods, and these P pools have been assumed to represent PIP, POP, DIP, and
72 DOP, respectively (Loh and Bauer, 2000). The pool size and relative composition of each
73 fraction vary horizontally and vertically (Loh and Bauer, 2000; Piper et al., 2016; Yoshimura
74 et al., 2007). Uncoupled behavior of POP and PIP in sinking particles was also reported in
75 Benitez-Nelson et al. (2004). Recent P fractionation studies in estuarine (Labry et al., 2013;
76 Lin et al., 2013) and Arctic waters (Lin et al., 2012; Piper et al., 2016) provide insights into
77 the dynamics among the different P pools. Asahi et al. (2014) has shown that the POP to PIP
78 ratio is a useful index to determine the origin of suspended particulate matter in the coastal
79 environment. Although very detailed fractionation methods for TPP (Cade-Menun and Paytan,
80 2010; Miyata and Hattori, 1986; Ruttenberg, 1992) and TDP pool (Kolowitz et al., 2001;
81 Young and Ingall, 2010) have been developed, these studies have shown relatively simple
82 fractionation techniques to be useful for a better understanding of the P cycle, especially when
83 large amounts of samples are difficult to obtain notably in open ocean waters.

84 An issue is that chemically fractionated P pools are operationally defined pools and

85 do not necessarily correspond to the conceptual P pools (Fig. 1). As a typical DIP species, PO_4
86 has been measured by the molybdenum blue colorimetric method. However this method does
87 not measure polyphosphates but does measure some part of acid-labile DOP. Thus the P
88 fraction measured using the molybdenum blue technique does not represent DIP and is termed
89 SRP (Benitez-Nelson, 2000). Although the difference between TDP and SRP has been
90 assumed to be DOP, this fraction contains polyphosphates, and thus this P fraction should be
91 termed as SNP (Benitez-Nelson, 2000). Similarly is the case for TPP fractionations. For
92 suspended particulate matter samples on glass fiber filters, Labry et al. (2013) concluded that
93 the high temperature combustion and acid hydrolysis method for TPP of Solórzano and Sharp
94 (1980) and the acid hydrolysis method for PIP of Aspila et al. (1976) are the best methods to
95 estimate the POP and PIP pools. However, since polyphosphate is only partially hydrolyzed
96 (41–49 %) by the method of Aspila et al. (1976) (Labry et al., 2013), intracellular
97 polyphosphates are significantly underestimated in PIP_{op} pools and included in POP_{op} pools
98 (Fig. 1). Therefore we hereafter term the four operationally fractionated P pools as PIP_{op} ,
99 POP_{op} , SRP, and DOP_{op} . Care is required to discuss chemically fractionated P data with
100 conceptual P pools, but simple chemical fractionations still have an advantage to obtain useful
101 insights for a better understanding of the marine P cycle.

102 A phytoplankton bloom is a typical platform to study the dynamics of bioactive
103 elements with the rapid buildup of phytoplankton biomass. Although C and N dynamics have
104 been directly measured during phytoplankton blooms (Biddanda and Benner, 1997; Carlson et
105 al., 2000; Wetz and Wheeler, 2003); P fractionated studies are scarce. We reported DOP_{op}
106 production and decomposition during phytoplankton blooms in the subarctic Pacific
107 (Yoshimura et al., 2014), but the dynamics of fractionated particulate P pools have not
108 received sufficient research attention to enable a clearer understanding of P biogeochemistry.
109 Fractionations of particulate P may be a key strategy to understand why P is preferentially
110 regenerated from particulate materials produced by phytoplankton relative to C and N (Engel
111 et al., 2002; Paytan et al., 2003; Yoshimura et al., 2009). Observations for the relationship
112 between the temporal dynamics of different P pools and that of phytoplankton will provide
113 insights to improve our understanding of the role of P in the marine environment.

114 The present study focused on measuring the dynamics of the four P pools during a
115 model phytoplankton bloom in the open ocean. This study was conducted during a mesoscale
116 in situ iron (Fe) enrichment experiment in the eastern subarctic Pacific, subarctic ecosystem
117 response to iron enrichment study (SERIES). This was the first experiment to demonstrate the

118 dynamics of an Fe induced phytoplankton bloom from its evolution through to its termination
119 during the 26-day observation using data from three research vessels (Boyd et al., 2004). We
120 observed the phytoplankton bloom aboard the F.R.V. *Kaiyo-Maru* from day 15 to day 26,
121 which corresponded to the peak and decline phase of the phytoplankton bloom (Saito et al.,
122 2006). We discuss temporal variations of the four operationally defined P pools of PIP_{op},
123 POP_{op}, SRP, and DOP_{op} with that of the phytoplankton biomass in terms of chlorophyll-*a*
124 (Chl-*a*) concentrations during the peak and decline phases of the SERIES phytoplankton
125 bloom.

126

127 **2. Materials and methods**

128 *2.1. SERIES experiment*

129 A mesoscale in situ Fe enrichment experiment (SERIES) was conducted in the
130 Alaskan gyre of the eastern subarctic Pacific (50.14°N, 144.75°W), north west of Ocean
131 Station Papa, from 9 July to 4 August 2002 by R.Vs. *J.P. Tully*, *El Puma*, and *Kaiyo-Maru*
132 (Boyd et al., 2004). Dissolved Fe (387 kg of Fe as FeSO₄) and the inert tracer SF₆ were
133 injected into the surface mixed layer over an area of 8.5 × 8.5 km from the *J.P. Tully* on 9 July
134 2002 (denoted as day 0). The Fe infusion increased the in situ concentration of dissolved Fe
135 from < 0.1 nmol L⁻¹ to > 1 nmol L⁻¹ (Boyd et al., 2004). A second release of dissolved Fe
136 (102 kg of Fe) without SF₆ was conducted aboard the *J.P. Tully* in response to the declining
137 dissolved Fe levels on 16 July 2002 (day 7), increasing the in situ concentration of dissolved
138 Fe to ~0.6 nmol L⁻¹ (Boyd et al., 2004). The Fe enriched patch was tracked by elevated SF₆
139 (until day 13) or decreased fCO₂ (after day 15) for 26 days (Law et al., 2006).

140

141 *2.2. Field samplings*

142 Seawater samplings for the fractionated P measurements were conducted inside
143 (IN-patch) and outside (OUT-patch) the Fe enriched patch during days 15–26 aboard the
144 *Kaiyo-Maru*. Samples for P fractionations were not obtained during days 0–14 during the
145 observations of the other vessels. The positions of IN- and OUT-patch stations were
146 determined to verify the center of the patch and outside the patch, respectively, by nighttime
147 horizontal patch surveys (Law et al., 2006). Since the samples collected from OUT-patch
148 were not traced with SF₆, we mainly focus on the IN-patch data. Discrete water samples were
149 collected at predetermined depths (2, 5, 10, 20, 30, 50, 75, 100, 125, 150, and 200 m) in 10 L

150 Niskin-X bottles attached on a Kevlar wire or 12 L Niskin-X bottles attached to a
151 CTD-carousel multiple sampler system. For dissolved P, subsamples were drawn from the
152 Niskin bottles by gravity-filtration through an in-line 47 mm Whatman GF/F filter
153 (precombusted at 450 °C for 4 h), attached directly to the Niskin bottle's spigot. The filtered
154 subsamples were collected in acid-cleaned polypropylene bottles, and stored at -20 °C until
155 analysis on land. For particulate P analysis, subsamples (600–3000 mL aliquots) were filtered
156 through duplicate precombusted and acid-washed 25 mm GF/F filters under gentle vacuum at
157 < 0.01 MPa; one for TPP and one for PIP_{op}. After the filtrations, the filters were washed with
158 0.17 mol L⁻¹ Na₂SO₄ (Solórzano and Sharp, 1980), and stored at -20 °C until analysis on land.
159 For Chl-*a* analysis, subsamples (116 mL aliquots) were filtered through 25 mm GF/F filter
160 under gentle vacuum, and the Chl-*a* on the filter was extracted immediately with 90 %
161 acetone at 4 °C in the dark for 24 h.

162

163 2.3. Chemical analyses

164 P concentrations were measured for four operationally defined pools, DOP_{op}, SRP,
165 POP_{op} and PIP_{op}. The concentration of DOP_{op} was estimated as the difference between total
166 dissolved P (TDP) and SRP concentrations. SRP was measured manually by the molybdenum
167 blue method (Hansen and Koroleff, 1999) using a 50 mm path length quartz cell and a
168 spectrophotometer (U-2001, Hitachi). The calibration was performed using KH₂PO₄
169 (Suprapur, Merck). Samples for the TDP analysis were autoclaved in an acid potassium
170 persulfate (N and P analysis grade, Wako) solution at 123 °C for 120 min (Hansen and
171 Koroleff, 1999; Ridal and Moore, 1990). TDP concentrations were measured as SRP after
172 removing excess free chlorine by placing the samples in a hot water bath for 2 h. Analyses for
173 SRP and TDP were done in triplicate for each sample, and the mean ± 1 standard deviation
174 (SD) was reported for SRP and DOP_{op}. The detection limit for SRP and TDP, given as three
175 times the standard deviation of 10 blank measurements, was 0.01 μmol L⁻¹. Accuracy of our
176 SRP measurements was confirmed using reference materials for nutrients (Aoyama et al.,
177 2012). The precision of the DOP_{op} concentration for a single sample analysis was typically ±
178 0.02 μmol L⁻¹ (ranging ± 0.00 μmol L⁻¹ to 0.04 μmol L⁻¹). Although we cannot evaluate the
179 accuracy of our DOP_{op} measurements due to a lack of appropriate RMs for DOP_{op} analysis
180 (Yoshimura, 2013), a good comparability in our analytical results was confirmed with stable
181 DOP_{op} results (± 0.01 μmol L⁻¹) of the successive measurements of a batch of aged surface
182 seawater sample (Yoshimura and Sharp, 2010). Coefficient of variance for DOP_{op} analysis

183 was 0–23 % for 0–50 m depth samples (roughly $> 0.10 \mu\text{mol L}^{-1}$) and 0–100 % for 75–200
184 m depth samples (roughly $< 0.10 \mu\text{mol L}^{-1}$). TPP was measured as SRP after
185 high-temperature combustion and acid hydrolysis of the filter samples as described by
186 Solórzano and Sharp (1980). PIP_{op} was extracted from filter samples with 1 N HCl at 20 °C in
187 the dark for 24 h and quantified as SRP (Aspila et al., 1976). Analyses for TPP and PIP_{op} were
188 done on single samples that were analyzed in duplicate and the mean \pm range is reported.
189 POP_{op} concentrations were calculated as the difference between TPP and PIP_{op} . A higher
190 precision ($\pm < 1 \text{ nmol L}^{-1}$) was obtained for TPP and PIP_{op} analyses due to the ca. 30 fold
191 concentration factor used in the analytical procedure. Please note that these operationally
192 defined P pools (PIP_{op} , POP_{op} , SRP, and DOP_{op}) do not necessarily correspond to the
193 conceptual P pools (PIP, POP, DIP, and DOP).

194 Chl-*a* concentrations were measured onboard with a fluorometer (Model 10-AU,
195 Turner Designs) with the acidification method of Holm-Hansen et al. (1965). Temporal
196 changes in vertical profiles of Chl-*a* have already been presented in Saito et al. (2006).

197

198 **3. Results and discussion**

199 *3.1. Development and decline of the SERIES phytoplankton bloom*

200 As demonstrated in Saito et al. (2006), our observations during days 15–26
201 corresponded to the peak on day 17 and thereafter the declining phase of the phytoplankton
202 bloom. The SERIES experiment started with a maximum surface Chl-*a* concentration of 0.3
203 $\mu\text{g L}^{-1}$ before Fe enrichment (Marchetti et al., 2006). The surface Chl-*a* concentration reached
204 6.3 $\mu\text{g L}^{-1}$ on day 15 and then decreased to 1.2 $\mu\text{g L}^{-1}$ on day 26 in our observation in the
205 IN-patch, indicating the development and decline of an Fe-induced phytoplankton bloom in
206 contrast to the lower Chl-*a* concentrations in the OUT-patch (0.4–0.5 $\mu\text{g L}^{-1}$ with
207 occasionally high values of 1.8 $\mu\text{g L}^{-1}$ on day 12 and 1.7 $\mu\text{g L}^{-1}$ on day 26). Temporal
208 changes in Chl-*a* concentration occurred in the upper 50 m layer (Fig. 2a), thus the bloom
209 dynamics can be shown as the changes in inventories in 0–50 m depth for Chl-*a* as well as P
210 pools. This corresponds to the variation of upper mixed layer depth (10–38 m) during days
211 2–19 (Marchetti et al., 2006) and of the main pycnocline (30–45 m) during our observation
212 (Saito et al., 2006). When the data obtained by Institute of Ocean Sciences (IOS), Fisheries
213 and Oceans Canada (F.A. Whitney, Personal communication) were combined with our data,
214 the Chl-*a* inventory integrated for 0–50 m depth in the IN-patch was shown to have gradually

215 increased and peaked on day 17, and then decreased until day 26 (Fig. 3a).

216 The SERIES bloom was terminated by simultaneous limitation of diatom growth by
 217 both Fe and silicic acid (Boyd et al., 2005). Boyd et al. (2005) showed that Fe limitation was
 218 detected on day 13 and silicic acid depletion was observed by day 17. On the other hand,
 219 surface SRP concentrations decreased from $1.3 \mu\text{mol L}^{-1}$ on day 0 to $0.5 \mu\text{mol L}^{-1}$ on day 19
 220 in the IN-patch (Fig. 2e) and ranged $0.9\text{--}1.2 \mu\text{mol L}^{-1}$ in the OUT-patch, indicating P
 221 sufficient conditions throughout the experiment even at the peak of the bloom. SRP inventory
 222 for 0–50 m depth decreased with the Chl-*a* increase and then increased toward an initial level
 223 in our observation in the IN-patch (Fig. 3b). Although bacterial activities will have played a
 224 role for the rapid increase in the SRP inventory during the declining phase, data on bacterial
 225 processes reported are available only until day 19 (Hale et al., 2006). Unlike with Chl-*a*, SRP
 226 inventories were not distinguishable between IN and OUT-patch before our observation (Fig.
 227 3b). This is explained by the intrusion of a high nutrient water mass beneath the surface mixed
 228 layer of the IN-patch (Timothy et al., 2006), which was also detected in our SRP vertical
 229 profiles (Fig. 2e). Our data confirmed that phytoplankton grew under conditions that were not
 230 limited by P in both IN and OUT-patch.

231

232 3.2. P partitioning during the peak of the SERIES bloom

233 Since particulate P has been measured as TPP in past studies (e.g. Yoshimura et al.,
 234 2014), this is the first report to demonstrate partitioning of consumed SRP into POP_{op} , PIP_{op} ,
 235 and DOP_{op} during the growth phase of a bloom. Although we have no data for the P pools
 236 other than SRP in the initial stage of the SERIES bloom, we considered that data on day 15 in
 237 the OUT-patch approximate the values in the initial stage in the IN-patch. This assumption is
 238 supported by the comparable values of Chl-*a* inventories of 16.5 mg m^{-2} for 0–50 m on day 0
 239 in the IN-patch observed by IOS and of 17.0 mg m^{-2} on day 15 in the OUT-patch observed in
 240 the present study (Fig. 2a). With this simple assumption, to estimate the net productions of P
 241 pools from the initial stage to the peak of the bloom on day 17, the values on day 15 in the
 242 OUT-patch were subtracted from those on day 17 in the IN-patch. We estimated net
 243 productions of POP_{op} , PIP_{op} , and DOP_{op} at the peak of the bloom as 4.1 ± 0.2 , 1.8 ± 0.1 , and
 244 $0.9 \pm 0.9 \text{ mmol m}^{-2}$, and their proportions as 60, 27, and 13 %, respectively. Our data show
 245 that POP_{op} and PIP_{op} were newly produced at a ratio of 7:3. The PIP_{op} pool in phytoplankton
 246 was assumed to be composed of intracellular stored P as PO_4 , pyro-, and polyphosphate

247 (Labry et al., 2013; Paytan et al., 2003), and adsorbed P onto phytoplankton cell surfaces (Fu
 248 et al., 2005) (Fig. 1). Note that a significant part of pyro- and polyphosphate would be
 249 measured as POP_{op}, which is basically composed of P-containing cell components such as
 250 phosphoesters and nucleotides (Fig. 1). Although a part of the P incorporated into
 251 phytoplankton cells can be converted into DOP_{op} through several possible processes of
 252 autotrophic and heterotrophic activities as described for dissolved organic C (Nagata, 2000),
 253 observed DOP_{op} production was within the range of uncertainty of analytical precision (Fig.
 254 4d). Considering DOP/Chl-*a* production ratios during subarctic phytoplankton blooms are
 255 reported as 0.0027–0.0044 mol/g (Yoshimura et al., 2014), we confirmed that the estimated
 256 DOP production of 0.016–0.026 $\mu\text{mol L}^{-1}$ under a condition of 6 $\mu\text{g L}^{-1}$ of Chl-*a* production
 257 in SERIES surface waters is within the level of the precision of our DOP analyses as
 258 mentioned in section 2.3.

259 Note that the amount of net production of POP_{op}, PIP_{op}, and DOP_{op} (6.8 mmol m^{-2})
 260 at the peak of the SERIES bloom did not match with the net consumption of SRP (16.8 mmol
 261 m^{-2}) even if we consider TPP sinking export flux of 2.6 mmol m^{-2} during day 3.5 to day 19.5,
 262 estimated from the data for particulate organic C sinking flux of 278 mmol m^{-2} reported in
 263 Timothy et al. (2006) and the Redfield ratio of C:P = 106:1 (Redfield et al., 1963). Net
 264 increased copepod biomass in 0–200 m in the IN-patch was estimated to be 0.25–0.54 mmol P
 265 m^{-2} using reported data on an increased C biomass of 0.6 gC m^{-2} (Tsuda et al., 2006) and C:P
 266 molar ratios of 93–204 for *Pseudocalanus* sp. (Gismervik, 1997), and does not compensate
 267 the imbalanced P budget. Lateral patch evolution and patch dilution with surrounding waters
 268 have significant impacts on materials in the IN-patch of SERIES, as also shown for another in
 269 situ Fe enrichment experiment (Law et al., 2006). Timothy et al. (2006) showed that
 270 consideration of patch expansion improves the balance of the budgets of C, N, and silicon, but
 271 was unable to show balanced budgets for them. Since similarly for P this is a likely outcome,
 272 we do not further discuss quantitative analysis on the P budget in this paper.

273

274 3.3. Uncoupled behavior of POP and PIP between before and after the SERIES 275 bloom

276 After the peak of the bloom, we found that POP_{op} and PIP_{op} dynamics were not
 277 coupled. This was typically observed in the different timing of the peaks of POP_{op} (day 17;
 278 Fig. 4b) and PIP_{op} (day 19; Fig. 4c), two days later than the peak of Chl-*a* and POP_{op}. In

279 contrast to a significant correlation between POP_{op} and $\text{Chl-}a$ (Spearman's rank correlation, p
280 < 0.05) during our observation period, PIP_{op} did not show a clear correlation with $\text{Chl-}a$
281 (Spearman's rank correlation, $p > 0.1$). In addition, rapid change in each proportion in the TPP
282 pool occurred between before and after the peak of the bloom (Fig. 5). While PIP_{op} in the
283 IN-patch until the peak of the bloom and OUT-patch composed 25 % of the TPP pool, the
284 proportions increased significantly (ANOVA, Tukey's test, $p < 0.01$) to 50 % on average
285 during days 19–26 in the IN-patch (Fig. 5). The uncoupled behavior of POP_{op} and PIP_{op}
286 between before and after the peak of the bloom is a highlight of this study.

287 Physiological status of phytoplankton can explain the difference in the PIP_{op}
288 proportion. Phytoplankton communities showed a relatively high net growth rate ($0.2\text{--}0.3$
289 day^{-1}) on days 15–17 of the IN-patch (Boyd et al., 2005) and may show a relatively high
290 growth rate in the OUT-patch due to low Fe adapted small-sized phytoplankters and closely
291 balanced algal growth-microzooplankton grazing systems in Fe limited waters, as shown in
292 Liu et al. (2002). Under these conditions, we found a similar PIP_{op} proportion of 20–30 % for
293 the IN and OUT-patch although size compositions of $\text{Chl-}a$ were different with dominance of
294 $> 20 \mu\text{m}$ fraction on days 15–17 of the IN-patch and of $< 20 \mu\text{m}$ fraction in the OUT-patch
295 (Saito et al., 2006). Similar values for the PIP_{op} proportion of 10–20 % were observed in
296 North Pacific surface waters (Yoshimura et al., 2007), consistent with the relatively high
297 growth rate of ambient phytoplankton communities which is closely coupled with high
298 microzooplankton grazing rates (Calbet and Landry, 2004; Liu et al., 2002). On the other hand,
299 elevated PIP_{op} proportions of 50 % were observed for phytoplankton communities with a
300 depressed growth rate ($0.1\text{--}0.2 \text{day}^{-1}$) on days 19–26 of the IN-patch (Boyd et al., 2005),
301 although $> 20 \mu\text{m}$ fraction still dominated the communities (Saito et al., 2006). Although
302 higher PIP_{op} proportions are expected to be observed toward the end of the bloom, the
303 proportion peaked on day 19 and did not further increase during the declining phase (Fig. 5).
304 This is the period of rapid decrease in TPP pools (Fig. 4a), therefore more senescent
305 phytoplankton cells with higher PIP_{op} proportion would sink out or be decomposed. PIP_{op}
306 proportions observed in the declining phase are relatively close to the proportions reported for
307 sinking particulate matter with typical values of 50–70 % (Benitez-Nelson et al., 2007; Lyons
308 et al., 2011; Paytan et al., 2003). The sinking particles were mainly composed of detritus
309 including dead phytoplankton cells. PIP_{op} proportions in phytoplankton may increase with
310 depressed growth conditions.

311 In our observation, TPP inventory was relatively stable from day 17 at which POP_{op}

312 peaked to day 19 at which PIP_{op} peaked (Fig. 4a), suggesting that decreased POP_{op} was
313 replaced by PIP_{op} through intracellular P metabolisms. In phytoplankton cells, many enzymes
314 such as alkaline phosphatase, polyphosphate kinase, and exophosphatase work to transport P
315 between key P-containing molecules (Lin et al., 2016). If these enzymes are still active when
316 phytoplankton growth has slowed down, the enzyme activities contribute to the
317 remineralization of organic P compounds to PO₄. Since we must consider that POP_{op} contains
318 polyphosphates due to an analytical restriction, production of PO₄ from polyphosphate may
319 contribute to transform POP_{op} to PIP_{op}. This is like an intracellular autolysis of P compounds.
320 This is reasonable to explain why P is preferentially regenerated from particulate materials
321 relative to C and N as reported in past studies (Engel et al., 2002; Paytan et al., 2003;
322 Yoshimura et al., 2009). In addition, more PO₄ is scavenged onto phytoplankton cell surface
323 under the growth rate depressed conditions (Fu et al., 2005), leading to increased PIP_{op}
324 proportions. Days 17–19 represented the period with higher bacterial biomass and production,
325 but the substantial change did not occur between the values on day 17 and day 19 when a
326 rapid shift occurred in PIP_{op} proportions. Bacterial and zooplankton activities play roles to
327 remineralize POP_{op}, but these should be detected as PO₄ increase, which was observed as a
328 SRP inventory increase in our observations (Fig. 3b). Although increases in detritus
329 concentration may change PIP_{op} proportions, no data are available to estimate the
330 contributions of detritus in particulate matter during SERIES. Discussions of the dynamics of
331 particulate P pools with particulate organic C and N might enable a clearer understanding of
332 bioactive element cycles and their interactions, direct measurements of particulate organic C
333 and N during the SERIES experiment are not available to our knowledge. We suggest that
334 physiological status of the phytoplankton assemblage determines the POP_{op}:PIP_{op} ratio in the
335 TPP pool.

336 Another plausible explanation for the changes in POP_{op}:PIP_{op} ratios other than
337 physiological mechanisms is the floral shift in the dominant diatoms around day 17 from
338 *Chaetoceros* spp. and *Thalassiosira* spp. to *Thalassiothrix longissima* (Boyd et al., 2005).
339 Although different species could have different cellular POP_{op}:PIP_{op} ratios, this has not yet
340 been fully examined. Using different P fractionation methods, Miyata et al. (1986) reported
341 that the proportions of PIP (orthophosphate + acid-soluble polyphosphates + acid-insoluble
342 polyphosphates) to total cellular P have a range from ca. 30 % in P limited to 40 % in N
343 limited chemostat culture of *Skeletonema costatum* and from 24 % to 35 % in that of
344 *Heterosigma akashiwo*, indicating relatively small interspecific difference. On the other hand,

345 using ^{31}P NMR spectroscopy, Cade-Menun and Paytan (2010) has reported relatively wide
346 ranges of PIP proportion (orthophosphate + pyrophosphates + polyphosphates) to total
347 cellular P for 12 algal species from 40 % to 82 % under optimal growth conditions with some
348 significant change under stressed conditions. Since a direct comparison between the results
349 from different methods is not straightforward, further data are required for the method used in
350 this study. However note that similar $\text{POP}_{\text{op}}:\text{PIP}_{\text{op}}$ ratios were observed between days 15–17
351 of the IN-patch with dominance of $> 20 \mu\text{m}$ diatoms and the OUT-patch with dominance of $<$
352 $20 \mu\text{m}$ phytoplankton such as *Synechococcus*, Prasinophyceae, and Prymnesiophyceae
353 (Marchetti et al., 2006) (Fig. 5), suggesting that different phytoplankton species and groups
354 have a similar $\text{POP}_{\text{op}}:\text{PIP}_{\text{op}}$ ratio under P sufficient conditions in the studied area.

355

356 **4. Conclusion**

357 We found uncoupling of POP_{op} and PIP_{op} dynamics during the peak and decline
358 phase of the phytoplankton bloom. $\text{POP}_{\text{op}}:\text{PIP}_{\text{op}}$ ratio of suspended particulate matter may be a
359 key factor to understand subsequent P regeneration. In this regard, characterization of the
360 composition of PIP_{op} (PO_4 or polyphosphates) will enable a clearer understanding of P
361 biogeochemistry. TPP fractionation provides powerful insights to better understand the marine
362 P cycle. In addition, interspecific differences in cellular $\text{POP}_{\text{op}}:\text{PIP}_{\text{op}}$ ratio and changes in the
363 ratio during different growth stages of the phytoplankton from exponential growth through
364 stationary to decline phase should be clarified in future comprehensive studies including
365 culture experiments. Recent development of a sensitive analytical method for the
366 fractionations of POP_{op} and PIP_{op} reported by Ehama et al. (2016) will enable further progress
367 this research field.

368

369 **Acknowledgments**

370 We acknowledge the field assistance of the captain, officers, crew, and scientists
371 aboard the *Kaiyo-Maru*. We thank Institute of Ocean Sciences, Fisheries and Oceans Canada
372 for the usage of Chl-*a* and SRP data, K. Sugita for the assistance in the laboratory on land, C.
373 Norman for his help to improve the English of the manuscript. This work was supported by a
374 grant from CRIEPI (U00024).

375

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534 Figure Legends

535 Fig. 1. A diagram of the differentiations of typical P compounds into conceptually and
536 operationally defined P pools. Positions of each component among the operationally defined P
537 pools are approximately illustrated with reference to Labry et al. (2013) for particulate P and
538 to Baldwin (1998) and Thomson-Bulldis and Karl (1998) for dissolved P.

539

540 Fig. 2. Temporal variations in the vertical profiles for selected sampling days of chlorophyll-*a*,
541 particulate P, and dissolved P in the Fe enriched patch. Mean values for duplicate analysis for
542 particulate P and triplicate analysis for dissolved P are reported. Error bars are within the
543 symbols except for DOP_{op} and are not shown for DOP_{op} to simplify the figure. The figure for
544 Chl-*a* was redrawn using the data presented in Saito et al. (2006).

545

546 Fig. 3. Temporal variations in the 0–50 m inventories for chlorophyll-*a* and soluble reactive P
547 (SRP) in the IN-patch and OUT-patch. Chl-*a* and SRP data during day 0 to day 14 are
548 provided by Institute of Ocean Sciences, Fisheries and Oceans Canada (F.A. Whitney,
549 Personal communication). The data in gray background during day 15 to day 26 was obtained
550 in our study. Chl-*a* inventory was calculated based on the data presented in Saito et al. (2006).

551

552 Fig. 4. Temporal variations in the 0–50 m inventories for the particulate P and dissolved
553 organic P (DOP_{op}) in the IN-patch and OUT-patch. Integrated values were calculated using
554 results of each depth (mean \pm range of duplicate analysis for particulate P and mean \pm 1
555 standard deviation of triplicate analysis for DOP_{op}). Error bars are within the symbols except
556 for DOP_{op}.

557

558 Fig. 5. Temporal variations in the relative composition of fractionated particulate P in the
559 0–50 m inventories for the IN-patch and OUT-patch.

Fig. 1

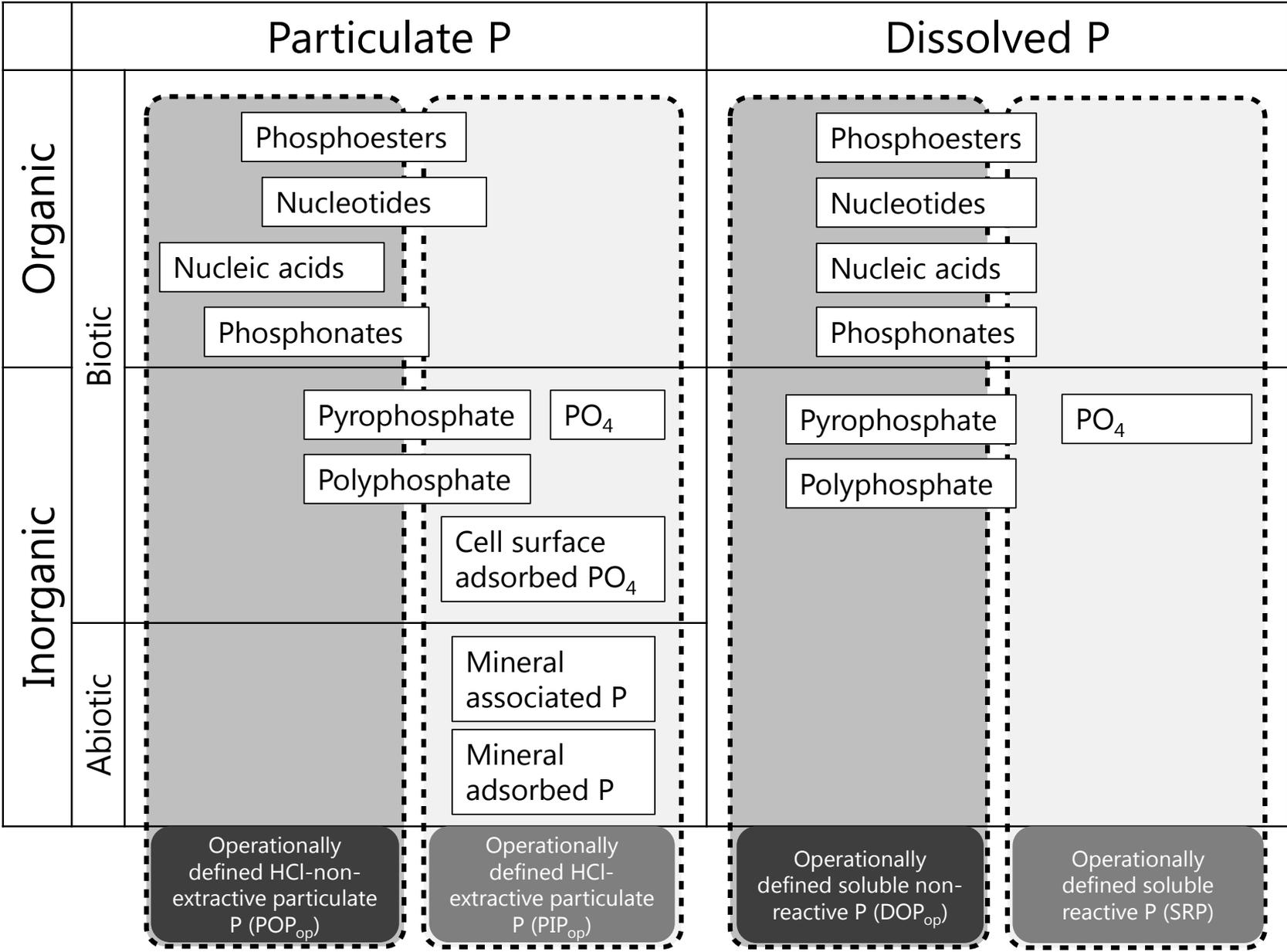


Fig. 2

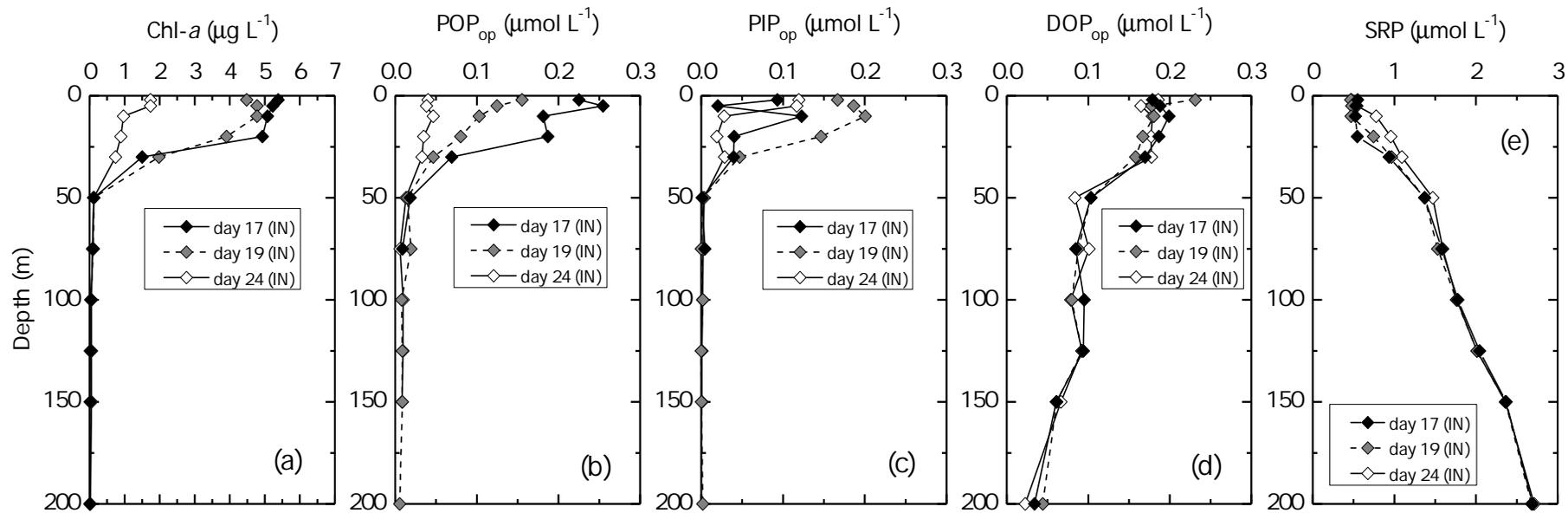


Fig. 3

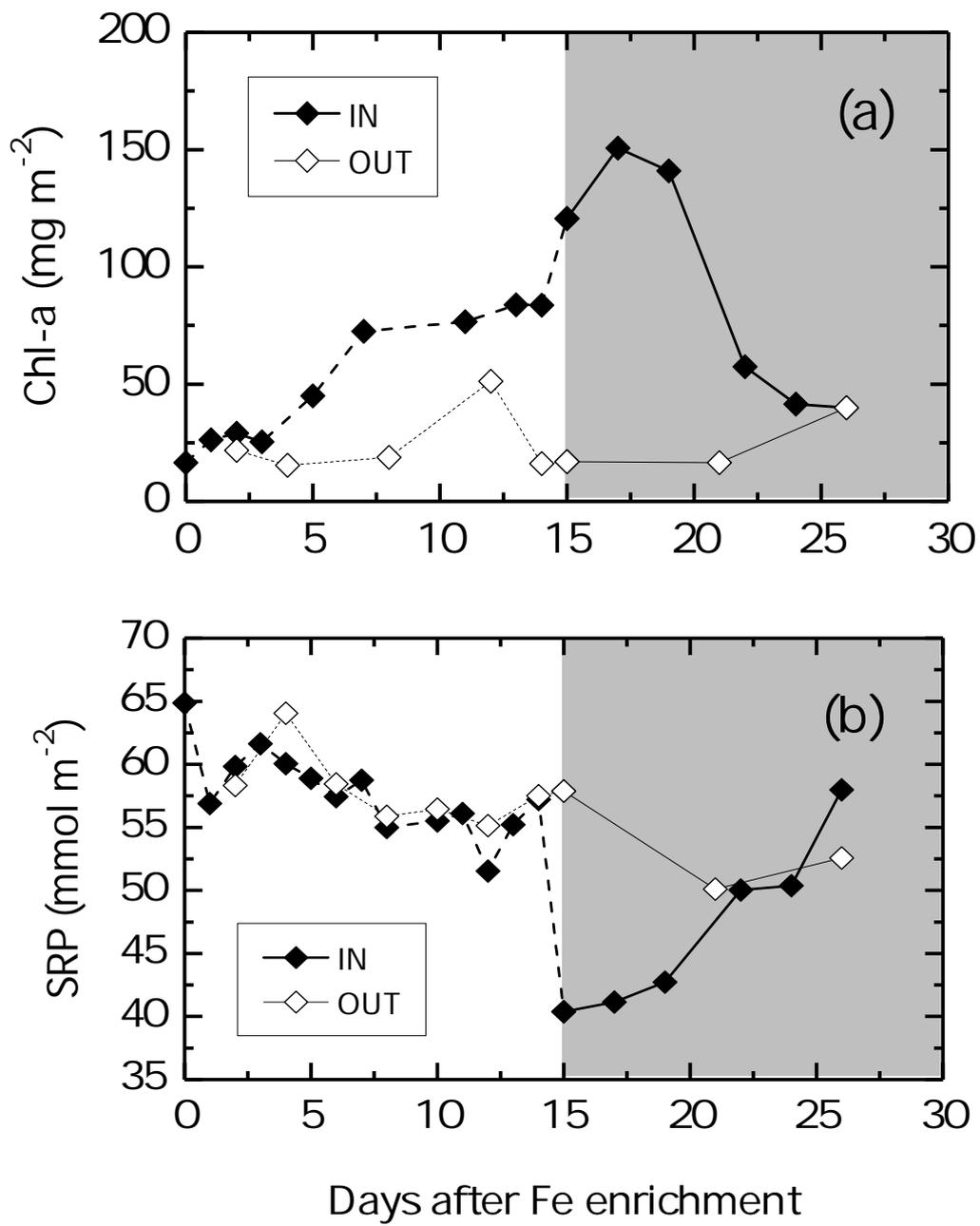


Fig. 4

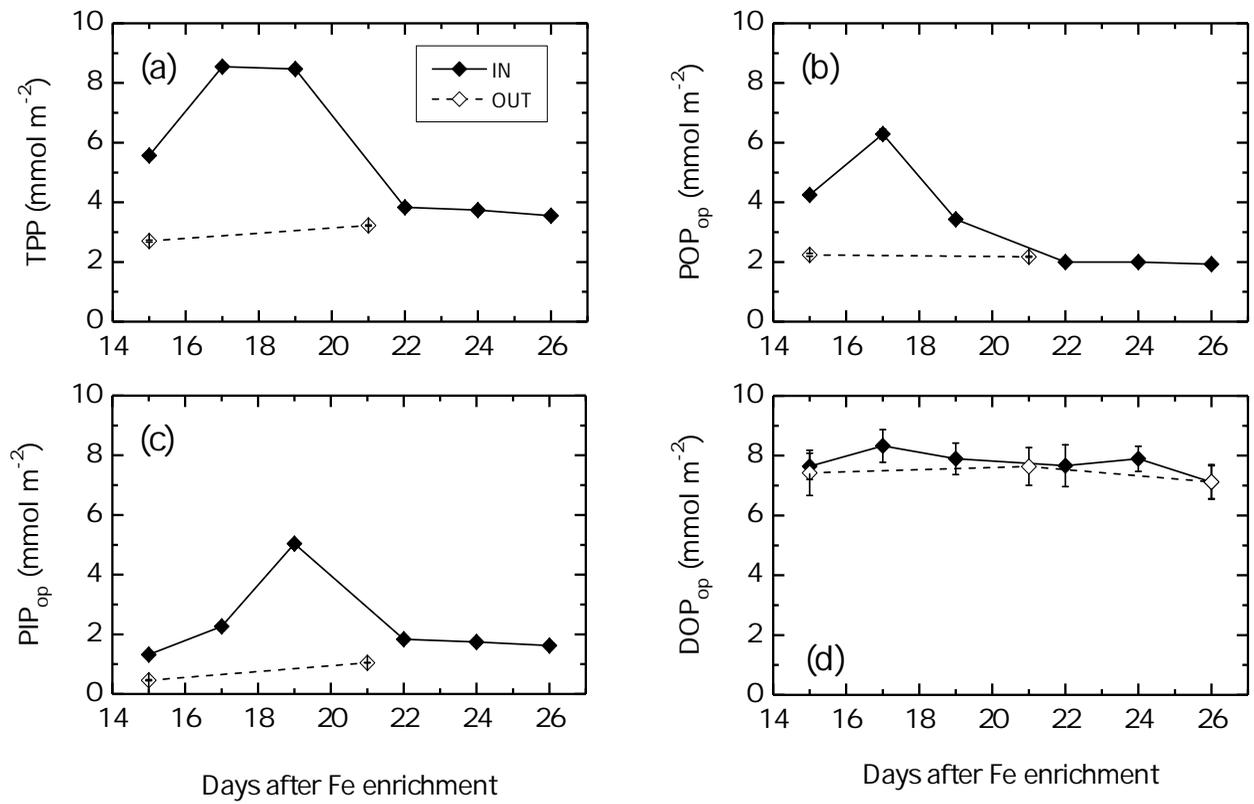


Fig. 5

