



- 1 Exploring the distance between nitrogen and phosphorus
- 2 limitation in mesotrophic surface waters using a sensitive
- 3 bioassay
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13 Abstract

14 The balance in microbial consumption of nitrogen and phosphorus was investigated in 15 samples collected in two mesotrophic coastal environments: the Baltic Sea (Tvärminne 16 field station) and the North Sea (Espegrend field station). For this, we have refined a bioassay based on the response in alkaline phosphatase activity (APA) over a matrix of 17 18 combinations in nitrogen and phosphorus additions. This assay not only provides 19 information on which element (N or P) is the primary limiting, but also gives a 20 quantitative estimate for the excess of the secondary limiting element (P^* or N^* , 21 respectively), as well as the ratio between N and P consumption over short time scales 22 (days). As expected for a Baltic Sea late spring-early summer situation, the Tvärminne 23 assays (n=5) indicated N-limitation with an average $P^*=0.30\pm0.10 \mu$ M-P, when incubated for 4 days. For short incubations (1-2 days), the Espegrend assays indicated P-24 25 limitation, but the shape of the response surface changed with incubation time, resulting 26 in a drift in parameter estimates toward N-limitation. Extrapolating back to zero 27 incubation time gave P-limitation with N^{*} \approx 0.9 μ M-N. The N:P ratio (molar) of nutrient 28 consumption varied considerably between investigated locations; from 2.3 ± 0.4 in the Tvärminne samples to 13±5 and 32±3 in two samples from Espegrend. Our assays 29 30 included samples from mesocosm acidification experiments, but statistically significant 31 effects of ocean acidification were not found by this method.

Keywords: alkaline phosphatase activity, bioassays, mesotrophic temperate seas,nutrient limitation, phytoplankton

34 **1 Introduction**

N to P balance is a core biogeochemical feature of aquatic systems as highlighted in Redfield's classical question of whether it is the chemistry of seawater that has determined the stoichiometry of the marine organisms, or biology is the cause for the "normal" 16:1 (molar) ratio between N and P in seawater (Redfield et al., 1963). The





1 issue of surface ocean nutrient limitation is as acute as ever (Moore et al., 2013), since it 2 has bearings on phenomena ranging from the global carbon cycle, where it plays a key 3 role in the dynamics of the ocean's biological pump (Ducklow et al., 2001); via basin 4 scale issues such as N deficiency in Arctic water of Pacific origin (Lehmann et al., 2005), 5 P deficiency in Eastern Mediterranean deep waters (Krom et al., 1991) and the North 6 Atlantic gyre (Mather et al., 2008); via regional issues such as the question of P and/or N 7 removal from the Baltic Sea (Elmgren and Larsson, 2001; Granéli et al., 1990; Räike et 8 al., 2003); to local ecosystem characteristics such as P-deficient brackish layer 9 overlaying potentially more N-limited marine waters in the fjords of western Norway 10 (Thingstad et al., 1993). The classical idea of predominantly N-limitation in marine 11 systems (as opposed to predominantly P-limitation in limnic systems) (Hecky and 12 Kilham, 1988) has also become considerably more nuanced, not only due to the cases 13 mentioned above, but also with the identification of the High Nutrients Low Chlorophyll (HNLC) areas as being iron-limited (Franck et al., 2003), phosphorus and iron as co-14 15 limiting elements of nitrogen fixation in the tropical North Atlantic (Mills et al., 2004) 16 and N₂ fixers having a competitive advantage in oligotrophic P-starved regions (Landolfi 17 et al., 2015). While some of the mechanisms behind these apparent deviations from 18 Redfield stoichiometry seem to be well understood, there are others which lack 19 generally accepted explanations.

20 In deep waters with most of the bioavailable N and P converted to NO_3 and PO_4 , the 21 chemical determination of N or P in excess of the Redfield ratio may be relatively 22 straight forward. In biogeochemistry this excess is referred to as N* and P* (Gruber and 23 Sarmiento, 1997). In productive surface waters this is a more complex issue. Not only 24 may both N and P accumulate in pools of DON and DOP with different grades of 25 bioavailability, but organisms may also have flexible stoichiometry, and groups of 26 organisms may differ in stoichiometry, theoretically allowing for the adjustment of the 27 N:P ratio in nutrient consumption at both an individual and a community level 28 (Bertilsson et al., 2003; Geider et al., 2002; Krauk et al., 2006). A potential solution to the 29 chemically intractable problem of measuring a large suite of presumably bioavailable 30 pools is to use a quantitative bioassay, i.e. to ask the organisms how much of limiting 31 elements they can "see".

32 Microorganisms have evolved sophisticated physiological mechanisms to adapt to the 33 different forms of nutrient limitation (Geider et al., 1997; Ivančić et al., 2012; Thingstad 34 et al., 2005; Van Mooy et al., 2009). Activation of the genes found in the inducible Pho-35 operon (Torriani-Gorini, 1994) is an indicator of the organism responding to P 36 deficiency. One of the enzymes induced as a part of this operon is the extracellular 37 enzyme alkaline phosphatase, used to split the ester bound in phosphomonoesters 38 (Hoppe, 2003). Since the alkaline phosphatase activity (APA) can easily and sensitively 39 be determined fluorometrically (Perry, 1972), it is used as a convenient indicator of the 40 Pho-operon being de-repressed, and therefore of P deficiency in aquatic systems. This 41 method was further exploited by Thingstad and Mantoura (2005) in the oligotrophic 42 Eastern Mediterranean, showing that the concentration of added PO₄ needed for APA to disappear in a P-limited system, or alternatively NH4 needed to induce APA in an N-43 44 limited system, could be used as a bioassay to quantitatively estimate N* or P*. We here 45 expand this technique by using a matrix-setup including simultaneous gradients in both PO₄ and NH₄ additions. This is applied to samples from the coastal waters of western 46 47 Norway and the Baltic Sea, confirming that the assay gives informative results also in 48 temperate, mesotrophic environments.





1 2 Material and Methods

2 2.1 Study areas and sampling

3 The Baltic water was collected as integrated samples (depth 0–10 m) in Storfjärden near Tvärminne field station (59° 51.50' N, 23° 15.50' E) on 6 August 2012. The collection 4 5 was performed 45-30 days after the first-last CO_2 treatments and 50 days after the 6 mesocosm closure (Paul et al., 2015). Samples were collected from the fjord (417 µatm) 7 and mesocosms M1 (365 µatm), M3 (1007 µatm), M6 (821 µatm) and M7 (497 µatm); 8 where numbers in parentheses are average $f(CO_2)$ over the period Day 1-Day 43. The 9 mesocosms received no nutrient manipulations except the CO₂ treatments. Further 10 details about location and the experiment can be found in Paul et al. (2015).

11 The samples from western Norway were collected during a similar mesocosm 12 experiment in Raunefjorden close to Espegrend field station (60° 16.2' N, 5° 11.7' E). 13 From one mesocosm (MR) an integrated (depth 0-20 m) sample (1165 µatm) was 14 collected on 25 May 2015 corresponding to Day 22 after acidification treatment. The 15 fjord sample was collected at nearby landlocked location Kviturspollen (60° 15.8' N, 5° 16 15' E) at the depth of 1 m using a Niskin sampler on 3 June 2015. Samples were pre-17 filtered through gauze of 112 μ m mesh size to minimize the variability due to the 18 occasional large zooplankton.

19 2.2 Matrices of nitrogen and phosphorus additions

20 Samples were distributed in 15 mL Falcon® polypropylene tubes (BD Biosciences®) 21 organized in 10x10 or 8x8 columns x rows (Tvärminne and Espegrend, respectively). 22 PO_4 (KH₂PO₄ 10 μ M) was added in final concentrations from 0 to 290 nM-P in steps of 23 32.2 nM (Tvärminne) and from 0 to 105 nM-P in steps of 15 nM (Espegrend). Each of the 24 columns received additions of NH₄ (NH₄Cl 200 µM) in final concentrations from 0 to 964 25 nM-N in steps of 107 nM-N (Tvärminne) and from 0 to 2100 nM-N in steps of 300 nM-N 26 (Espegrend). The tubes were incubated in light:dark (16 h:8 h) at 17–18°C (Tvärminne) 27 and in light:dark (12 h:12 h) at 16.5°C (Espegrend), both at irradiance of 78 µmol 28 photons m⁻² s⁻¹. Incubation at Tvärminne lasted 4 days for all samples, whilst APA assays 29 for Espegrend were repeated as given in each case.

30 2.3 Alkaline phosphatase activity

APA in the Baltic Sea was measured by Varian Cary Eclipse fluorometer, while APA in the coastal waters of the western Norway was measured using a PerkinElmer Enspire 2300 plate reader. Measurements of APA were done according to Perry (1972) but modified to the use of fluorescence plate reader by pipetting 200 μL subsamples from each Falcon tube into the wells containing the substrate. Results are expressed in relative fluorescence units (RFU). The need for intercalibration of different fluorometers was thus avoided.

38 2.4 Fitting the response surface

Thingstad and Mantoura (2005) fitted sigmoidal functions to the observed APAresponses; either a decreasing function parallel to the P-addition axis in the case of a Plimited system, or an increasing function parallel to the N-addition axis in the case of Nlimitation. To avoid this pre-fitting choice of function, we here have instead started with





- 1 the assumption that the *P*,*N*-plane is split into a P-limited and an N-limited part by the 2 straight line $N = N_0 + rP$ where a negative value of the intercept N_0 corresponds to the
- $-\frac{N_0}{N_0}$
- 3 excess-N (N^{*}) present in a P-limited system and \overline{r} is the amount of phosphate needed
- 4 to shift the system to N-limitation. Conversely, a positive value of the intercept N_0 would 5 correspond to the amount of N required to shift an N-limited system into P-limitation,
- 6 while $\frac{N_0}{r}$ corresponds to the excess-P (P*) in this N-limited system. The shift from P- to
- 7 N-limitation, and therefore the expression of APA in a point P,N, is assumed to be a
- 8 function of the distance *Z* between this point and the line (Fig. 1). The sigmoidal function

9 fitted is $APA_{est}(P,N) = \frac{A}{1 + e^{sZ}} - B$. Here, the exponential function replaces the term $(Z)^s$

 $\left(\frac{Z}{Z_0}\right)^s$ adopted by Thingstad and Mantoura (2005) from standard toxicology. This 10 standard expression is undefined for $Z_0=0$ and therefore not applicable with our 11 12 approach. Visual inspection of residuals in graphs (see Fig. 2A, B) did not suggest 13 systematic deviances between response surfaces fitted with this function and the 14 observed data. Alternative fitting functions have therefore not been explored. With five 15 parameters to fit (r, N_0, s, A, B) , this leaves 95 and 59 degrees of freedom for the 16 Tvärminne and Espegrend set-ups, respectively. The fitted surface APAest has a 17 maximum A-B obtained for co-ordinates combining low P with high N (large negative Z) 18 and $APA_{est} = (A/2)$ -B along the line $N = N_0 + rP$ separating the P- and N-limited regions. The 19 parameter *s* defines the steepness of transition between the two regions perpendicular 20 to this line. B is the background APA_{est} found for high-P, low-N (large positive Z) co-21 ordinates. The fitting was done using the "fit" function in Matlab® with its default 22 Levenberg-Marquardt algorithm providing the parameter estimates with 95% 23 confidence intervals (c.i.) (code included in SI).

24 3 Results

Two examples of the fitted response surface, one from Tvärminne (Fjord) (Fig. 2A) and one from Espegrend (MR) (Fig. 2B) are shown to illustrate the difference in shape of the response in situations apparently N-limited (Tvärminne) and P-limited (Espegrend), with estimated P*=0.3 μ M-P and N*=0.4 μ M-N, respectively. All assays are summarized in Table 1.

30 For the two Espegrend samples, the change in shape of the response surface with 31 incubation time was explored (Fig. 3). For both samples, N_0 increased with incubation 32 time ($p \le 0.05$, Table 2), i.e. the assay results drifted towards increasing N deficiency 33 when using longer incubation times. In the sample MR, r and s decreased significantly 34 over time (Table 2). Using linear regression, the parameter estimates can be 35 extrapolated back to zero incubation time. With this technique the average P^* for the 36 Tvärminne samples, based on a single incubation time, was 0.3 μ M-P, and the average N^{*} 37 for the two Espegrend samples, based on backward extrapolation, was 0.9μ M-N.

The assays from Tvärminne mesocosms include an f(CO₂) gradient. Linear regressions of

- N_0 (p=0.55), r (p=0.63) (Fig. 4) and s (p=0.19) (not presented) on f(CO₂) gave no
- 40 indication of any statistically significant effect of the 45 days exposure of the systems to
- different CO₂-levels. Compared to a Redfield N:P value of 16, all the Tvärminne samples





- 1 gave low r (2.3±0.5; mean over samples±sd), while the two Espegrend samples gave r of
- 2 13±2 (Kviturspollen) and 32±3 (MR) (mean±sd, both over incubation times).

3 4 Discussion

This study extends the demonstrated applicability of this type of assay from its previous
use in warm oligotrophic waters (Thingstad and Mantoura, 2005) to mesotrophic
temperate environments. We modified the technique so that no *a priori* assumptions are
now required as to whether the system investigated is N- or P-deficient.

8 We explored the use of this modified assay in two environments with anticipated 9 differences in ambient N:P stoichiometry. The Tvärminne mesocosm experiment was 10 planned with the expectation of an N-limited spring-summer situation as characteristic 11 in the Baltic Sea (Granéli et al., 1990; Rolff and Elfwing, 2015; Thomas et al., 2003), 12 subsequently transiting from N-limitation towards an N- and P-co-limited situation as 13 the result of "new" N being added through late summer blooming of diazotrophic cyanobacteria (Lignell et al., 2003). This bloom did not occur during the whole 14 15 Tvärminne experiment and N-limitation at the time of sampling has been confirmed by 16 Nausch et al. (2016) who studied the microbial P-cycle just before our experiment. The 17 finding of positive N_0 -estimates for all 5 samples (Table 1) is in line with this. The 18 Tvärminne assays were performed after 4 days of incubation when the APA-responses 19 emerged. The conclusion of N-limitation is therefore confounded by the potential drift in 20 parameter estimates as observed for the Espegrend samples (Fig. 3). The drift obviously 21 complicates the use of this assay since there may be no single incubation time that gives 22 a "correct" set of parameter values. Since the drift seems to be reasonably linear for all 23 parameters (Fig. 3), a promising option is to extrapolate the linear regressions back to 24 time 0. In our case this gives negative N_0 values of -0.8 (-1.4,-0.2) and -1.0 (-2.4,0.4) for 25 the samples from MR and Kviturspollen, respectively (intercept with 95% c.i.); 26 suggesting initial P-limitation. This conclusion is in accordance with the expectation 27 since the top layer of the fjords in western Norway has been shown to be P-deficient 28 (Thingstad et al., 1993).

29 The mechanisms behind the drift in parameter estimates have not been studied further 30 here. Two, not mutually exclusive, scenarios may, however, illustrate the options: 1) The 31 microbial food web in the incubated tubes remineralizes P faster than N (Garber, 1984). 32 The assay may then correctly reflect the succession of the limiting nutrient in the sense 33 that the bioavailable pools in the tubes change with incubation time; 2) N added in 34 excess of P in the upper P-limited part of the P,N-plane is used by the organisms to 35 produce alkaline phosphatase (rather than biomass). This would lift the response 36 surface for high values of added N and move the fitted line towards higher N_{0} , i.e. 37 towards N-limitation. The use of extra N to produce exo-enzymes for acquisition of P 38 from DOP has recently been argued for, but then with N-fixation as the N-source 39 (Landolfi et al., 2015).

40 The r values representing the ratio of N- and P-consumption are comparable between all 41 the Tvärminne samples $(2.3\pm0.5, n=5 \text{ different samples})$, indicating good reproducibility 42 of the assay for similar water samples. This low value compared to the Redfield value of 43 16 was, however, strikingly different from the Espegrend samples with a Redfield-like 44 13 ± 2 (Kviturspollen) and a high 32 ± 3 (MR), both over incubation times. A similar 45 phenomenon was noted by Thingstad and Mantoura (2005) using this method to study 46 in-out differences in a Lagrangian experiment where orthophosphate was added to the 47 P-deficient surface system in the Eastern Mediterranean. While their P-limited out-





- 1 sample gave an $r=15\pm 2$, the inside system, when driven to N deficiency by the phosphate 2 addition, gave $r=3.0\pm0.2$. Interestingly, we also here found the lower-than-Redfield r 3 values in the probably N-limited samples from Tvärminne. From the limited number of 4 determinations available, the linkage between N deficiency and low r values seems 5 consistent, but the underlying mechanism is not immediately obvious. One could argue 6 that, in a P-deficient system, the organisms present would be expected to have marked 7 luxury consumption of any added P (Thingstad, 2005) (and vice versa for N) (Leonardos 8 and Geider, 2004). As r represents the ratio between utilization of added N and added P. 9 luxury uptake seems to lead to an expected effect on r opposite to that observed. One 10 could speculate that organisms in N-deficient environments are selected for (or adapted 11 to) low N:P requirements. Although this may be in qualitative agreement with our data,
- 12 it seems doubtful that the large range in r (~2 to ~30) can be explained in this manner.

13 5 Conclusions

14 We have demonstrated the extension of the APA assay from the warm oligotrophic to 15 the temperate, mesotrophic waters. This technique has the advantage over traditional nutrient-limitation bioassays because it determines not only the most limiting element, 16 17 but also estimates the quantitative excess in bioavailable forms of the secondary limiting 18 element (N^*, P^*) ; this without any requirement for a determination of the large variety of 19 chemical and/or physical forms in which these excess nutrients exist. The assay is 20 complicated by a drift in parameter estimates with incubation time. A backward 21 extrapolation to zero incubation time appears promising.

22 Data availability

Original data are given in Supplementary Information (SI) for each assay. SI also
 contains the Matlab[®] program for fitting the response surfaces as shown in Fig. 2.

25 Acknowledgements

The cooperation between Enis Hrustić and Professor Tron Frede Thingstad was partially realized through the Erasmus+ training at University of Bergen, Norway. The mesocosm studies in Tvärminne and Espegrend were funded by BMBF projects SOPRAN Phase II (FKZ 03F0611) and BIOACID II (FKZ 03F06550). MSc Johanna Oja is thanked for laboratory assistance in the Tvärminne experiment.

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- Table 1: Estimates (with 95% c.i.) of the intercept (N_0) and the slope (r) of the line 1
- 2 $N=N_0+rP$ separating the N- and P-limited regions as illustrated in Fig. 1. *s* represents the
- steepness of transition from N- to P-limitation, perpendicular to the line. R²-values are
- 3 4 for the fitted response surfaces.
- 5 6

	N₀ (μM-N)	r (µМ-N:µМ-Р)	<i>s</i> (μM ⁻¹)	R ²				
Tvärminne (Baltic Sea): (incubation time 4 days)								
Fjord	0.79 (0.75, 0.83)	2.5 (2.2, 2.8)	29 (23, 35)	0.674				
M1	0.34 (0.30, 0.38)	2.9 (2.7, 3.2)	43 (31, 56)	0.622				
M3	0.76 (0.69, 0.83)	2.2 (1.9, 2.5)	19 (15, 24)	0.664				
M6	0.64 (0.56, 0.72)	2.2 (1.9, 2.6)	19 (13, 25)	0.569				
M7	0.70 (0.66, 0.75)	1.6 (1.4, 1.8)	20 (16, 25)	0.635				
Mesocosm Raunefjorden								
Incubation time								
(days):								
1	-0.41 (-0.60, -0.22)	40 (36, 44)	64 (52,76)	0.965				
2	-0.63 (-1.02, -0.25)	34 (28, 40)	50 (31, 70)	0.764				
3	-0.21 (-0.47, 0.06)	31 (27, 36)	48 (33, 63)	0.843				
4	-0.13 (-0.34, 0.08)	31 (27, 34)	47 (35, 59)	0.933				
4.5	-0.13 (-0.30, 0.04)	30 (27, 33)	51 (40, 63)	0.951				
5	0.42 (0.15, 0.69)	26 (23, 29)	36 (25, 47)	0.907				
Kviturspollen								
2	-0.09 (-0.26, 0.08)	9.9 (8.5, 11.4)	25 (17, 33)	0.940				
3	0.58 (0.46, 0.70)	12 (11, 13)	20 (15, 25)	0.972				
4	1.83 (0.72, 2.95)	14 (13, 16)	15 (8, 22)	0.946				
5	1.65 (1.32, 1.99)	15 (14, 17)	25 (20, 30)	0.963				
7	2.84 (0.10, 5.59)	14 (11, 16)	18 (9, 27)	0.855				

7





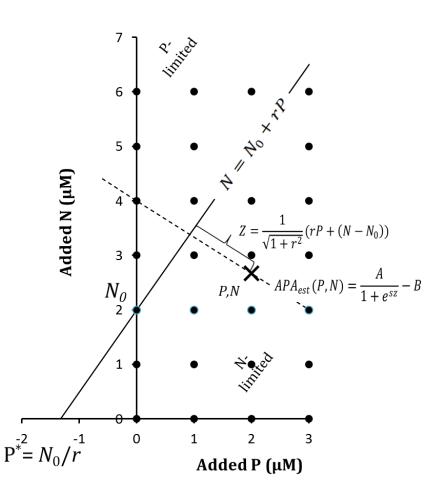
Table 2. Linear regressions of parameter estimates against incubation time for the
 Espegrend samples (see Fig. 3). Extrapolation to zero time is given (Day 0).

Espegrend	samples (see Fig. 3). Ext	trapolation to zero	time is given (Day	0).			
	Intercept	Slope	р	R ²			
	(Day 0)	of linear	(H₀: slope≠0)				
		regression					
MR							
N_0	-0.8 (-1.4,-0.2)	0.19 (0.01,0.37)	0.05	0.674			
r	41 (36,46)	-2.8 (-4.1,-1.5)	0.004	0.896			
S	65 (48,81)	-4.7 (-9.4,0.0)	0.05	0.656			
Kviturspollen							
N_0	-1.0 (-2.4,0.4)	0.6 (0.3,0.9)	0.01	0.917			
r	10 (3,16)	0.8 (-0.7,2.3)	0.2	0.479			
S	24 (6,42)	-0.8 (-4.7,3.1)	0.6	0.121			

3 H₀ null hypothesis









3

4 5

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7 Figure 1. Illustration of the fitting algorithm used. With APA measured over a (here) 4x7 matrix of 8 combinations in additions of P and N (black circles), the objective is to find the line that splits the 9 P,N-plane in an upper P-limited region with high APA and a lower N-limited region with low APA. This is done by least square fitting of the surface $APA_{est} = \frac{A}{1+e^{sZ}} - B$ to the APA-values measured 10 in each grid point. APA_{est}(P,N) is a sigmoidal function of the distance $Z = \frac{1}{\sqrt{1+r^2}}(rP + (N - N_0))$ 11 12 from the point P,N (marked X) to the line. The situation illustrated represents an N-limited system with the positive N-axis intercept (N_0) and excess-P (P^*) represented by the negative P-axis 13 14 intercept N_0/r .

15



13



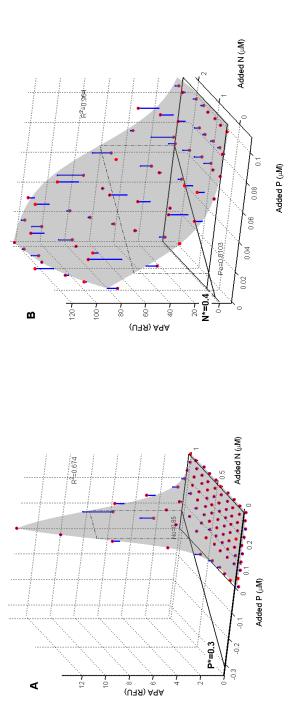


Figure 2. Measured APA values (red), fitted response surface (grey) and residuals (blue) for assays: (A) Fjord from Tvärminne and (B) MR from Espegrend (Day 1); illustrating situations interpreted as N-limited with $P^*=0.3 \mu$ M-P and a P-limited with $N^*=0.4 \mu$ M-N, respectively.

0 m 4 u





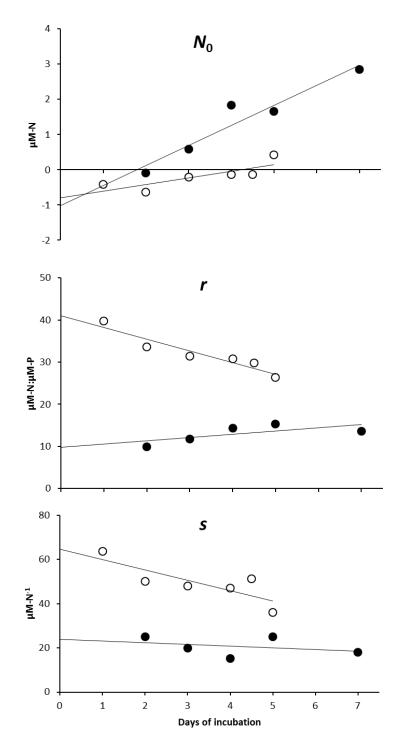
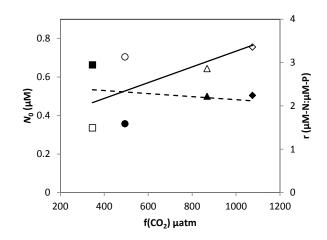


Figure 3. Change in parameter estimates with incubation time for the two samples from western

Norway. Kviturspollen has filled symbols, mesocosm MR has open symbols.







1 2

Figure 4. Scatterplots between $f(CO_2)$ and estimates of N_0 (open symbols, solid regression line) and

3 r (closed symbols, dotted regression line) for Tvärminne mesocosms M1 (squares), M3 (diamonds),

4 M6 (triangles) and M7 (circles). Regression slopes are not significant (p=0.27 and 0.79, 5 respectively).