

Supplement A. Environmental data of the sampling stations.

Table A1. Environmental data for the sampling stations. Values presented are the integrated average over the euphotic zone.

Cruise	Station	Temperature (°C)	Salinity (psu)	PAR ($\mu\text{E}/\text{m}^2/\text{s}$)	Nitrite (μM)	Nitrate (μM)	Phosphate (μM)	Silicate (μM)	Chlorophyll <i>a</i> (mg/m^3)
May 2010	St. 1	18.97	28.34	5.50	1.05	7.00	0.45	14.70	0.66
May 2010	St. 5	21.82	33.29	762.00	0.56	1.20	0.26	2.60	0.68
May 2010	St. 9	23.11	34.56	813.50	0.15	5.20	0.43	6.50	0.56
Dec. 2010	St. 1	17.42	29.88	14.10	0.23	14.80	0.69	22.80	0.69
Dec. 2010	St. 5	20.50	33.06	0.10	0.16	4.90	0.37	10.40	0.51
Dec. 2010	St. 7	20.48	34.09	0.01	0.15	4.00	0.47	9.70	0.96
Dec. 2010	St. 9	24.71	34.56	163.30	0.09	3.60	0.40	8.60	0.40
Jun. 2011	St. 1	22.87	33.87	5.39	0.83	2.52	0.41	6.31	1.52
Jun. 2011	St. 2	23.24	34.08	24.40	0.74	2.08	0.39	3.94	1.12
Jun. 2011	St. 5	24.33	34.21	82.84	0.14	0.23	0.08	1.85	0.47
Aug. 2011	St. 1	24.52	34.08	6.23	0.90	1.58	0.33	7.76	1.29
Aug. 2011	St. 3	25.19	33.96	75.97	0.86	1.17	0.25	5.62	0.82
Aug. 2011	St. 5	26.95	33.77	5.62	0.28	0.52	0.12	6.24	0.23
Aug. 2011	St. 11	22.38	34.41	15.71	0.14	3.24	0.30	6.46	0.42

Jul. 2011	St. E_1	22.43	34.24	5.71	0.18	9.13	0.33	7.55	0.38
Jul. 2011	St. E_12	23.01	34.21	16.00	0.03	2.97	0.18	6.04	0.26
Jul. 2011	St. E_19A	19.15	30.52	12.41	0.46	22.34	0.85	27.15	1.49
Jul. 2011	St. E_29	21.16	32.96	0.04	0.24	10.17	0.51	15.44	1.53
Jul. 2011	St. E_30	20.48	33.47	0.03	0.28	12.71	0.97	20.13	2.55
Sep. 2011	St. E_1	20.63	34.46	11.95	0.39	0.08	4.33	0.42	7.83
Sep. 2011	St. E_24	22.45	34.19	0.01	0.79	0.09	3.32	0.47	7.77
Sep. 2011	St. E_29	20.73	33.14	0.01	1.52	0.47	14.30	1.44	32.05
Sep. 2011	St. E_30	21.95	30.85	1.58	11.56	0.99	13.51	0.98	18.05
Oct. 2011	St. 1	23.39	29.29	14.50	0.40	24.07	0.82	22.74	0.26
Oct. 2011	St. 7	21.02	34.26	0.01	0.15	5.05	0.41	8.85	NA
Oct. 2011	St. 9	20.04	34.41	32.72	0.06	5.30	0.30	8.43	0.27

Supplement B. Biological data of the sampling stations.

Table B1. Summary statistics for the dilution experiments and Normalized Biomass Size Spectra (NBSS).

Cruise	Station	Average particles processed in T_0	Particle density (ind./ml)	Biomass ($\mu\text{g/L}$)	Average p-value of dilution exp. across size classes	NBSS slope	r^2 of NSSS
May 2010	St. 1	6168	2694	188.539	0.402	-0.727	0.854*
May 2010	St. 5	7423	3448	192.977	0.293	-0.664	0.865*
May 2010	St. 9	5742	1985	199.861	0.082	-1.064	0.785*
Dec. 2010	St. 1	6563	2248	349.784	0.462	-1.470	0.964*
Dec. 2010	St. 5	2868	1071	161.569	0.264	-1.233	0.922*
Dec. 2010	St. 7	670	253	317.033	0.179	-0.635	0.857*
Dec. 2010	St. 9	690	266	161.564	0.150	-0.785	0.888*
Jun. 2011	St. 1	2771	735	1091.136	0.177	-0.754	0.965*
Jun. 2011	St. 2	14286	587	815.715	0.248	-0.570	0.665*
Jun. 2011	St. 5	443	130	763.249	0.074	-0.458	0.882*
Aug. 2011	St. 1	1669	513	1417.715	0.118	-0.673	0.953*
Aug. 2011	St. 3	1381	369	2252.591	0.085	-0.497	0.923*
Aug. 2011	St. 5	255	76	390.693	0.167	-0.533	0.846*
Aug. 2011	St. 11	468	169	76.721	0.128	-0.834	0.943*

Jul. 2011	St. E_1	220	65	75.035	0.205	-0.659	0.912*
Jul. 2011	St. E_12	71	28	27.098	0.290	-0.540	0.813*
Jul. 2011	St. E_19A	1364	547	384.590	0.183	-0.880	0.944*
Jul. 2011	St. E_29	922	369	349.928	0.068	-0.851	0.893*
Jul. 2011	St. E_30	9302	2939	4334.308	0.521	-0.813	0.882*
Sep. 2011	St. E_1	296	111	68.792	0.195	-0.644	0.913*
Sep. 2011	St. E_24	171	61	72.329	0.169	-0.675	0.860*
Sep. 2011	St. E_29	1299	450	299.164	0.460	-0.798	0.905*
Sep. 2011	St. E_30	3513	1243	281.763	0.089	-1.150	0.895*
Oct. 2011	St. 1	3520	1205	279.075	0.330	-1.190	0.874*
Oct. 2011	St. 7	946	342	105.449	0.224	-1.099	0.962*
Oct. 2011	St. 9	1206	443	151.017	0.373	-1.183	0.995*

*indicates significant NBSS slope

Table B2. Size-specific growth rate and grazing mortality scaling on body size for each sampling station.

Cruise	Station	μ scaling slope	SE of μ scaling slope	p-value of μ scaling slope	m scaling slope	SE of m scaling slope	p-value of m scaling slope
May 2010	St. 1	0.112	0.053	0.077	0.247	0.150	0.151
May 2010	St. 5	0.057	0.123	0.667	0.010	0.266	0.971
May 2010	St. 9	0.379	0.091	0.006	0.092	0.036	0.042
Dec. 2010	St. 1	0.180	0.019	<0.001	0.298	0.133	0.066
Dec. 2010	St. 5	0.129	0.045	0.036	-0.199	0.351	0.596
Dec. 2010	St. 7	0.076	0.046	0.159	-0.347	0.185	0.119
Dec. 2010	St. 9	0.206	0.066	0.014	0.116	0.060	0.089
Jun. 2011	St. 1	-0.023	0.030	0.461	0.135	0.067	0.081
Jun. 2011	St. 2	-0.054	0.009	<0.001	0.000	0.117	0.997
Jun. 2011	St. 5	0.014	0.021	0.525	0.094	0.060	0.170
Aug. 2011	St. 1	-0.014	0.020	0.485	-0.036	0.040	0.393
Aug. 2011	St. 3	-0.026	0.022	0.267	-0.116	0.064	0.100
Aug. 2011	St. 5	-0.344	0.209	0.200	-0.223	0.167	0.275
Aug. 2011	St. 11	0.196	0.048	0.010	0.027	0.026	0.344
Jul. 2011	St. E_1	0.070	0.169	0.702	0.187	0.113	0.175
Jul. 2011	St. E_12	0.082	0.015	0.005	-0.118	0.049	0.073

Jul. 2011	St. E_19A	0.085	0.052	0.151	0.076	0.141	0.611
Jul. 2011	St. E_29	0.086	0.057	0.182	0.003	0.051	0.961
Jul. 2011	St. E_30	-0.010	0.067	0.894	0.569	0.577	0.380
Sep. 2011	St. E_1	-0.056	0.084	0.534	0.147	0.107	0.220
Sep. 2011	St. E_24	0.079	0.055	0.193	0.244	0.087	0.026
Sep. 2011	St. E_29	0.352	0.144	0.092	0.595	0.355	0.192
Sep. 2011	St. E_30	0.415	0.130	0.085	0.526	0.111	0.041
Oct. 2011	St. 1	0.388	0.030	<0.001	0.223	0.129	0.145
Oct. 2011	St. 7	0.267	0.032	<0.001	0.415	0.081	0.001
Oct. 2011	St. 9	0.021	0.101	0.852	0.370	0.089	0.026

Supplement C. Results of analysing the six reduced data sets.

Table C1. Results of analyzing the six reduced data sets showing the general scaling relationship of size specific growth rate (μ') and grazing mortality (m) versus body size. In addition, the coupling of grazing mortality and growth rate is also examined. All the data sets show qualitatively consistent results as the whole data set. Reduced data set 1: removing size classes with the regression p-value larger than 0.25 in dilution experiments; Reduced data set 2: removing stations with the regression p-value for the whole community larger than 0.25; Reduced data set 3: first removing the stations with the regression p-value for the whole community larger than 0.25 and then removing the size classes with the regression p-value larger than 0.25 in the remaining stations; Reduced data set 4: removing the stations with average regression p-value of all size classes larger than 0.25; W/O "colony small cells": removing colony small cells from calculation; W/O "chain-forming diatom": removing chain-forming diatom from calculation.

Reduced data set 1	Estimate	SE	p-value	Reduced data set 2	Estimate	SE	p-value
μ - size GLMM	0.094	0.015	<0.001	μ - size GLMM	0.094	0.017	<0.001
m - size GLMM	0.137	0.018	<0.001	m - size GLMM	0.124	0.026	<0.001
$m - \mu'$ GLMM	0.777	0.086	<0.001	$m - \mu'$ GLMM	0.865	0.095	<0.001
residuals of m - size GLMM~ residuals of μ' - size GLMM	0.441	0.065	<0.001	residuals of m - size GLMM~ residuals of μ' - size GLMM	0.359	0.045	<0.001
Reduced data set 3	Estimate	SE	p-value	Reduced data set 4	Estimate	SE	p-value

μ - size GLMM	0.090	0.017	<0.001	μ - size GLMM	0.078	0.018	<0.001
m - size GLMM	0.138	0.019	<0.001	m - size GLMM	0.091	0.025	<0.001
$m - \mu'$ GLMM	0.778	0.092	<0.001	$m - \mu'$ GLMM	0.798	0.098	<0.001
residuals of m - size GLMM~ residuals of μ' - size GLMM	0.480	0.073	<0.001	residuals of m - size GLMM~ residuals of μ' - size GLMM	0.382	0.050	<0.001
W/O "colony small cells"	Estimate	SE	p-value	W/O "chain-forming diatom"	Estimate	SE	p-value
μ - size GLMM	0.049	0.017	0.004	μ - size GLMM	0.082	0.016	<0.001
m - size GLMM	0.101	0.031	0.001	m - size GLMM	0.122	0.030	<0.001
$m - \mu'$ GLMM	0.518	0.118	<0.001	$m - \mu'$ GLMM	0.792	0.117	<0.001
residuals of m - size GLMM~ residuals of μ' - size GLMM	0.614	0.093	<0.001	residuals of m - size GLMM~ residuals of μ' - size GLMM	0.658	0.008	<0.001

Table C2. Results of univariate model analyses for the six reduced data sets. The results here are qualitatively the same as the results of the whole data set.

	Independent variables	Biological anticipation	Reduced data set 1		Reduced data set 2		Reduced data set 3		Reduced data set 4	
			Coefficient	p-value	Coefficient	p-value	Coefficient	p-value	Coefficient	p-value
Model 1	m_S	+	0.024	0.654	0.197	0.337	0.193	0.611	-0.006	0.515
Model 2	m_L	-	-0.073	0.082	-0.019	0.492	-0.045	0.090	-0.028	0.163
Model 3	μ_S'	-	0.163	0.997	0.342	0.990	0.245	0.924	0.196	0.928
Model 4	μ_L'	+	-0.090	0.929	-0.049	0.950	-0.077	0.995	-0.062	0.999
Model 5	I_S'	+	-0.057	0.909	-0.302	0.951	-0.257	0.904	-0.249	0.910
Model 6	I_L'	-	-0.051	0.367	0.016	0.552	-0.047	0.748	0.023	0.598
Model 7	m_S/m_L	+	0.085	0.035*	0.403	0.027*	0.415	0.061	0.277	0.058
Model 8	μ_S'/μ_L'	-	0.149	1.000	0.698	0.999	0.745	0.999	0.556	0.999
Model 9	I_S'/I_L'	+	-0.028	0.736	-0.072	0.912	-0.097	0.866	-0.022	0.745

Table C2. (Continued)

	Independent variables	Biological anticipation	Data set W/O "colony small cells" category		Data set W/O "chain-forming diatom" category	
			Coefficient	p-value	Coefficient	p-value
Model 1	m_S	+	0.139	0.444	0.141	0.394
Model 2	m_L	-	-0.024	0.436	-0.028	0.194
Model 3	μ_S'	-	-0.021	0.774	0.226	0.984
Model 4	μ_L'	+	0.032	0.242	-0.040	0.963
Model 5	I_S'	+	-0.103	0.730	-0.251	0.940
Model 6	I_L'	-	-0.015	0.902	-0.187	0.862
Model 7	m_S/m_L	+	0.303	0.044*	0.321	0.023*
Model 8	μ_S'/μ_L'	-	0.404	0.967	0.587	0.999
Model 9	I_S'/I_L'	+	0.004	0.939	-0.013	0.592

* indicates the model that gives biologically reasonable and significant result.

Supplement D. Relationship between NBSS slope and relative grazing mortality of small versus large phytoplankton (m_S/m_L)

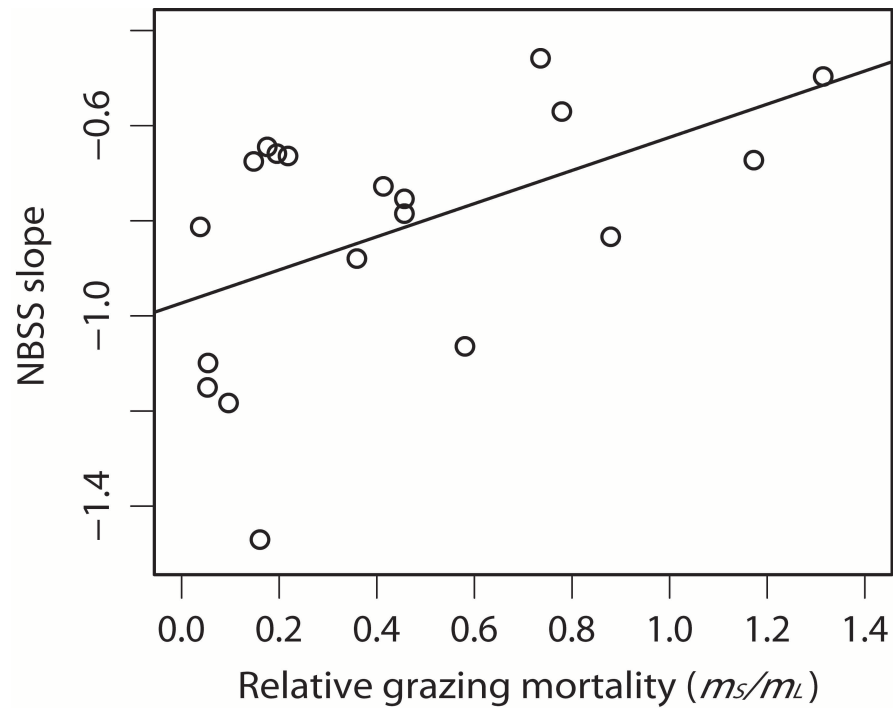


Figure D1. Scatter plot showing the relationship between NBSS slope and relative grazing mortality of small versus large microphytoplankton (m_S/m_L). The solid line indicates the significant regression line (slope=0.348; $p=0.026$)