



Carbon and nitrogen fluxes in the marine coccolithophore
Emiliana huxleyi grown under different nitrate
concentrations

by

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List of contents

Publications	4
Abstract	5
List of selected abbreviations	7
List of figures	10
List of tables	15
1. Introduction	17
1.1 The biogeochemical role of <i>Emiliana huxleyi</i> in the ocean	17
1.2 Nitrogen-nitrate in the ocean	20
1.3 Nitrate assimilation	21
1.4 Carbon skeletons for amino acid synthesis	25
1.5 Inorganic carbon assimilation	28
1.6 Photosynthesis and respiration.....	29
1.7 Effects of nitrogen limitation on <i>Emiliana huxleyi</i>	30
1.8 Objectives of the study.....	32
2. Material and methods	33
2.1 Culture conditions.....	33
2.2 Carbonate chemistry	33

2.3 Cellular carbon and nitrogen.....	34
2.4 Macromolecular composition	34
2.5 Coccolith morphology	35
2.6 Enzyme extraction	36
2.7 Nitrate reductase	36
2.8 Nitrite reductase	37
2.9 Glutamine synthetase	37
2.10 Glutamate synthase	39
2.11 Phosphoenolpyruvate carboxykinase	39
2.12 Purification of glutamine synthetase isoforms	40
2.13 Photosynthesis and inorganic carbon fluxes	41
2.14 Statistical analysis	42
3. Results	43
3.1 Carbonate chemistry	43
3.2 Growth parameters	44
3.4 Macromolecular composition	48
3.5 Nitrate assimilation	50
3.6 Glutamine synthetase isoforms	54
3.7 Photosynthesis and respiration	62
3.8 Inorganic carbon fluxes	65

4. Discussion	68
4.1 Growth parameters	68
4.2 Macromolecular pools	69
4.3 Nitrate assimilation	70
4.4 Photosynthesis and respiration.....	72
4.5 Inorganic carbon uptake.....	73
4.6 Intracellular concentrations of nitrate and carbon dioxide.....	74
4.7 Conclusions.....	77
6. Literature cited	78
7. Acknowledgements	91

Publications

Publications

- Kaffes, A, Thoms S, Trimborn S, Rost B, Langer G, Richter K-U, Köhler A, Norici A, Giordano M. Carbon and nitrogen fluxes in the marine coccolithophore *Emiliana huxleyi* grown under different nitrate conditions, in preparation

Poster presentations

- Kaffes, A., Trimborn, S., Rost, B., Langer, G., Richter, K.-U., Norici, A., Giordano, M., Thoms, S. (2009). Effects of NO_3^- availability on carbon and nitrogen assimilation in the marine coccolithophore *Emiliana huxleyi*, 9th International Phycological Congress, Tokyo 2009, Japan.

Abstract

Coccolithophores are important primary producers and the most prolific calcifiers in the ocean. Since carbon (C) and nitrogen (N) assimilation are tightly connected, N availability has a crucial impact on photosynthesis and calcification, and therefore affects the biogeochemical function of coccolithophores in the sea. In general, N availability limits primary production in many areas in the ocean. Furthermore, N availability is expected to decrease in the near future, due to increased thermal stratification in the surface ocean and reduced nutrient input from deeper waters. Keeping this in mind, we examined the effects of NO_3^- availability on C and N assimilation in the marine coccolithophore *Emiliana huxleyi* under saturating light and present day pCO_2 . Although growth rates and C to N ratios were similar in low and high NO_3^- -grown cells, cellular organic C and N, and also inorganic C decreased under low NO_3^- . Growth at low NO_3^- caused a higher proportion of fixed C to be allocated to lipids relative to carbohydrates and proteins. Low NO_3^- availability led to downregulation of nitrate reductase (NR), nitrite reductase (NiR) and phosphoenolpyruvate carboxykinase (PEPCK) maximal activities ($V_{\max}$), up-regulation of glutamine synthetase (GS) and glutamate synthase (GOGAT) $V_{\max}$, and down-regulation of net photosynthesis and HCO_3^- uptake. The apparent Chl α specific efficiency of light harvesting (α) and CO_2 uptake remained similar, irrespective of the NO_3^- concentration in the culture medium. The affinity of NR for NO_3^- , NiR for NO_2^- and GS for Glu, as well as the affinities of inorganic C uptake were higher under low NO_3^- . This study suggests that maintenance of a constant C to N ratio under low and high NO_3^- requires a concerted regulation of the intracellular CO_2 and NO_3^- concentrations to equilibrate the fluxes through the C and N assimilation pathway.

List of selected abbreviations

α_{chl}	Apparent Chl <i>a</i> specific efficiency of light harvesting
Acetyl-CoA	Acetyl Coenzyme A
ADP	Adenosine phosphate
CA	Carbonic anhydrase
CCM	Carbon concentrating mechanism
Chl	Chlorophyll
Ci	Inorganic carbon
DIC	Dissolved inorganic carbon
EDTA	Ethylenediaminetetraacetic acid
Fer _{red}	Reduced ferredoxin
FTIR	Fourier transform infrared spectroscopy
Glu	Glutamate
Gln	Glutamine
GS	Glutamine synthetase
Gly	Glycine
GOGAT	Glutamate synthase

HEPES	4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid
IR	Infrared
ICDH	Isocitrate dehydrogenase
kDa	Kilo Dalton
K _m	Michaelis-Menten constant
LD	Light-dark
MDH	Malate dehydrogenase
mRNA	Messenger ribonucleic acid
NADPH	Nicotinamide adenine dinucleotide phosphate
NADH	Nicotinamide adenine dinucleotide
NiR	Nitrite reductase
NiT	Nitrite transporter
NR	Nitrate reductase
NT	Nitrate transporter
OAA	Oxalacetate
OG	Oxoglutarate
PDC	Pyruvate dehydrogenase complex
PEP	Phosphoenolpyruvate

PEPC	Phosphoenolpyruvate carboxylase
PEPCK	Phosphoenolpyruvate carboxykinase
PGA	Phosphoglycerate
PIC	Particulate inorganic carbon
POC	Particulate organic carbon
PON	Particulate organic nitrogen
RuBP	Ribulose biphosphate
RubisCO	Ribulose biphosphate carboxylase/oxygenase
Ser	Serine
TA	Total alkalinity
TCA	Tricarboxylic acid
TPC	Total particulate carbon
TrioseP	Triose phosphate
V _{max}	Maximal activity

List of figures

Fig. 1 The biological C pumps: Photosynthetic production of organic matter in the surface layer and its subsequent transport to depth (organic C pump), generates a CO₂ sink in the ocean. In contrast, calcium carbonate production and its transport to depth (carbonate counter pump), releases CO₂ in the surface layer. The relative strengths of these two processes (rain ratio) largely determine the biologically-mediated ocean atmosphere CO₂ exchange (in Rost and Riebesell, 2004). 18

Fig. 2 Seawater pH and the dissolved carbon dioxide (CO₂) and carbonate ion (CO₃²⁻) concentrations in the surface layer of the ocean assuming a “business as usual” (IS92a) anthropogenic CO₂ emission scenario (Houghton et al., 1995). Dashed lines represent the predicted changes in carbonate chemistry if CO₂ emissions are reduced according to the Kyoto Protocol (in Rost and Riebesell, 2004; modified after Wolf-Gladrow et al., 1999). 19

Fig. 3 The progressive increase in atmospheric CO₂ affects the marine biota indirectly through rising mean global temperatures, thereby causing increased surface ocean stratification. This in turn reduces the nutrient input from deeper layers and increases the light availability due to shoaling of the upper mixed layer (in Rost and Riebesell, 2004). 19

Fig. 4 The nitrate (NO₃⁻) assimilation pathway including NO₃⁻ reduction to NH₄⁺ and incorporation of NH₄⁺ into C skeletons in the chloroplast. NT: nitrate transporter; NR: nitrate reductase; NiT: nitrite transporter; NiR: nitrite reductase; NH₄⁺: ammonium; GS: glutamine synthetase; GOGAT: glutamate synthase ATP: adenosine triphosphate; NADH: nicotinamide

adenine dinucleotide; red Fer: reduced Ferredoxin; Glu: glutamate; Gln: glutamine and 2OG: 2 oxoglutarate. 23

Fig. 5 Cytosolic ammonium assimilation. NH_4^+ : ammonium; GS: glutamine synthetase; ATP: adenine triphosphate; NADH: nicotinamine adenine dinucleotide; Glu: glutamate; Gln: glutamine; Rubisco: ribulose bisphosphate carboxylase/oxygenase; RuBP: ribulose bisphosphate; Gly: glycine; Ser: serin; PGA: phosphoglycerate; TrioseP: triose phosphate. . 26

Fig. 6 The GS/GOGAT cycle produces glutamine and glutamate which are the starting compounds for the synthesis of all remaining amino acids, nucleic acids and chlorophylls (Lea et al., 1993). 27

Fig. 7 The C skeletons needed for amino acid synthesis are mainly provided via the TCA cycle in form of 2-oxoglutarate (2OG). PEPCK: phosphoenolpyruvate carboxykinase; PEPC: phosphoenolpyruvate carboxylase; PKc: pyruvate kinase C; PDC: pyruvate dehydrogenase complex; ICDH: isocitrate dehydrogenase; Glu: glutamate; Gly: glycine; TrioseP: triose phosphate; OAA: oxalacetate; AcetylCoA: acetyl coenzyme A, PEP: phosphoenolpyruvate. 27

Fig. 8 Carbon concentrating mechanisms (CCMs) in eukaryotic algal cells. Bicarbonate (HCO_3^-) is taken up by specific HCO_3^- transporters, whereas CO_2 is mainly taken up by diffusion. An active CO_2 system seems to exist, but it is poorly described till now. The enzyme carbonic anhydrase (CA) catalyzes the conversion between CO_2 and HCO_3^- . Various isoenzymes of CA have been described, which are located in the chloroplast stroma, the thylakoid membrane, the cytosol, the mitochondrion (not shown here) and the extracellular space. RubisCO: ribulose bisphosphate carboxylase/oxygenase. 29

- Fig. 9 Effects of high (■) and low (□) NO_3^- on particulate organic carbon (POC) production (A), particulate organic nitrogen (PON) production (B) and particulate inorganic carbon (PIC) production (C) in *E. huxleyi* cells . Values represent the means of triplicate incubations ($\pm$ SD).. 46
- Fig. 10 Scanning electron microscopy images of coccoliths of *E. huxleyi*. All scalebars 2 μm . All specimens in distal view. (a) normal coccolith. (b) incomplete coccolith. (c) malformed coccolith. (d) incomplete and malformed coccolith)... 47
- Fig. 11 Percentage of normal, malformed, incomplete, and incomplete and malformed (inc/malf) coccoliths under high (■) and low (□) NO_3^- . Values represent an average of duplicates ($\pm$ SD)..... 48
- Fig. 12 The ratios of proteins to lipids (A), proteins to carbohydrates (B) and lipids to carbohydrates (C) in cells under high (■) and low (□) NO_3^- . Values represent an average of triplicates ($\pm$ SD).... 49
- Fig. 13 Kinetic curve of nitrate reductase (NR) for NO_3^- (A) and NADH (B) in cells grown under high (■) and low (□) NO_3^- . All values represent the means of triplicates..... 51
- Fig. 14 Kinetic curve of nitrite reductase (NiR) for NO_2^- in cells grown under high (■) and low (□) NO_3^- . All values represent the means of triplicates..... 52

- Fig. 15 Kinetic curve of GS for glutamate (Glu) in cells grown under high (■) and low (□) NO_3^- . One unit is equal to 1 fmol L-Glu γ -monohydroxamate. All values represent the means of triplicates..... 54
- Fig. 16 Kinetic curve of GS for ATP (A) and NH_2OH (B) in cells grown under high (■) and low (□) NO_3^- . One unit is equal to 1 fmol L-Glu γ -monohydroxamate. In the kinetics of GS for ATP (n=3) and for NH_2OH (n=4).... 55
- Fig. 17 Purification of the GS isoforms in cells grown under high (■) (A) and low (□) (B) NO_3^- using a Sephacryl S-200 HR column. GS active fractions are shown with arrows... 56
- Fig. 18 Relative (A) and specific (B) activities of the GS_1 and GS_2 isoforms in high (■) and low (□) NO_3^- -grown cells after separation on the Sephacryl S-200 HR column. GS_1 : cytosolic isoform; GS_2 : chloroplastic isoform..... 57
- Fig. 19 Purification of the GS_2 isoform in cells grown under high (■) (A) and low (□) (B) NO_3^- using a MonoQ column. GS active fractions are shown with arrows..... 58
- Fig. 20 Purification of the GS_1 isoform in cells grown under high (■) (A) and low (□) (B) NO_3^- using a MonoQ column. GS active fractions are shown with arrows. The GS_1 isoform could not be further purified in the MonoQ column under low NO_3^- 59

Fig. 21 An aliquot of the glutamine synthetase (GS) active fractions after separation on a MonoQ 5/50 GL column was applied on a 10 % SDS-PAGE gel. A band at approximately 45 kDa was visualized after silver staining (Std: protein standard – ovalalbumin: 45 kDa)..... 61

Fig. 22 Kinetic curve of net photosynthesis for CO₂ (A) and DIC (B) in cells grown under high (■) and low (□) NO₃⁻. For net photosynthesis in cells grown under high (H) NO₃⁻ (n=5). All other values represent the means of triplicates..... 64

Fig. 23 The effect of various light intensities (PFD) on net photosynthesis in cells grown under high (■) and low (□) NO₃⁻. All values represent the means of triplicates..... 65

Fig. 24 Kinetics of HCO₃⁻ uptake in cells grown under high (■) and low (□) NO₃⁻. For HCO₃⁻ uptake in cells grown under high (H) NO₃⁻ (n=5). All other values represent the means of triplicates..... 66

Fig. 25 Kinetics of gross CO₂ (A) and net CO₂ (B) uptake in cells grown under high (■) and low (□) NO₃⁻. For gross and net CO₂ uptake in cells grown under high (H) NO₃⁻ (n=5). All other values represent the means of triplicates..... 67

List of tables

Tab. I Major nitrogen (N) species in the sea. Approximate average values are given for oceanic waters and appropriate ranges are given for coastal and estuarine waters. All values are listed in $\mu\text{g atom N L}^{-1}$ (in Sharp, 1983)..... 20

Tab. II The energetics of CO_2 and N assimilation (simplified). Table I outlines the ATP and reductant requirements of the various processes associated with photosynthetic C fixation and the assimilation of N into protein (in Turpin, 1991).The equations are not balanced.. 31

Tab. III Carbonate chemistry in the culture medium of the high (H) and low (L) NO_3^- treatments.TA: total alkalinity; DIC: dissolved inorganic carbon; pCO_2 : CO_2 partial pressure.....43

Tab. IV Growth rate (μ), cell volume (CV), particulate organic carbon (POC), particulate inorganic carbon (PIC), particulate organic nitrogen (PON), protein content and C to N ratio in cells grown under high (H) and low (L) NO_3^- . The values represent the means of triplicate incubations ($\pm$ SD). 45

Tab. V Kinetics V_{max} and K_m of NR and NiR in cells grown under high (H) and low (L) NO_3^- . All values represent the means of triplicates ($\pm$ SD)..... 50

.

Tab. VI Kinetics $V_{\max}$ and K_m of GS, as well as the activities ($V_{\max}$) of GOGAT and PEPCK in cells grown under high (H) and low (L) NO_3^- . All values represent the means of triplicates ($\pm$ SD). One unit is equal to 1 fmol L-Glu γ -monohydroxamate... 53

Tab. VII Purification of GS isoforms in *E. huxleyi* cells grown under high (H) and low (L) NO_3^- (GS₁: cytosolic GS isoform; GS₂: chloroplastic GS isoform). GS activity was measured according to the GS synthetase assay. Relative GS activity is given in nmol L-Glu γ -monohydroxamate $\text{min}^{-1} \text{mL}^{-1}$. Specific GS activity is given in nmol L-Glu γ -monohydroxamate $\text{min}^{-1} \text{mg protein}^{-1}$ 60

Tab. VIII Kinetics $V_{\max}$ and $K_{1/2}$ of net photosynthesis, apparent Chl α specific efficiency of light harvesting (α), kinetics of gross CO_2 uptake, net CO_2 uptake and HCO_3^- uptake, as well as respiration in cells grown under high (H) and low (L) NO_3^- . For net photosynthesis, gross and net CO_2 uptake, HCO_3^- uptake and respiration in cells grown under high (H) NO_3^- (n=5). All other values represent the means of triplicates ($\pm$ SD)..... 63

1. Introduction

1.1. The biogeochemical role of *Emiliana huxleyi* in the ocean

The marine coccolithophore *Emiliana huxleyi*, which is distributed worldwide except for the polar regions (Winter et al., 1994), is considered to be one of the major producers of calcite in the ocean (Baumann et al., 2004). Coccolithophores influence the carbonate pump by producing calcium carbonate, but as important marine primary producers, they also affect the organic carbon (C) pump through photosynthesis. The vertical transport of a part of the organic material produced via photosynthesis causes a drawdown of CO₂ in the surface ocean (organic C pump), whereas calcification displays an opposite role in the carbonate chemistry of the surface ocean by causing a net release of CO₂ during the formation and export of calcium carbonate (carbonate counter pump) (Fig. 1). The formation of calcite in the surface ocean and its sinking to deeper layers modifies the alkalinity of the upper water layer. The *rain ratio* represents the relationship of the particulate inorganic C to the particulate organic C in exported biogenic material, and may determine the biologically mediated CO₂ exchange between the surface ocean and the overlying atmosphere (Fig. 1).

Anthropogenic emissions increase the amount of CO₂ in the atmosphere, which in turn leads to changes in the seawater carbonate chemistry of the surface ocean (Fig. 2). The concentration of CO₂ increases in the surface ocean and as a consequence the carbonate ion (CO₃²⁻) concentration and the pH decrease. An acidification of the upper ocean is already observed nowadays and is going to increase in the future (Fig. 2). Photosynthesis and calcification in *E. huxleyi* are sensitive to changes in the seawater carbonate chemistry (Rost et al., 2003; Iglesias-Rodriguez et al., 2008).

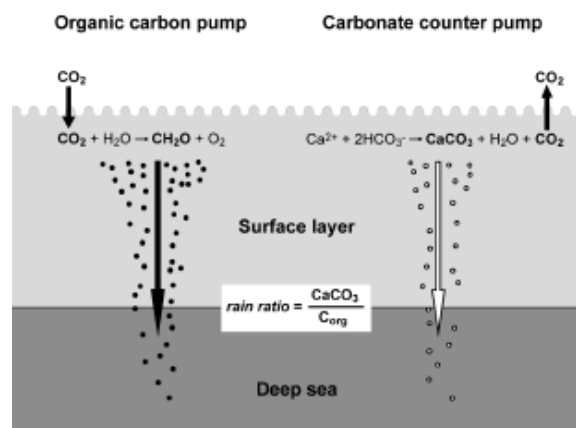


Figure 1. The biological carbon (C) pumps: Photosynthetic production of organic matter in the surface layer and its subsequent transport to depth (organic C pump), generates a CO_2 sink in the ocean. In contrast, calcium carbonate production and its transport to depth (carbonate counter pump), releases CO_2 in the surface layer. The relative strengths of these two processes (rain ratio) largely determine the biologically-mediated ocean atmosphere CO_2 exchange (in Rost and Riebesell, 2004).

The ongoing increase in atmospheric CO_2 affects the marine environment directly through changes in the seawater carbonate chemistry of the surface ocean and, indirectly through rising mean global temperatures (Intergovernmental Panel on Climate Change 2007). Increased temperatures in the surface ocean will increase thermal stratification and reduce therewith nutrient input from deeper waters, and will also have an impact on the light conditions of the upper water layer (Fig. 3).

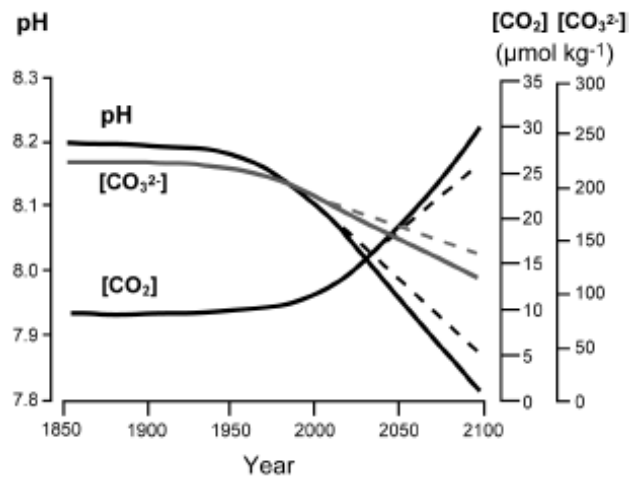


Figure 2. Seawater pH and the dissolved carbon dioxide (CO_2) and carbonate ion (CO_3^{2-}) concentrations in the surface layer of the ocean assuming a “business as usual” (IS92a) anthropogenic CO_2 emission scenario (Houghton et al., 1995). Dashed lines represent the predicted changes in carbonate chemistry if CO_2 emissions are reduced according to the Kyoto Protocol (in Rost and Riebesell, 2004; modified after Wolf-Gladrow et al., 1999).

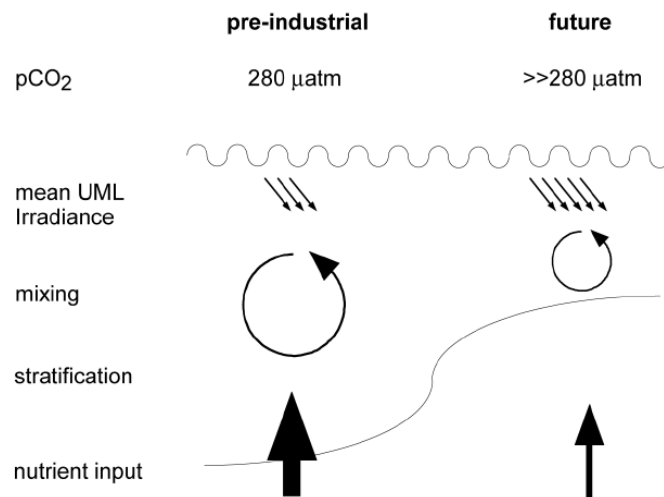


Figure 3. The progressive increase in atmospheric CO_2 affects the marine biota indirectly through rising mean global temperatures, thereby causing increased surface ocean stratification. This in turn reduces the nutrient input from deeper layers and increases the light availability due to shoaling of the upper mixed layer (in Rost and Riebesell, 2004).

1.2. Nitrogen-nitrate in the ocean

A high amount of the assimilated C in algae is devoted to N metabolism for the synthesis of proteins, nucleic acids and chlorophylls. Therefore the concentration and form of N in seawater should affect C assimilation in coccolithophores thereby altering their role in the global C cycle. Nitrogen can be found in the sea in form of inorganic and organic compounds (Sharp, 1975). Inorganic N includes molecular nitrogen (N_2), nitrate (NO_3^-), nitrite (NO_2^-), ammonium (NH_4^+), nitrous oxide (N_2O) and amine (NH_2). N_2 is by far the most abundant N compound (Tab. I), but can be utilized almost only by cyanobacteria in the ocean (Carpenter, 1975), and a small part by marine plants and their symbiotic bacteria (Capone, 1975). Besides N_2 , NO_3^- is the most abundant species of inorganic N in the sea, whereas NO_2^- and NH_4^+ are less abundant altogether, but are of local significance (Tab. I). Organic N compounds (Williams, 1975), such as urea, amino acids, nucleic acids, nucleotides and nucleic bases contain both C, and N, may be also of great importance in certain areas in the sea (Tab. I).

Table I. Major nitrogen (N) species in the sea. Approximate average values are given for oceanic waters and appropriate ranges are given for coastal and estuarine waters.

All values are listed in $\mu\text{g atom N L}^{-1}$ (in Sharp, 1983).

Species	Surface oceanic (0–100 m)	Deep oceanic (>100 m)	Coastal	Estuarine
Nitrogen gas N_2	800	1150	700-1100	700-1100
Nitrate NO_3^-	0.2	35	0-30	0-350
Nitrite NO_2^-	0.1	<0.1	0-2	0-30
Ammonium NH_4^+	<0.5	<0.1	0-25	0-600
Dissolved organic N	5	3	3-10	5-150
Particulate organic N	0.4	<0.1	0.1-2	1-100

In many areas in the sea, N limitation limits primary production (Dugdale, 1967; Howarth, 1988). As being one of the most common forms of inorganic N, NO_3^- plays a crucial role in marine primary productivity (Dugdale and Göring, 1967; Kanda, 1995). Nitrate concentrations vary in the ocean (Tab. I) with low NO_3^- concentrations ($\sim 5 \text{ nmol L}^{-1}$) in oligotrophic areas and higher NO_3^- concentrations ($\sim 30 \text{ } \mu\text{mol L}^{-1}$) in upwelling systems, as reported by Rees et al. (2006) for the Atlantic Ocean. Although, *E. huxleyi* is capable of utilizing also dissolved organic N (Ietswaart et al., 1994; Palenik and Henson, 1997), NO_3^- is an important N source for the growth of this coccolithophore (Page et al., 1999).

1.3. Nitrate assimilation

Nitrate assimilation includes NO_3^- transport into the cell, the cytosolic reduction of NO_3^- to NO_2^- , the NO_2^- transport into the chloroplast and its subsequent reduction to NH_4^+ (Fig. 4). Ammonium is then incorporated into C skeletons in the chloroplast or the cytosol (Fig. 5). Nitrate assimilation by marine phytoplankton has been studied, either by measuring the reduction in NO_3^- concentration in the culture medium (Dortch et al., 1982) or the incorporation of NO_3^- containing the isotope ^{15}N in algal biomass (Lomas and Glibert, 2000).

The transport of NO_3^- into the cell is mediated by NO_3^- transporter(s) (Nrt(s)) in the plasma membrane (Fig. 4) and it was suggested to be a key step controlling the amount of NO_3^- assimilated by cells (Galvan and Fernandez, 2001). Four different gene systems (*Nrt2.1/Nar2*, *Nrt2.2/Nar2*, *Nrt2.3*, *Nrt2.X*) are responsible for NO_3^- transport, and some of them also for NO_2^- transport, in *Chlamydomonas reinhardtii* (Quesada et al., 1994; Quesada and Fernandez, 1994; Galvan et al., 1996; Rexach et al., 1999; Fernandez and Galvan, 2007). These NO_3^- transport systems can be distinguished in high and low affinity NO_3^- transporters. Apart from one NO_3^- transporter which is constitutively expressed in *C. reinhardtii*, NO_3^- was found to

induce the expression of all other NO_3^- transport systems, whereas NH_4^+ represses them (Galvan et al., 1996; Rexach et al., 1999). High-affinity NO_3^- transporter genes (*Nrt2*) have been described in *E. huxleyi* and *Nrt2* gene transcripts were found to be induced in the presence of NO_3^- , but also under N starvation (Song and Ward, 2007).

Nitrate reductase (NR) is the first enzyme in the NO_3^- reduction pathway and is located in the cytosol (Fig. 4). NR catalyzes the two electron reduction of NO_3^- to NO_2^- . The energy for this reductive step is supplied by the oxidation of NADH/NADPH. Although characterization of the gene coding for NR is still missing in *E. huxleyi*, the molecular mass and kinetics of the purified NR enzyme have been recently described in this phytoplankton species (Iwamoto and Shiraiwa, 2003). Native *E. huxleyi* NR has an overall mass of 514 kDa and is composed of six 85 kDa homologous subunits (Iwamoto and Shiraiwa, 2003). The K_m for NADH and NO_3^- of purified NR were 40 μM and 104 μM , respectively (Iwamoto and Shiraiwa, 2003). Nitrate seems to play an important role in regulating NR gene and protein expression, as well as NR activity. Nitrate acts positive on NR mRNA and protein expression in *C. reinhardtii* (Navarro et al., 1996). Furthermore, increasing NO_3^- concentrations were found to increase NR activity in the marine diatom *Thalassiosira pseudonana* (Berges and Harrison, 1995) as well as in the green algae *Dunaliella primolecta* and *Chlorella stigmatophora* (Hipkin et al., 1983). NR seems to be of great importance in the NO_3^- assimilation pathway through its involvement in the regulation of the $\text{NO}_3^-/\text{NO}_2^-$ transport as reported for *C. reinhardtii* cells (Navarro et al., 1996).

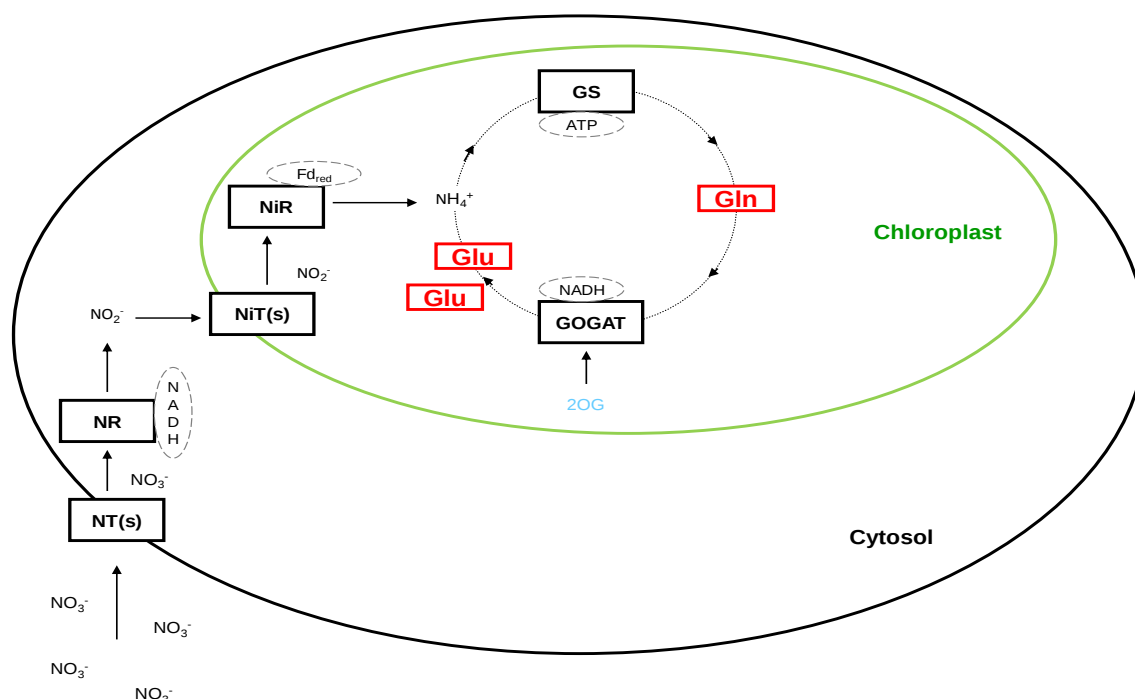


Figure 4. The nitrate (NO_3^-) assimilation pathway including NO_3^- reduction to NH_4^+ and incorporation of NH_4^+ into C skeletons in the chloroplast. NT: nitrate transporter; NR: nitrate reductase; NiT: nitrite transporter; NiR: nitrite reductase; NH_4^+ : ammonium; GS: glutamine synthetase; GOGAT: glutamate synthase ATP: adenosine triphosphate; NADH: nicotinamide adenine dinucleotide; Fer_{red} : reduced Ferredoxin; Glu: glutamate; Gln: glutamine and 2OG: 2 oxoglutarate.

The transport of NO_2^- into the chloroplast is mediated via NO_2^- transporters in the chloroplast envelope (Fig. 4). Nitrite transporters have not been studied yet in *E. huxleyi* neither at the gene nor at the protein level. Two high-affinity NO_2^- transporters have been identified in *C. reinhardtii* and their expression was found to be induced by NO_3^- and repressed by NH_4^+ (Rexach et al., 1999). Rexach et al. (2000) reported the existence of the *Nar1* gene which codes for a transmembrane protein (NAR1) involved in the transport of NO_2^- from the cytosol into the chloroplast. This protein was found to be essential for growth under NO_3^- limiting conditions and seems to control the amount of NO_3^- incorporated under limiting CO_2

conditions (Rexach et al., 2000). Galvan et al. (2002) have reported the existence of various *Nar1* genes involved in NO_2^- transport and Mariscal et al. (2006) showed that the *Nar1* gene family is differentially regulated in response to the C and N source.

Nitrite is reduced to NH_4^+ in the chloroplast and the reaction is catalyzed by the NiR which utilizes reduced ferredoxin. As in the case of NO_2^- transporters, the NiR enzyme has not been studied yet in the coccolithophore *E. huxleyi*. Maximal activity of NiR in *C. reinhardtii* was detectable only in the presence of NO_3^- and NH_4^+ was found to repress NiR activity (Galvan et al., 1991). The gene *Nii1* coding for NiR in *C. reinhardtii* was found to be located in the same gene cluster with other NO_3^- assimilating genes and it was found to be regulated in response to light and N availability (Quesada et al., 1998). The kinetics of NR and NiR as well as optimal temperature of NR and NiR activities were studied in diatoms and flagellates by Lomas and Glibert (2000). The authors suggested that differences in the affinity of NR for NO_3^- and the temperature optimum of NR activity may in part explain the timing of the occurrence of diatom and flagellate blooms.

Ammonium is incorporated into C skeletons mainly via the GS/GOGAT cycle (Lea et al., 1990), which is located in both chloroplast (Fig. 4) and cytosol (Fig.5). The GS/GOGAT cycle produces glutamine (Gln) and glutamate (Glu), which are the starting compounds for the synthesis of all rest amino acids, nucleic acids and chlorophylls (Fig. 6). Two GS isoenzymes, a cytosolic (GS_1) and chloroplastic (GS_2) one, have been identified in *E. huxleyi* (Maurin and Le Gal, 1997). Both isoforms are homohexamers with molecular masses of 402 kDa for GS_1 and 501 kDa for GS_2 , whereas the molecular masses of the subunits of GS_1 and GS_2 were estimated to be 61 and 70 kDa, respectively (Maurin and Le Gal, 1997a). The same authors reported that the K_m for hydroxylamine (NH_2OH) was approximately 3 mM for both

GS isoforms, but GS₂ had higher affinity for Gln than GS₁. The genes coding for the GS and GOGAT enzymes have not been studied yet in *E. huxleyi*, and in the case of GOGAT no data is available for the expression pattern of the protein and its activity. Decreasing NO₃⁻ concentrations were found to stimulate GS and GOGAT activity in various phytoplankton cells (Tischner and Hüttermann, 1980; Menacho and Vega, 1989; Maurin and Le Gal, 1997b) and to increase the affinity of GS for NH₄⁺ in *E. huxleyi* (Maurin and Le Gal, 1997b). GS activity was also found to exhibit a marked diel periodicity in *E. huxleyi* with maximal values during the dark period and a minimum during the light period (Maurin-Defosse and Le Gal, 1998).

1.4. Carbon skeletons for amino acid synthesis

The C skeletons for amino acid synthesis in the GS/GOGAT cycle are mainly provided via the tricarboxylic acid (TCA) cycle (Elfiri and Turpin, 1986; Weger and Turpin, 1989) (Fig. 7). The removal of intermediates of the TCA cycle requires that the cycle is replenished of its intermediates via anaplerotic reactions (Beardall and Raven, 1990; Norici and Giordano, 2002). Furthermore, the provision of inorganic carbon (Ci) for these reactions may be crucial to sustain amino acid and protein synthesis (Norici and Giordano, 2002). Tsuji et al. (2009) reported that β -carboxylases including phosphoenolpyruvate carboxykinase (PEPCK) contribute to the anaplerotic reaction by providing C skeletons for amino acid and lipid synthesis. Although the exact role of β -carboxylases in response to N availability has not been

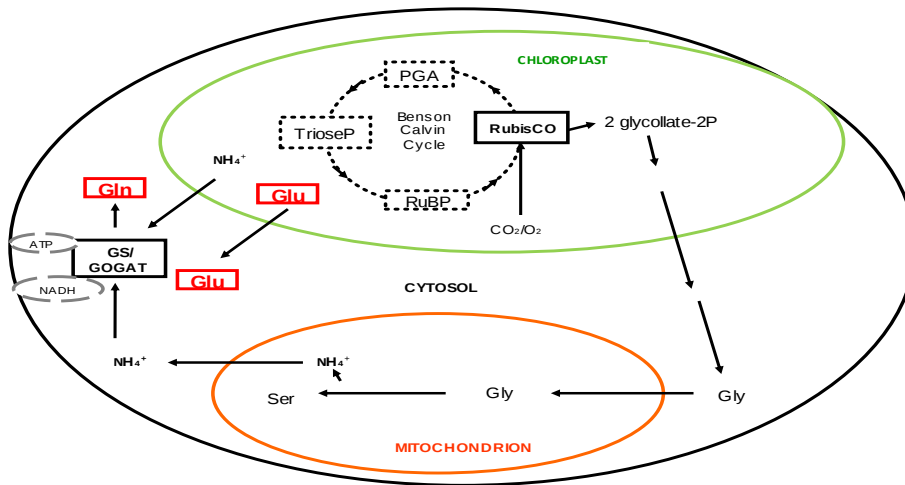
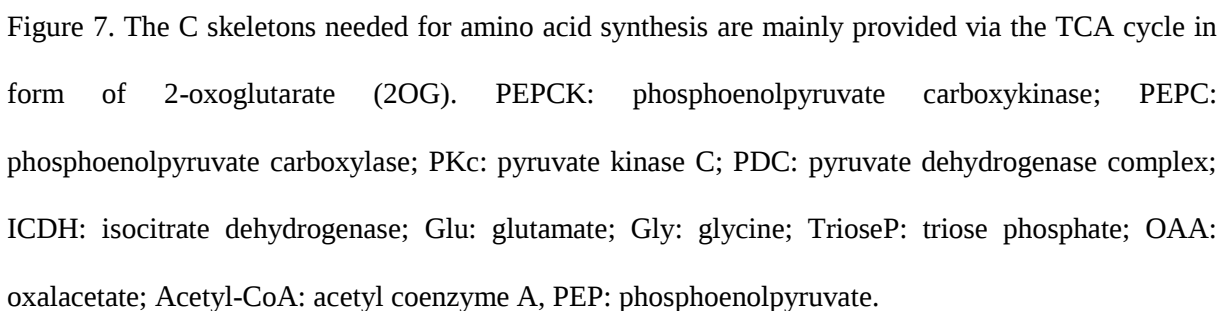
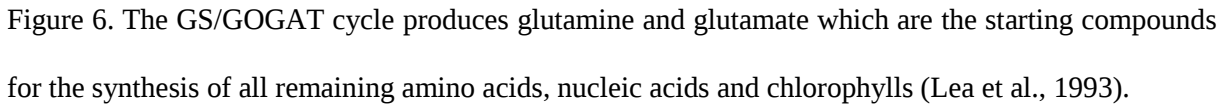


Figure 5. Cytosolic ammonium assimilation. NH_4^+ : ammonium; GS: glutamine synthetase; ATP: adenine triphosphate; NADH: nicotinamine adenine dinucleotide; Glu: glutamate; Gln: glutamine; RubisCO: ribulose biphosphate carboxylase/oxygenase; RuBP: ribulose biphosphate; Gly: glycine; Ser: serin; PGA: phosphoglycerate; TrioseP: triose phosphate.

studied yet in *E. huxleyi*, NO_3^- supply to N limited *Selenastrum minutum* led to increased β -carboxylase activities in this species (Elfiri and Turpin, 1987). Furthermore, Norici et al. (2002) reported that the expression of phosphoenolpyruvate carboxylase (PEPC) isoforms in *Dunaliella salina* responds to the variation in the C skeleton demand deriving from changes in the chemical form and availability of N. In general, the capacity for N assimilation increases in cells under N limitation, as reported for N limited cells of *Selenastrum minutum* (Birch et al., 1986; Elfiri and Turpin, 1986).



1.5. Inorganic carbon assimilation

The inorganic C (Ci) acquisition has been suggested to play an important role in the evolution and ecology of marine phytoplankton (Badger et al., 1998; Tortell, 2000; Giordano et al., 2005). The main carboxylating enzyme, ribulose1,5-bisphosphate carboxylase/oxygenase (RubisCO) is characterized by a low affinity for its substrate and microalgae cells have to invest an important amount of energy in order to enhance CO₂ concentration in the carboxylation site of RubisCO and avoid Ci limitation (Badger et al., 1998). Therefore, cells have developed special mechanisms, i.e. carbon concentrating mechanisms (CCMs), including the active uptake of CO₂ or HCO₃⁻ or both as well as the enzyme carbonic anhydrase (CA), which accelerates the otherwise slow conversion rate between HCO₃⁻ and CO₂ (Fig. 8). The β -carboxylase PEPC may also contribute, together with active HCO₃⁻ uptake, to the CCM as reported for diatoms (C₄-like pathway) (Reinfelder et al., 2000, 2004). The effects of N availability on the regulation of the CCM have still to be studied in *E. huxleyi*. Nevertheless, the CCM of this species was found to be regulated in response to light and CO₂ (Rost et al., 2003; Rost et al., 2006; Trimborn et al., 2007). In general, the CCMs have been suggested to improve N-use efficiency in microalgae, mainly by increasing the achieved rate of CO₂ fixation per unit N in Rubisco (Raven, 1997; Beardall et al., 1998), thereby controlling the cellular elemental ratios, specifically the C to N ratio (Beardall and Giordano, 2002).

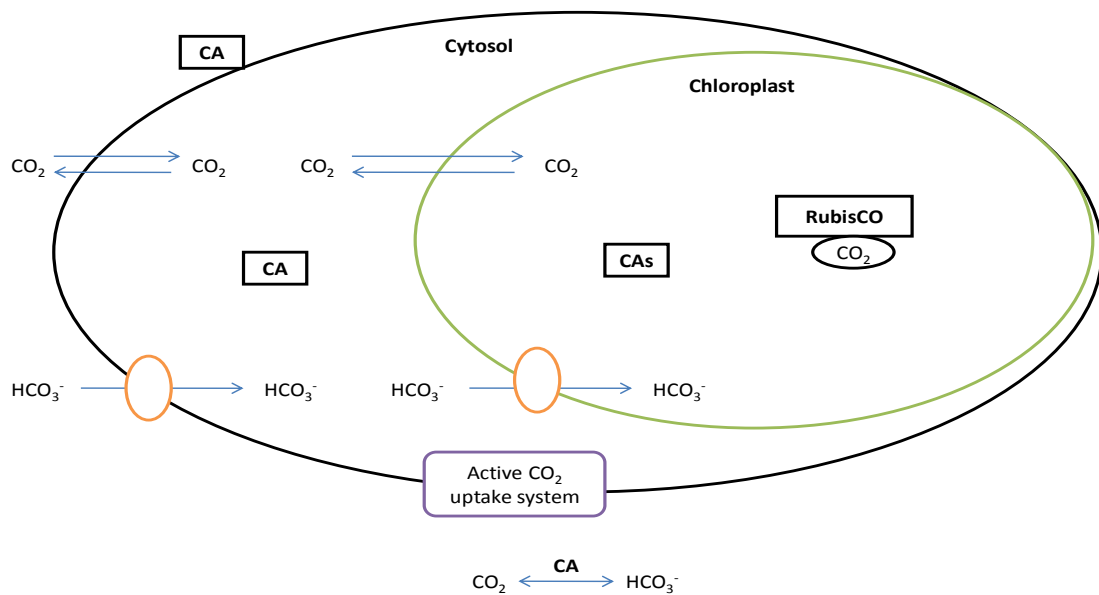


Figure 8. Carbon concentrating mechanisms (CCMs) in eukaryotic algal cells. Bicarbonate (HCO_3^-) is taken up by specific HCO_3^- transporters, whereas CO_2 is mainly taken up by diffusion. An active CO_2 system seems to exist, but it is poorly described till now. The enzyme carbonic anhydrase (CA) catalyzes the conversion between CO_2 and HCO_3^- . Various isoenzymes of CA have been described, which are located in the chloroplast stroma, the thylakoid membrane, the cytosol, the mitochondrion (not shown here) and the extracellular space. RubisCO: ribulose biphosphate carboxylase/oxygenase.

1.6. Photosynthesis and respiration

In photosynthetic organisms, all ATP and reductant (NADH/NADPH) come from photosynthesis directly through the light-dependent reactions or indirectly through respiration of fixed C and this enables cells to afford the energy requirements of C and N assimilation (Tab. II). Nitrogen limitation was found to decrease the rate of photosynthesis in the diatom *Phaeodactylum tricornutum* (Osborne and Geider, 1986). The most obvious effect, which causes the decrease in photosynthesis under N limitation, is the decline in nitrogenous

photosynthetic pigments such as chlorophylls (Chl) and phycobilins (Falkowski et al., 1989; Herzig and Falkowski, 1989; Plumley et al., 1989). Furthermore, the carotenoid to chl α ratio increases dramatically under N limitation (Plumley et al., 1989). Nitrogen limitation also decreases thylakoid stacking and absorptivity (Rhiel et al., 1985, 1986). Interestingly, N limitation was found to increase, decrease or maintain the apparent Chl specific efficiency of light harvesting (α_{chl}), depending on the investigated organism and the degree of N limitation (Osborne and Geider, 1986; Kolber et al., 1988; Herzig and Falkowski, 1989). Similarly to photosynthetic pigments, the enzymes of photosynthetic C fixation are also affected under N limitation and a decline in RubisCO per cell was reported in previous studies (Lapointe and Duke, 1984; Falkowski et al., 1989; Beardall et al., 1991). Respiration was shown to support NO_3^- and NO_2^- reduction during photosynthesis in *S. minutum* (Weger and Turpin, 1989). The impact of N availability on photosynthesis and respiration has still to be tested in the marine coccolithophore *E. huxleyi*.

1.7. Effects of nitrogen limitation on *Emiliania huxleyi*

The effect of NO_3^- availability on C and N assimilation in *E. huxleyi* cells was the subject of several studies in the past (Paasche, 1998; Fritz, 1999; Riegmann et al., 2000). Cellular particulate organic C (POC) and N (PON) have been shown to decrease with increasing NO_3^- limitation (Riegmann et al., 2000). According to Riegmann et al. (2000), various levels of N limitation did not have any influence on cellular inorganic C (PIC) but cellular PIC production decreased due to decreasing growth rate. In contrast, cellular PIC was found to increase with increasing NO_3^- limitation in cyclostat cultures (Fritz, 1999). Moreover, *E. huxleyi* cells were covered by multiple layers of coccoliths at low NO_3^- concentrations (Lindschooten et al., 1991; Fritz, 1999). Nitrate limitation has also been shown to decrease the calcite content of coccoliths (Paasche et al., 1998).

Table II. The energetics of CO₂ and N assimilation (simplified). Table I outlines the ATP and reductant requirements of the various processes associated with photosynthetic C fixation and the assimilation of N into protein (in Turpin, 1991).

The equations are not balanced.

Process	Reactions	ATP/e ⁻ requirement
1. Photosynthetic CO ₂ fixation	$\text{CO}_2 + \text{RuBP} + 3\text{ATP} + 4\text{e}^- \rightarrow \text{RuBP} + 1/3 \text{ triose-P}$	0.75
2. N transport	$\text{N}_{\text{out}} + \text{ATP} \rightarrow \text{N}_{\text{in}}$	
3. NO ₃ ⁻ reduction (NR)	$\text{NO}_3^- + 2\text{e}^- \rightarrow \text{NO}_2^-$	
4. NO ₂ ⁻ reduction (NiR)	$\text{NO}_2^- + 6\text{e}^- \rightarrow \text{NH}_4^+$	
5. NH ₄ ⁺ assimilation (GS/GOGAT)	$\text{NH}_4^+ + \text{Glu} + \alpha\text{-KG} + \text{ATP} + 2\text{e}^- \rightarrow 2\text{Glu}$	
6. Amino acid interconversion (transaminase)	$\text{Glu} + \text{keto-acid} \rightarrow \text{amino acid} + 2\text{-OG}$	
7. Protein synthesis	$\text{Amino acid} + 4\text{ATP} \rightarrow \text{polypeptide}$	
8. NO ₃ ⁻ assimilation into protein (2 + 3 + 4 + 5 + 7)	$\text{NO}_3^- + \text{keto-acid} + 6\text{ATP} + 10\text{e}^- \rightarrow \text{protein}$	0.6
9. NO ₂ ⁻ assimilation into protein (2 + 4 + 5 + 7)	$\text{NO}_2^- + \text{keto-acid} + 6\text{ATP} + 8\text{e}^- \rightarrow \text{protein}$	0.75
10. NH ₄ ⁺ assimilation into protein (2 + 5 + 7)	$\text{NH}_4^+ + \text{keto-acid} + 6\text{ATP} + 2\text{e}^- \rightarrow \text{protein}$	3

1.8. Objectives of the study

Although biochemical studies have been conducted on sole proteins such as NR (Iwamoto and Shiraiwa, 2003) and GS (Maurin and Le Gal, 1997) in *E. huxleyi*, knowledge on the effect of NO_3^- availability on the kinetics of inorganic C/N acquisition and assimilation is still lacking for this common phytoplankton species. The main goal of this study is to describe the C and N fluxes from the cellular to the environmental level, with starting point the intracellular processes involved in C/N metabolism and end point the elemental ratios which define the impact of the species on its environment. Therefore the regulation of NR and NiR kinetics which describe the capacity of the NO_3^- to NH_4^+ reduction pathway is investigated in response to NO_3^- availability. Ammonium assimilation into C skeletons via the GS/GOGAT cycle is studied by measuring the kinetics of GS and GOGAT activity, but also through determination of the GS isoforms in *E. huxleyi* cells grown under different NO_3^- conditions. The acquisition of inorganic C in cells is investigated via the kinetics of CO_2 and HCO_3^- uptake in response to NO_3^- availability. The impact of the NO_3^- concentration in the culture medium on photosynthesis, respiration and allocation of fixed C into the macromolecular pools is also studied in cells of *E. huxleyi*. Furthermore the elemental composition of cells, and especially cellular organic C and N as well as inorganic N, is also determined in cells grown at different NO_3^- conditions. Finally, the intracellular NO_3^- and CO_2 concentrations in the active sites of NR and RubisCO needed for adequate cellular growth are calculated in *E. huxleyi* cells in response to NO_3^- availability. With all these measurements, the hypotheses whether cellular C and N fluxes require a controlled regulation in response to NO_3^- availability and whether different NO_3^- conditions have an impact on the allocation of the macromolecular pools and growth in *E. huxleyi* cells are tested.

2. Materials and methods

2.1. Culture conditions

The coccolith-bearing strain B 92/11 (J. C. Green 1990, Plymouth Marine Laboratory) of *E. huxleyi* was grown in semi continuous cultures in 10-L polycarbonate flasks. Experiments were carried out under 16:8 h light-dark (LD) cycle at a constant temperature of 15°C. The applied mean photon flux density was 240 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$. The culture flasks were aerated with air containing CO₂ partial pressure (pCO₂) of 37.5 Pa and placed on a shaker, to keep the cells in suspension. The growth medium was prepared from sterile-filtered (0.2 μm pore-size cellulose-acetate filters, Sartorius) natural seawater. Nutrient additions, with the exception of iron and NO₃⁻, were made according to the f/2 recipe (Guillard and Ryther, 1962). The iron concentration in the culture medium was 8 nM and was sufficient to cover the iron need of the relative low cell densities in our experiment, as determined in preliminary experiments. The initial NO₃⁻ concentrations in the growth medium were ca. 280 μM in the high and ca. 10 μM in the low NO₃⁻ treatment. Over the course of the experiment, cell densities were kept between 3×10^4 and 3×10^5 cells mL⁻¹ by regular dilutions. This dilution pattern was kept throughout the whole experiment, which lasted 1 year. The cells were acclimated to the experimental conditions for approximately 2 weeks before measurements. Growth rates were calculated from daily cell counts by means of a cell counter (COULTER MULTISIZER 3, Beckmann).

2.2. Carbonate chemistry

Samples for total alkalinity (TA) measurements were filtered (nominal pore size 0.6 μm GFF filters, Whatman), and the filtrate was stored at 4°C and measured by means of potentiometric

titration with an average precision of $\pm 8 \mu\text{eq kg}^{-1}$ (Brewer et al., 1986). TA was calculated from linear Gran plots (Gran, 1952). Samples for dissolved inorganic carbon (DIC) were sterile-filtered (0.2 μm pore size cellulose-acetate syringe filters, Sartorius, Göttingen, Germany) and stored free of air bubbles at 4°C until they were measured with an average precision of $\pm 10 \mu\text{mol kg}^{-1}$ (QUAATRO, Seal Analytical). Values for pCO_2 were calculated using the program CO₂sys (Lewis and Wallace, 1998). Equilibrium constants of Mehrbach et al. (1973) refitted by Dickson and Millero (1987) were chosen.

2.3. Cellular carbon and nitrogen

Samples for total particulate C (TPC), POC and PON were filtered onto precombusted (500°C, 12 h) filters (nominal pore size 0.6 μm GFF filters, Whatman) and stored in precombusted (500°C, 12 h) petri-dishes at -20°C. Filters for POC were treated before analysis with 200 μl HCl (0.1 M) to remove all inorganic C. TPC, POC and PON samples were measured on an elemental analyzer mass spectrometer (ANCA-SL 2020, Secron). Values for PIC were calculated as the difference between TPC and POC. POC, PON and PIC production was measured by multiplying the cellular POC, PON and PIC content with the growth rate.

2.4. Macromolecular composition

Cells for fourier transform infrared spectroscopy (FTIR) analyses were harvested by centrifugation at 1500 x g for 15 minutes and washed twice with an isoosmotic solution of ammonium formate, to minimize the carry-over of IR absorbing medium components. Cells were resuspended in ammonium to obtain the concentration that was preliminary determined to afford the best signal to noise ratio. Aliquotes of 50 μL of these suspensions were deposited on silica windows (Crystran Ltd., Poole, UK) and desiccated in an oven at 60°C for at least 3

hours. Silica windows on which 50 μL of ammonium formate solution were deposited were treated as the samples and used as blanks (Domenighini and Giordano, 2009).

FTIR spectra were acquired with a Tensor 27 FTIR spectrometer (Bruker Optics, Ettlingen, Germany) as described in Domenighini and Giordano (2009). All spectra were baseline corrected prior to any analysis applying the Rubberband correction function of the OPUS 6.0 software (Bruker Optics, Ettlingen, Germany) and then normalized to the amide I band ($\sim 1650\text{ cm}^{-1}$). Bands were assigned to the macromolecular pools according to Giordano et al. 2001. The sum of the integrals of the absorption bands at $\sim 1160\text{ cm}^{-1}$, $\sim 1080\text{ cm}^{-1}$ and $\sim 1030\text{ cm}^{-1}$ was used as a proxy for carbohydrates. The amide I peak (1650 cm^{-1}) was used as an indicator of protein relative abundance. The $\sim 1740\text{ cm}^{-1}$ feature was used for lipids. Relative ratios of protein, carbohydrates and lipids were calculated from the bands' integrals, using the OPUS 6.5 software (Bruker Optik GmbH, Ettlingen, Germany). The calculation of the band integrals was performed after deconvolution of the spectrum from 1800 cm^{-1} to 1000 cm^{-1} using the Peak fit function of OPUS 6.5; to minimize subjective assessments, the main peaks on which deconvolution was based were identified by the application to each spectrum of a second derivative, with 9 smoothing points.

2.5. Coccolith morphology

Samples for scanning electron microscope analysis were filtered on polycarbonate filters (0.2 μm pore size, Whatman), dried in a drying cabinet at 60°C for 24 hours, then sputter-coated with gold-palladium. Imaging was performed with a digital scanning field-emission electron microscope (XL-30, Philips). Four categories were used to describe the morphology of *E. huxleyi* coccoliths: 'normal', 'malformed', 'incomplete', and 'incomplete and malformed'. An

average of approx. 350 coccoliths was analyzed per sample. For determination of the number of visible coccoliths per cell an average of approx. 50 cells per sample was counted.

2.6. Enzyme extraction

The cells were sampled about 4 hours after the beginning of the light period. Cells were concentrated by centrifugation at 2,772 *g* for 6 min at 4°C (Megafuge 1.0 R, Heraeus), frozen in liquid N₂ and stored at – 80°C. The cells were gently disrupted using a hand operated glass homogenizer (Wheatman) at 4°C. All extracts were centrifuged at 16,060 *g* for 15 min at 4°C (Biofuge fresco, Heraeus), in order to spin down the calcite. The supernatant was centrifuged again at 61,740 *g* for 20 min at 4°C (OPTIMA™ LE-80K Ultracentrifuge, Beckmann). The resulting supernatant was used to study enzyme kinetics and measure total protein amount. The protein amount in extracts was measured according to the method of Bradford (1976), using BSA as standard. All solutions were made in Milli-Q H₂O (Milli-Q academic A10, Millipore) and the chemicals were purchased by Sigma-Aldrich (St. Louis, MO, USA).

2.7. Nitrate reductase

Nitrate reductase activity was determined by NO₃[–] reduction to NO₂[–] and subsequent colorimetric measurement of the NO₂[–] produced according to the method described by Giordano et al. (2005b). The extraction buffer for NR (buffer A) contained 50 mM HEPES (pH 7.5), 1 mM EDTA, 1 mM dithiothreitol, 0.1% Triton X-100 (v/v), 0.3% polyvinylpolypyrrolidone (w/v) and 10 mM MgSO₄. The amount of NO₂[–] was measured according to the method of Snell and Snell (1949) in a microplate reader (SPECTRAFLUOR, Tecan) at 535 nm. Controls included the same reaction mixture, but extraction buffer was used instead of extract. The amount of NO₂[–] in the tubes was quantified according to a NO₂[–]

standard curve from 0 to 40 μM . Nitrate concentrations from 0.005 to 20 mM and NADH concentrations from 5 to 400 μM were used for the determination of NR kinetic parameters.

2.8. Nitrite reductase

Activities of NiR were determined using an assay similar to Wray and Filner (1970), which is based on the colorimetric measurement of the NO_2^- left in the reaction mixture. Buffer A was used for NiR extraction and the reaction was run in open tubes. The assay mixture contained in a total volume of 500 μL : 100 μL of 150 mM phosphate buffer (pH 7.5), 20 μL Milli-Q H_2O , 250 μL of 32 mM NaNO_2 , 5 μL of a 100 mM methyl viologen solution, 50 μL of extract and 75 μL of fresh $\text{Na}_2\text{S}_2\text{O}_4$ (25 mg mL^{-1} $\text{Na}_2\text{S}_2\text{O}_4$ in 0.29 M NaHCO_3) solution. The reaction mixture was incubated for 15 min at 30°C and was initiated by adding the sodium dithionite solution. The reaction was stopped by vigorous shaking until the dithionite was completely oxidized and the solution turned from dark blue to colourless. Controls included the same reaction mixture, but extraction buffer was used instead of extract. The amount of NO_2^- was measured in a microplate reader at 535 nm according to the method of Snell and Snell (1949) with a standard curve from 0 to 40 μM NO_2^- . Nitrite concentrations from 0.25 to 16 mM were used in the assay, in order to determine the NiR kinetic parameters.

2.9. Glutamine synthetase

Glutamine synthetase activity was determined from the formation of L-Glu γ -monohydroxamate as described by O'Neal and Joy (1973), and Oaks et al (1980). The extraction buffer for GS (buffer B) contained 50 mM HEPES (pH 7.5), 1 mM EDTA, 2 mM dithiothreitol, 0.1% Triton X-100 (v/v), 0.3% polyvinylpyrrolidone (w/v) and 10 mM MgSO_4 . The extracts were desalted by applying them on a Sephadex column (PD-10

DESALTING COLUMN, GE Healthcare, Uppsala, Sweden). The extracts were eluted using 50 mM HEPES (pH 7.5). An assay mixture containing 160 mM HEPES (pH 7.5), 4.8 mM hydroxylamine (pH 7.5), 3.2 mM MgSO₄ and 0.16 mM EDTA was prepared. The reaction was conducted in a 96-well plate in a final volume of 200 µL. To a 76.4 µL aliquot of the assay mixture, we added 25.6 µL of 6.25 mM ATP (pH 7.5) and 76.8 µL extract. The reactants were incubated for 5 min at 30°C. At last, 21.2 µL of 0.6 M Glu were added, the microplate was incubated for 20 min at 30°C. The reaction was terminated by adding 40 µL of a FeCl₃ solution [2.5% FeCl₃ (w/v), 5% TCA (w/v), 1.5 N HCl]. The reactants were centrifuged at 16,060 g for 2 min. Controls were run, in which 50 mM HEPES (pH 7.5) was used instead of extract. A second control was run, in which the reaction was stopped by adding the FeCl₃ solution immediately after the Glu addition. The amount of L-Glu γ-monohydroxamate was measured in a microplate reader at 535 nm with a standard curve from 0 to 25 µM γ-L-Glu monohydroxamate. Glu concentrations from 0.5 to 63.6 mM, ATP concentrations from 0.025 to 3.2 mM and hydroxylamine concentrations from 0.02 to 4.8 mM were used to derive GS kinetic parameters.

The maximal enzyme activity ($V_{\max}$) and the Michaelis-Menten constant (K_m) for NR, NiR and GS were calculated from the Michaelis-Menten equation

$$V_o = \frac{V_{\max}[S]}{K_m + [S]}$$

where V_o is the enzyme activity, $V_{\max}$ is the maximum enzyme activity, $[S]$ is the substrate concentration and K_m is the Michaelis-Menten constant.

2.10. Glutamate synthase

Glutamate synthase activity was determined by the reduction in the absorbance of NADH similar to the methods of Singh and Srivastava (1986) and Lin and Kao (1996). Buffer A was used for GOGAT extraction. The assay mixture contained 125 mM HEPES (pH 7.5), 5 mM EDTA, 20 mM KCl and 875 μ M NADH. The reaction was conducted in a 96-well plate in a final volume of 250 μ L. To a 50 μ L aliquot of the assay mixture, we added 100 μ L extract, 50 μ L of 20 mM Gln and 50 μ L of 5 mM 2-oxoglutarate (2-OG). Controls included the same reaction mixture, in which the extract was substituted with extraction buffer. The reduction in NADH absorbance was measured for 2 min at 30 °C in a microplate reader (Spectrafluor, Tecan) at 340 nm, immediately after the addition of 2-OG.

2.11. Phosphoenolpyruvate carboxykinase

Phosphoenolpyruvate carboxykinase activity was determined through the coupled reaction with malate dehydrogenase (MDH), as described by Holsworth and Bruck (1977) and Johnston and Raven (1989). Buffer A, pH 7.9 was used for PEPCK extraction. The extracts were desalted by applying them onto a Sephadex column (PD-10 DESALTING COLUMN, GE Healthcare, Uppsala, Sweden) and proteins were eluted using 50 mM HEPES (pH 7.9). The assay mixture had the following composition: 50 mM HEPES (pH 7.9), 10 mM NaHCO_3 , 5 mM dithiothreitol, 200 μ M NADH and MDH (1 μ L mL^{-1} assay mixture). To 100 μ L extract, we added 67.75 μ L assay mixture, 12.5 μ L of 100 mM phosphoenolpyruvate, 25 μ L of 100 mM MgCl_2 , 12.5 μ L of 10 mM ADP and 32.25 μ L Milli-Q H_2O . PEPCK activity was measured from the rate of reduction of NADH absorbance for 60 min, at 30 °C, in a microplate reader (Spectrafluor, Tecan) at 340 nm. Two controls were run; one included Mn instead of Mg, the other contained Milli-Q H_2O instead of ADP. We did not measure PEPC

activity, because it was found to be one order of magnitude lower than PEPC activity in *E. huxleyi* strain PML 92/11 (data not shown).

2.12. Purification of the glutamine synthetase isoforms

We also verified whether different GS isoforms existed in *E. huxleyi* and if their relative amount was dependent on the growth NO_3^- concentration. The purification of GS isoforms was conducted by means of HPLC (Jasco PU-1580/LG-1580-04/DG-1580-54/MD-2010-plus/AS-1555/CO-1560, Jasco, Groß-Umstadt, Germany), similar to the GS purification procedure described in Maurin and Le Gal (1997a). Proteins were separated due to their molecular size, affinity to adenylates and charge. All solutions used during the purification procedure were air-free and each purification step was conducted at 4 °C. Proteins were extracted in buffer (B) at 4 °C. The crude extracts were desalted and proteins were concentrated by means of ultracentrifugation to a final volume of 200 μL (AMICON ULTRA CENTRIFUGAL FILTER UNITS, Sigma-Aldrich, St. Luis, MO, USA). The protein aliquots (3.5 and 3.6 mg mL^{-1} for high and low NO_3^- -grown cells, respectively) were initially run through a Sephacryl S-200 HR column (GE Healthcare, Uppsala, Sweden). The proteins were eluted with in HEPES [50 mM, pH 7.5) and fractions were collected at a flow rate of 1 mL min^{-1} . Detection of proteins was performed at 280 nm and two GS active fractions were detected. The GS active fractions were concentrated by ultrafiltration to a final volume of 200 μL . One aliquot was used for estimating GS activity, the rest was applied in a HiTrap Blue HP 1 mL column (GE Healthcare, Uppsala, Sweden) and eluted with buffer HEPES [50 mM, pH 7.5) at a flow rate of 1 mL min^{-1} . The active GS fractions were concentrated by ultracentrifugation to a final volume of 200 μL . One aliquot was used for estimating GS activity, the rest was applied to a MonoQ 5/50 GL column (GE Healthcare, Uppsala, Sweden) and proteins were fractionated using a linear salt gradient of 0-500 mM NaCl in buffer

HEPES (50 mM, pH 7.5) with a flow rate of 1 mL min⁻¹. At the end of the purification, aliquots of the GS active fractions were loaded on a 10% SDS-PAGE gel (Laemmli, 1970) and proteins were silver stained (SILVER STAIN KIT, Biorad).

2.13. Photosynthesis and inorganic carbon fluxes

Inorganic carbon (Ci) fluxes during steady-state photosynthesis were investigated by means of a sector-field multicollector mass spectrometer (ISOPRIME, GV Instruments, Manchester, UK). Net photosynthesis was measured by monitoring the O₂ concentration over consecutive LD intervals with increasing Ci concentrations (0 to ~ 4 mM). Light and dark intervals during the assay lasted 6 min. Simultaneous measurements of the CO₂ concentration enabled us to determine the CO₂ uptake and HCO₃⁻ uptake kinetics according to equations by Badger et al. (1994), using a photosynthetic quotient (PQ) of 1.4 (Williams and Robertson, 1991). All measurements were performed in f/2 medium with ca. 300 µM NO₃⁻ for high NO₃⁻-grown cells and ca. 10 µM NO₃⁻ for low NO₃⁻-grown cells and buffered with 50 mM HEPES (pH 8.0) at 15°C. Dextran-bound sulfonamide (DBS), a membrane-impermeable inhibitor of external carbonic anhydrase (CA), was added to the cuvette to a final concentration of 100 µM. Chl *a* concentration ranged between 0.5 and 1 mg mL⁻¹. The incident photon flux density was 300 µmol photons m⁻² s⁻¹. For further details on the method and calculation, we refer to Badger et al. (1994) and Rost et al. (2007). The kinetic parameters V_{max} and K_{1/2} were calculated from the Michaelis Menten equation.

Furthermore, net photosynthesis was measured under various light intensities (0 to 800 µmol photons m⁻² s⁻¹). While the same buffer (HEPES 50 mM, pH 8.0) and respective NO₃⁻ concentrations (300 versus 10 µM) were used as in the Ci flux assay measurements, the Ci

concentration were kept at ~ 2 mM. This way rates of photosynthesis, both expressed in terms of O_2 evolution and C fixation, could be measured as a function of PFD. The rate of photosynthesis (light-saturated photosynthetic rate) P was calculated from the equation

$P = P_m(1 - \exp(-P_1(x - P_2)))$. The apparent Chl α specific efficiency of light harvesting α was calculated from the equation $\alpha = P_m * P_1$, with units of $\mu\text{mol } O_2 (\text{mg Chl } \alpha)^{-1} \text{ h}^{-1} (\mu\text{mol photon m}^{-2} \text{ s}^{-1})^{-1}$.

2.14. Statistical analysis

All data was statistically analyzed using the T-test with a confidence level of 95%.

3. Results

3.1. Culture conditions

The cell densities of ca. 3×10^5 cells mL⁻¹ were found to reduce TA and DIC by 20% in the high and 10% in the low NO₃⁻ treatment, respectively (Tab. III). Cells at a density of 3×10^4 and 10^5 cells mL⁻¹ had no almost no impact on TA and DIC in the growth medium (Tab. III). This drift in DIC over the course of the experiment was generally low, ensuring only small changes in pH and pCO₂ and therefore is reported as a negligible effect on C and N assimilation in cells in our study.

Table III. Carbonate chemistry in the culture medium of the high (H) and low (L) NO₃⁻ treatments.

TA: total alkalinity; DIC: dissolved inorganic carbon; pCO₂: CO₂ partial pressure

NO ₃ ⁻ treatment	Cell density (cells mL ⁻¹)	TA (μeq kg ⁻¹)	DIC (μmol kg ⁻¹)	pCO ₂ (μatm)
H	30000	2277 ± 26	2047 ± 15	400.9
L	30000	2318 ± 21	2095 ± 6	431.5
H	100000	2242 ± 39	2024 ± 8	418.5
L	100000	2330 ± 50	2108 ± 39	438.2
H	300000	1917	1792 ± 5	~ 600
L	300000	2095	1914 ± 17	~ 500

3.2. Growth parameters

Although the growth rates were quite similar (1.1 to 1.2 d^{-1}) between high and low NO_3^- -grown cells (Tab. IV), low NO_3^- -grown cells accumulated less organic C and N. The cellular POC content was approximately one-third lower under low NO_3^- with values of $9.3 \pm 0.8\text{ pg cell}^{-1}$ in low and $13.9 \pm 2.4\text{ pg cell}^{-1}$ in high NO_3^- -grown cells (Tab. IV; T-test: $t = -3.27$, $p = 0.03$). Similarly, low NO_3^- -grown cells contained less PON ($1.3 \pm 0.1\text{ pg N cell}^{-1}$) than their high NO_3^- cells counterparts ($1.9 \pm 0.2\text{ pg N cell}^{-1}$) (Tab. IV; T-test: $t = -6.206$, $p = 0.0001$). Furthermore, cells in the low NO_3^- treatment were characterized by decreased protein content as well as cell size. The cellular protein content was about one-third lower when cells were cultured under low NO_3^- with values up to $1.2 \pm 0.1\text{ pg cell}^{-1}$ compared to a protein content of $1.7 \pm 0.1\text{ pg cell}^{-1}$ under high NO_3^- (Tab. IV; T-test: $t = -10.606$, $p = 0.0004$). The cell volume under low NO_3^- was ca. one-fourth lower ($272 \pm 37\text{ }\mu\text{m}^3$) compared to high NO_3^- ($394 \pm 46\text{ }\mu\text{m}^3$) (Tab. IV; T-test: $t = -4.606$, $p = 0.001$). Despite changes in POC and PON, the C to N ratio remained constant and was about 8.3 to 8.5 in both NO_3^- treatments (Tab. IV). POC production was significantly higher under high NO_3^- in comparison with low NO_3^- -grown cells, being $16.7 \pm 2.8\text{ pg C cell}^{-1}\text{ d}^{-1}$ and $10.4 \pm 1.2\text{ pg C cell}^{-1}\text{ d}^{-1}$, respectively (Fig. 9A, T-test: $t = 3.578$, $p = 0.023$). The PON production under high NO_3^- was $2.3 \pm 0.3\text{ pg N cell}^{-1}\text{ d}^{-1}$ whereas under low NO_3^- was $1.4 \pm 0.2\text{ pg N cell}^{-1}\text{ d}^{-1}$ (Fig. 9B, T-test: $t = -7.361$, $p = 0.00002$).

The cellular PIC content also decreased with decreasing NO_3^- availability. Low NO_3^- -grown cells contained $6.6 \pm 0.9\text{ pg C cell}^{-1}$ whereas high NO_3^- -grown cells had a PIC content of $8.7 \pm 0.9\text{ pg C cell}^{-1}$ (Tab. IV, T-test: $t = -2.763$, $p = 0.05$). The PIC to POC ratio was 0.6 to 0.7 (as can be calculated from the POC and PIC values in Tab. IV) and remained constant irrespective of the tested NO_3^- levels. PIC production was $10.5 \pm 1.1\text{ pg C cell}^{-1}\text{ d}^{-1}$ under high NO_3^- and $7.3 \pm 1\text{ pg C cell}^{-1}\text{ d}^{-1}$ under low NO_3^- (Fig. 9C, T-test: $t = -3.64$, $p = 0.021$).

Cells grown under low NO_3^- produced 0.2 ± 0.2 % normal coccoliths, 88.3 ± 0.5 % incomplete coccoliths, 0 % malformed coccoliths, and 11.5 ± 0.5 % incomplete and malformed coccoliths (Fig. 10 and 11). Cells grown under high NO_3^- produced 96.2 ± 1.9 % normal coccoliths, 2.4 ± 1.4 % incomplete coccoliths, 1.3 ± 0.5 % malformed coccoliths, and 0.1 ± 0.2 % incomplete and malformed coccoliths (Fig. 10 and 11). The number of visible coccoliths per cell was 6.3 ± 0.1 under high and 6.9 ± 0.4 under low NO_3^- .

Table IV. Growth rate (μ), cell volume (CV), particulate organic carbon (POC), particulate inorganic carbon (PIC), particulate organic nitrogen (PON), protein content and C to N ratio in cells grown under high (H) and low (L) NO_3^- .

The values represent the means of triplicate incubations ($\pm$ SD).

NO_3^- treatment	μ (d^{-1})	CV (μm^3)	POC (pg cell^{-1})	PIC (pg cell^{-1})	PON (pg cell^{-1})	Protein (pg cell^{-1})	C to N ratio
H	1.2 ± 0.1	394 ± 46	13.9 ± 2.4	8.7 ± 0.9	1.9 ± 0.2	1.7 ± 0.1	8.5 ± 0.5
L	1.1 ± 0.2	272 ± 37	9.3 ± 0.8	6.6 ± 0.9	1.3 ± 0.1	1.2 ± 0.1	8.3 ± 0.9

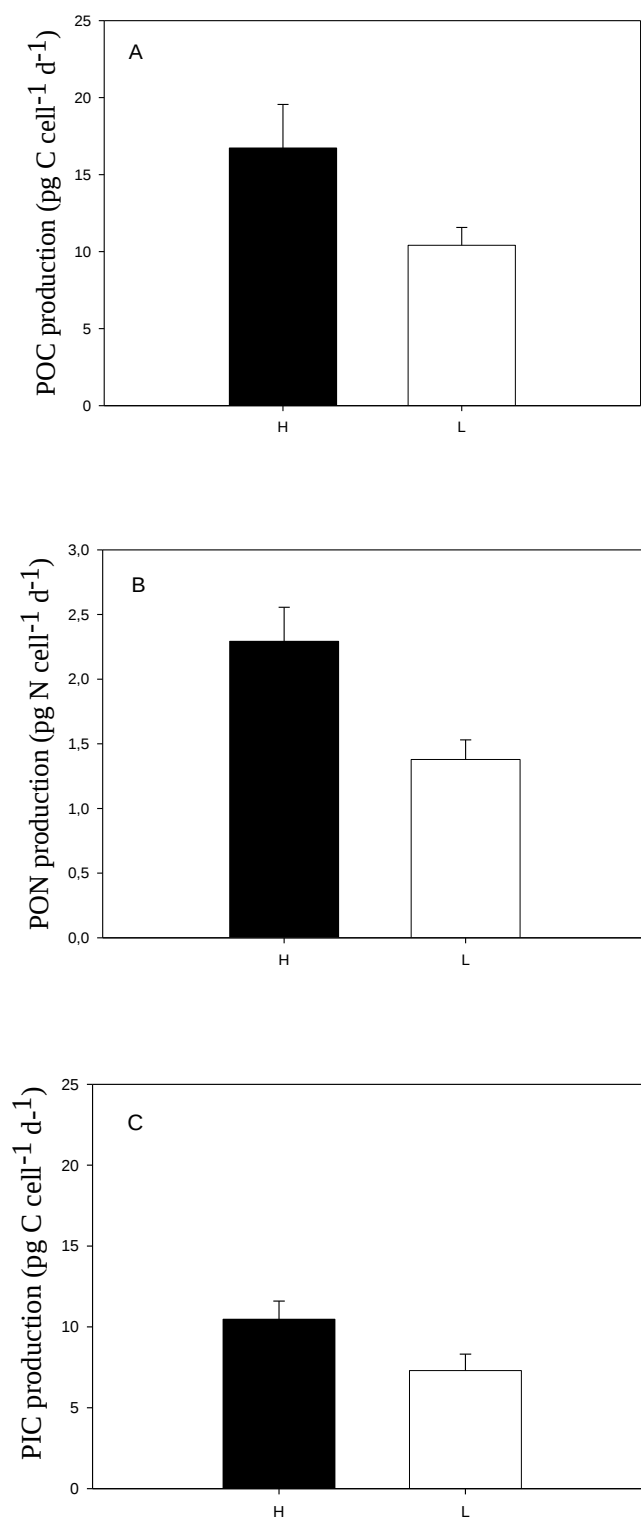


Figure 9. Effects of high ($\blacksquare$) and low ($\square$) NO_3^- on particulate organic carbon (POC) production (A), particulate organic nitrogen (PON) production (B) and particulate inorganic carbon (PIC) production (C) in *E. huxleyi* cells. Values represent the means of triplicate incubations ($\pm$ SD).

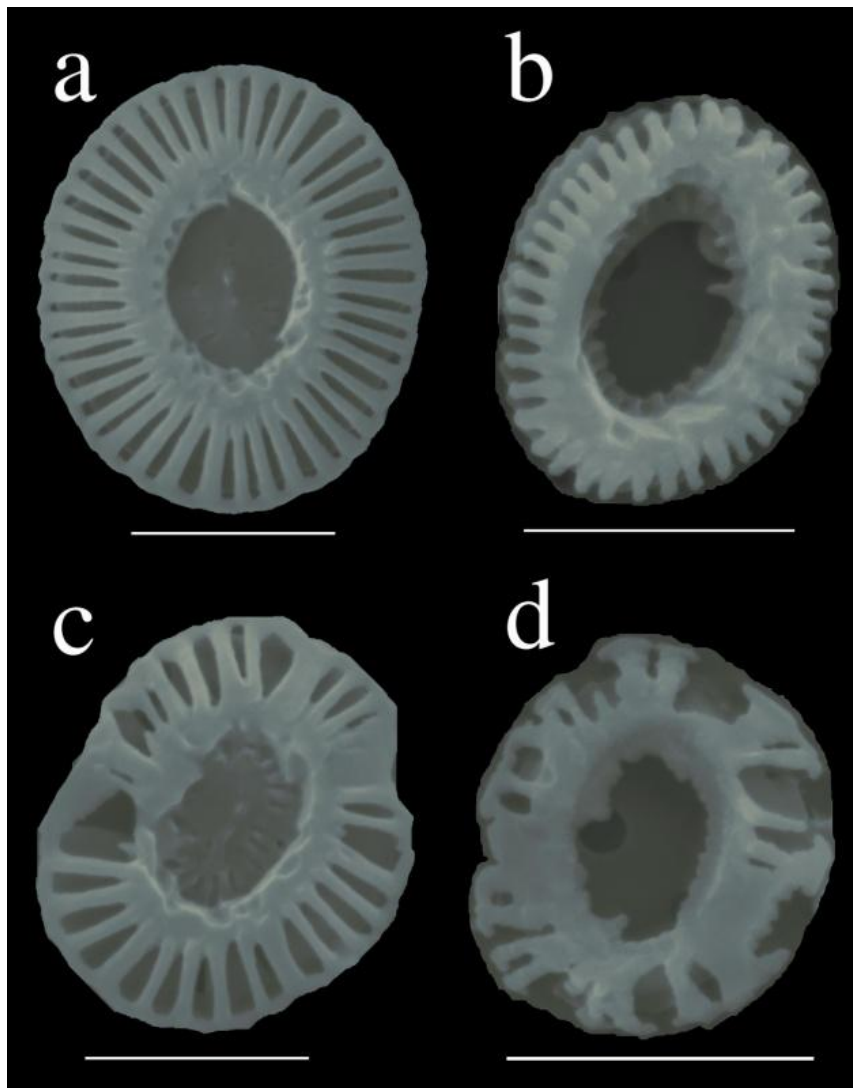


Figure 10. Scanning electron microscopy images of coccoliths of *E. huxleyi*. All scalebars 2 μm . All specimens in distal view. (a) normal coccolith. (b) incomplete coccolith. (c) malformed coccolith. (d) incomplete and malformed coccolith.

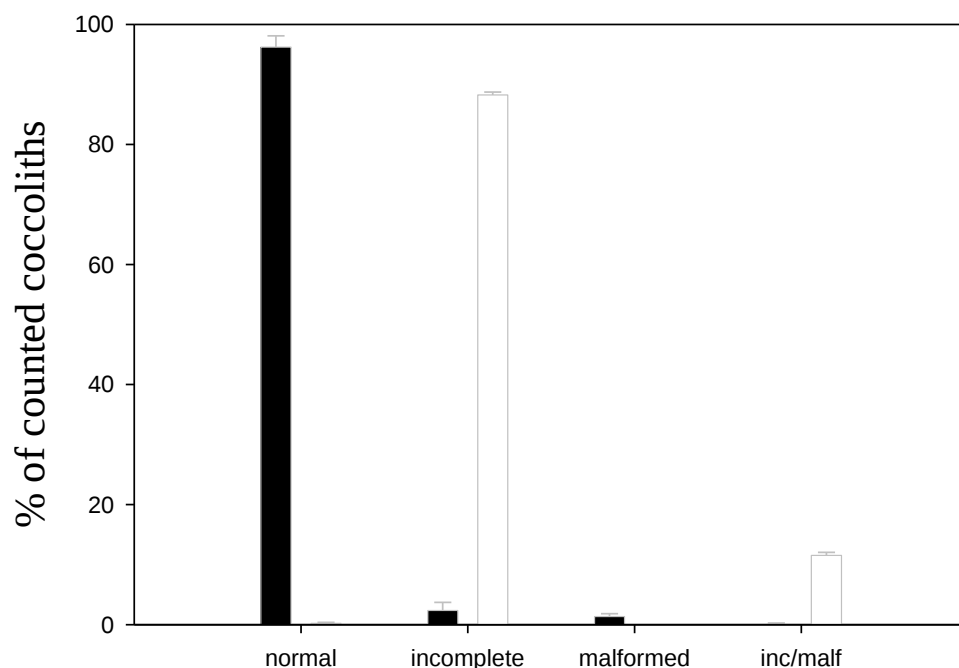


Figure 11. Percentage of normal, malformed, incomplete, and incomplete and malformed (inc/malf) cocoliths under high (■) and low (□) NO_3^- . Values represent an average of duplicates ($\pm$ SD).

3.3. Macromolecular composition

Nitrate availability caused an allocation of macromolecular pools in cells in our study. While the protein to lipid ratio decreased under low NO_3^- , the ratios of proteins to carbohydrates and lipids to carbohydrates increased. The protein to lipid ratio was 11.6 ± 2.2 under low NO_3^- compared to a value of 15.4 ± 1.2 under high NO_3^- (Fig. 12A). The ratio of proteins to carbohydrates was $3 \pm 0.3 (10^{-4})$ in low NO_3^- -grown cells and $2.1 \pm 0.3 (10^{-4})$ in high NO_3^- -grown cells (Fig. 12B). Similarly, the ratio of lipids to carbohydrates was $2.7 \pm 0.6 (10^{-6})$ under low NO_3^- and $1.4 \pm 0.1 (10^{-6})$ in cells under high NO_3^- (Fig. 12C).

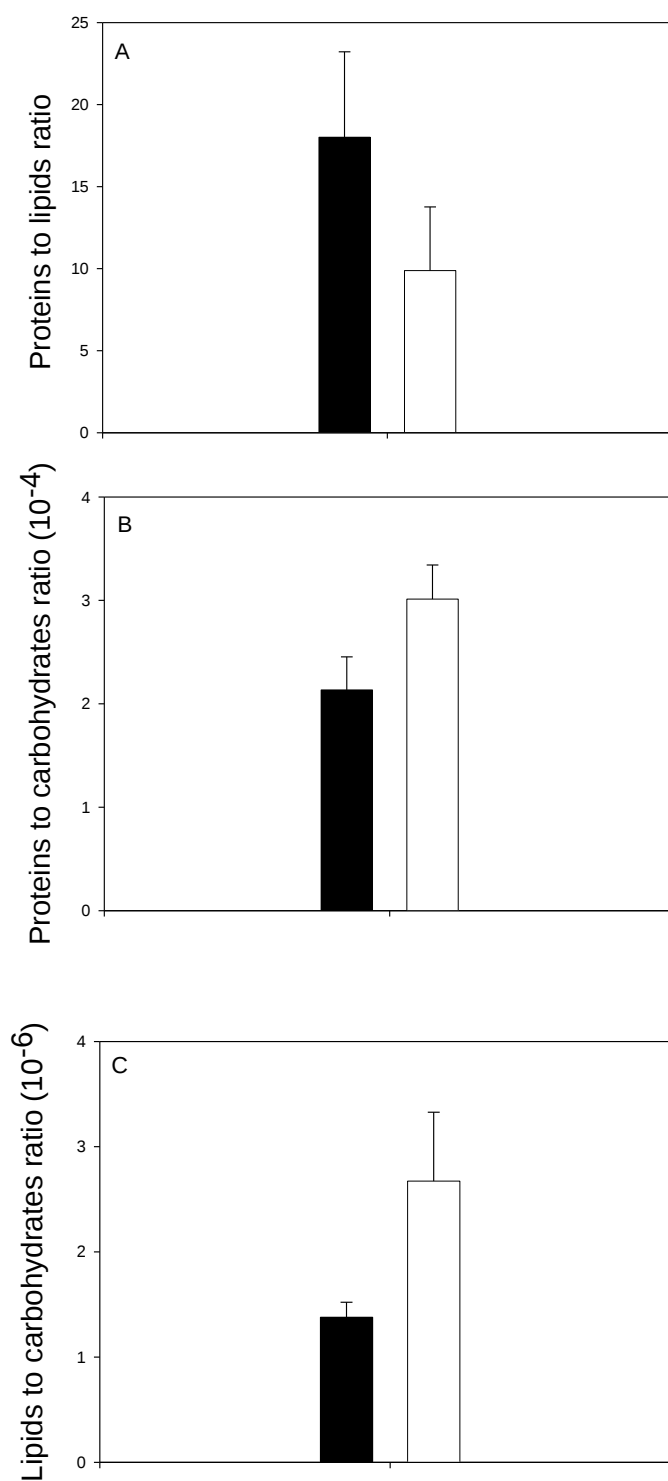


Figure 12. The ratios of proteins to lipids (A), proteins to carbohydrates (B) and lipids to carbohydrates (C) in cells under high (■) and low (□) NO_3^- . Values represent an average of triplicates ($\pm$ SD).

3.4. Nitrate assimilation

Nitrate availability was found to affect the kinetics of both NR and NiR. NR $V_{\max}$ was 1.7-fold lower in cells under low NO_3^- ($0.64 \pm 0.06 \text{ fmol NO}_2^- \text{ min}^{-1} \text{ cell}^{-1}$) compared to high NO_3^- -grown cells ($1.09 \pm 0.25 \text{ fmol NO}_2^- \text{ min}^{-1} \text{ cell}^{-1}$), when it was expressed on a cell basis (Tab. V; T-test: $t = -3.037$, $p = 0.038$). When expressed on a protein basis, cell volume, C or N, NR $V_{\max}$ was unaffected by the NO_3^- concentration in the medium. The Michaelis-Menten constant (K_m) of NR for NO_3^- differed between high and low NO_3^- -grown cells and was significantly lower in low NO_3^- cells. Cells at high NO_3^- had a K_m of NR for NO_3^- of $0.099 \pm 0.013 \text{ mM}$, whereas under low NO_3^- the K_m of NR for NO_3^- was $0.074 \pm 0.007 \text{ mM}$ (Tab. V; T-test: $t = -2.952$, $p = 0.041$; Fig. 13A). The K_m of NR for NADH was not significantly different between high and low NO_3^- -grown cells (Tab. IV; Fig. 13B). Regardless of the basis on which it was expressed, NiR $V_{\max}$ was lower under low NO_3^- availability in the growth

Table V. Kinetics $V_{\max}$ and K_m of NR and NiR in cells grown under high (H) and low (L) NO_3^- .

All values represent the means of triplicates ($\pm \text{SD}$).

NO_3^- treatment			H	L
	$V_{\max}$	$\text{fmol NO}_2^- \text{ min}^{-1} \text{ cell}^{-1}$	1.09 ± 0.25	0.64 ± 0.06
NR	$K_m(\text{NO}_3^-)$	mmol L^{-1}	0.099 ± 0.013	0.074 ± 0.007
	$K_m(\text{NADH})$	mmol L^{-1}	0.25 ± 0.013	0.206 ± 0.025
	$V_{\max}$	$\text{fmol NO}_2^- \text{ min}^{-1} \text{ cell}^{-1}$	0.56 ± 0.05	0.3 ± 0.01
NiR	$K_m(\text{NO}_2^-)$	mmol L^{-1}	3.14 ± 0.6	1.69 ± 0.43

medium (Tab. V). NiR $V_{\max}$ was 0.3 ± 0.01 $\text{fmol NO}_2^- \text{min}^{-1} \text{cell}^{-1}$ in cells grown under low NO_3^- and 1.9-fold higher (0.56 ± 0.05 $\text{fmol NO}_2^- \text{min}^{-1} \text{cell}^{-1}$) in cells grown under high NO_3^- (Tab. V; T-test: $t=-9.212$, $p=0.0007$). Also, the K_m of NiR for NO_2^- was substantially lower under low NO_3^- (1.69 ± 0.43 mM in low and 3.14 ± 0.6 mM in high NO_3^- -grown cells; Tab. V; T-test: $t=-3.408$, $p=0.027$; Fig. 14).

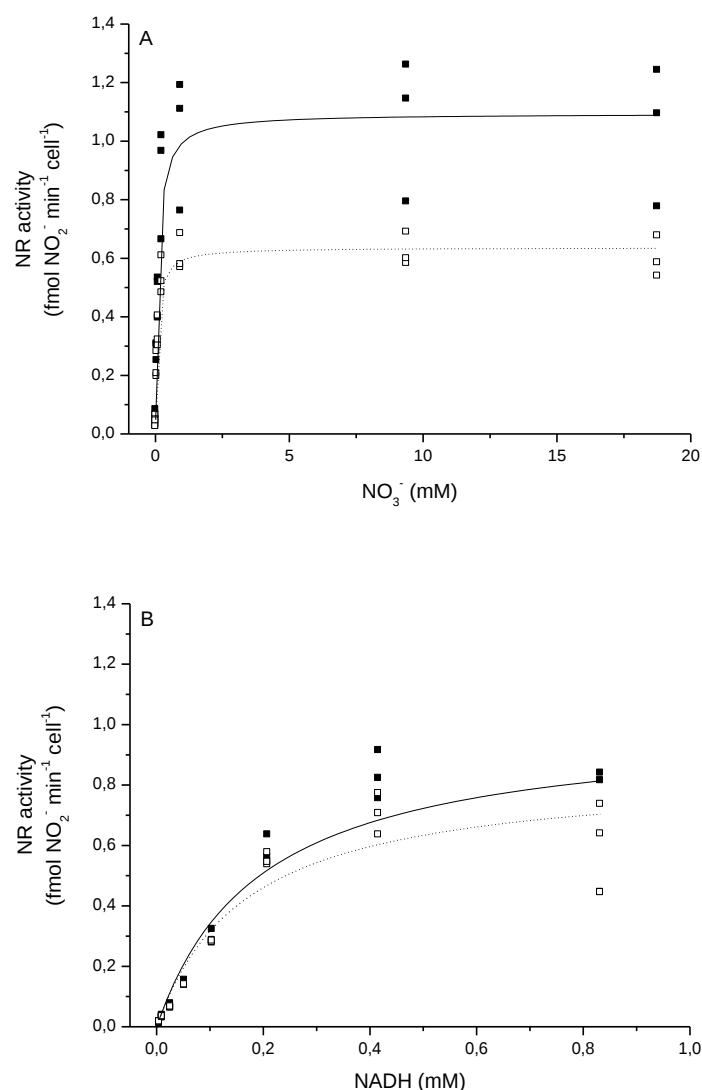


Figure 13. Kinetic curve of nitrate reductase (NR) for NO_3^- (A) and NADH (B) in cells grown under high (■) and low (□) NO_3^- . All values represent the means of triplicates.

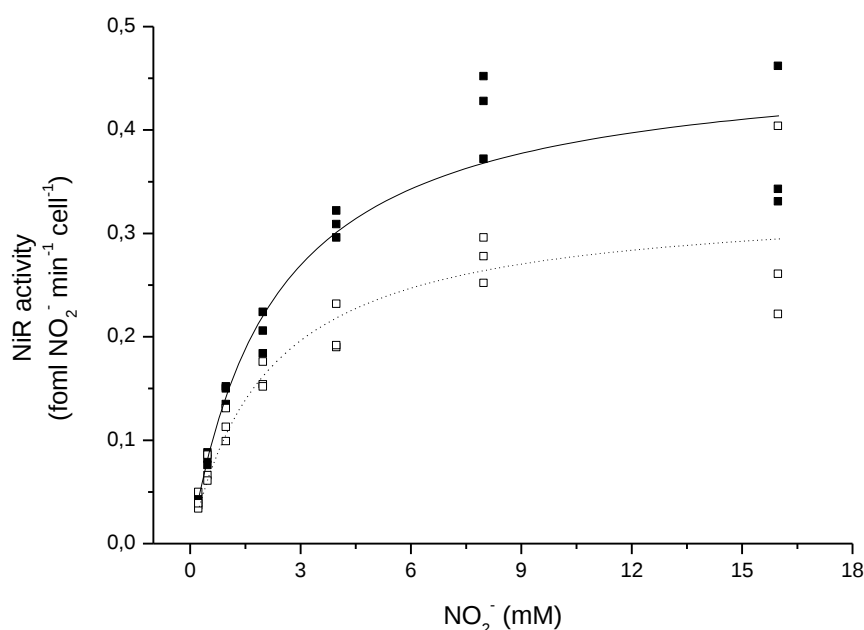


Figure 14. Kinetic curve of nitrite reductase (NiR) for NO₂⁻ in cells grown under high (■) and low (□) NO₃⁻. All values represent the means of triplicates.

The different NO₃⁻ concentrations also affected GS kinetics. GS $V_{\max}$ in low NO₃⁻-grown cells was about one-fourth higher than under high NO₃⁻ (Tab. VI; T-test: $t=-3.529$, $p=0.01$). GS $V_{\max}$ was higher in low NO₃⁻-grown cells and this relationship was independent of the type of normalization (protein, cell volume, cell C or N). The K_m for glutamate was two-fold lower in low NO₃⁻-grown cells (1.612 ± 0.37 mM) than in their counterparts (3.812 ± 1.63 mM) (Tab. VI; T-test: $t=-3.529$, $p=0.01$; Fig. 15). The K_m for ATP and NH₂OH were similar between high and low NO₃⁻-grown cells (Tab. VI; Fig. 16A and 16B). GOGAT $V_{\max}$ was 3.91 ± 0.76 fmol NADH min⁻¹ cell⁻¹ under low NO₃⁻ and 2.87 ± 0.18 fmol NADH min⁻¹ cell⁻¹ under high NO₃⁻, showing a significant difference in its activity between the two NO₃⁻ regimes (Tab. VI; T-test: $t = 2.639$, $p = 0.025$). When activities were expressed on protein, cell volume, C or N basis, GOGAT $V_{\max}$ was also higher in low NO₃⁻-grown cells. The $V_{\max}$ of the anaplerotic

enzyme PEPCK was about 1.6-fold lower in low NO_3^- grown-cells compared to high NO_3^- -grown cells, with values of $4.66 \pm 0.45 \text{ fmol NADH min}^{-1} \text{ cell}^{-1}$ under low and $7.29 \pm 0.28 \text{ fmol NADH min}^{-1} \text{ cell}^{-1}$ under high NO_3^- (Tab. VI; T-test: $t = -5.697$, $p = 0.001$). When normalized on protein, cell volume, C or N, however, PEPCK $V_{\max}$ did not differ between high and low NO_3^- cells.

Table VI. Kinetics $V_{\max}$ and K_m of GS, as well as the activities ($V_{\max}$) of GOGAT and PEPCK in cells grown under high (H) and low (L) NO_3^- .

All values represent the means of triplicates ($\pm$ SD). One unit is equal to 1 fmol L-Glu γ -monohydroxamate.

NO_3^- treatment			H	L
GS	$V_{\max}$	units $\text{min}^{-1} \text{ cell}^{-1}$	0.385 ± 0.11	0.57 ± 0.04
	K_m (ATP)	mmol L^{-1}	0.513 ± 0.157	0.344 ± 0.05
	K_m (NH_2OH) (n=4)	mmol L^{-1}	0.095 ± 0.035	0.092 ± 0.023
	K_m (Glu)	mmol L^{-1}	3.812 ± 1.63	1.621 ± 0.372
GOGAT	$V_{\max}$	$\text{fmol NADH min}^{-1} \text{ cell}^{-1}$	2.87 ± 0.18	3.91 ± 0.76
PEPCK	$V_{\max}$	$\text{fmol NADH min}^{-1} \text{ cell}^{-1}$	7.29 ± 0.28	4.66 ± 0.45

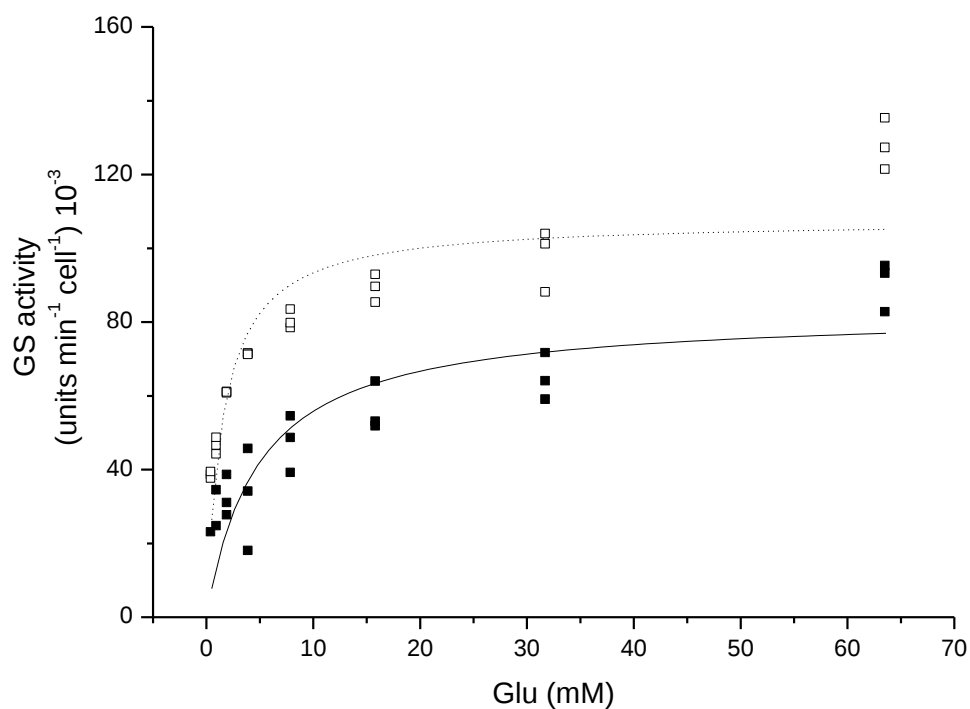


Figure 15. Kinetic curve of GS for glutamate (Glu) in cells grown under high (■) and low (□) NO₃⁻. One unit is equal to 1 fmol L-Glu γ -monohydroxamate. All values represent the means of triplicates.

3.5. Glutamine synthetase isoforms

After separation on the Sephacryl S-200 HR column, two fractions showed GS activity and were recognized as the chloroplastic (GS₂) and cytosolic (GS₁) isoform, according to their retention times (51-54 min for GS₂ and 62-65 min for GS₁, respectively, Fig. 17A and 17B).

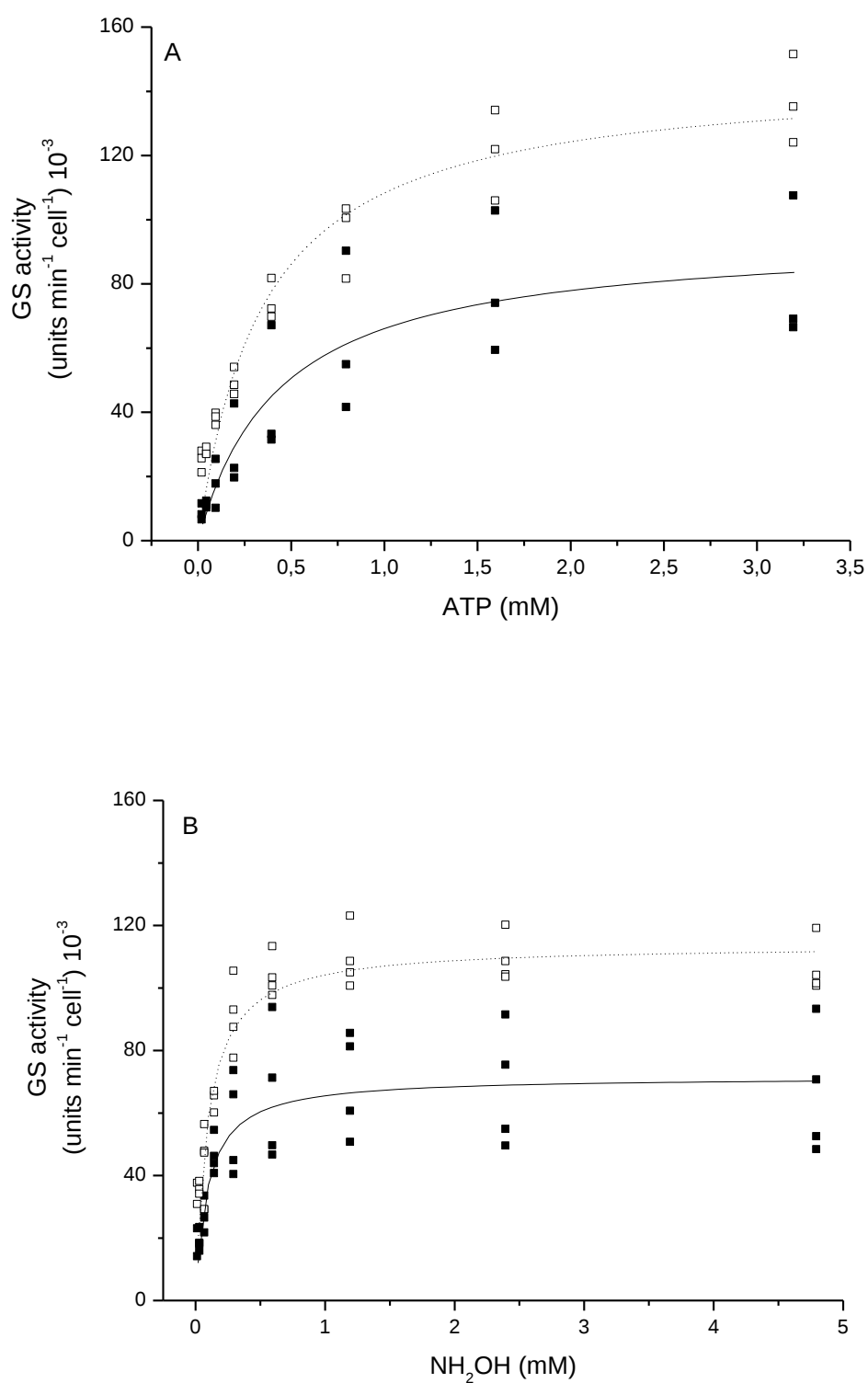


Figure 16. Kinetic curve of GS for ATP (A) and NH₂OH (B) in cells grown under high (■) and low (□) NO₃⁻. One unit is equal to 1 fmol L-Glu γ -monohydroxamate. In the kinetics of GS for ATP (n=3) and for NH₂OH (n=4).

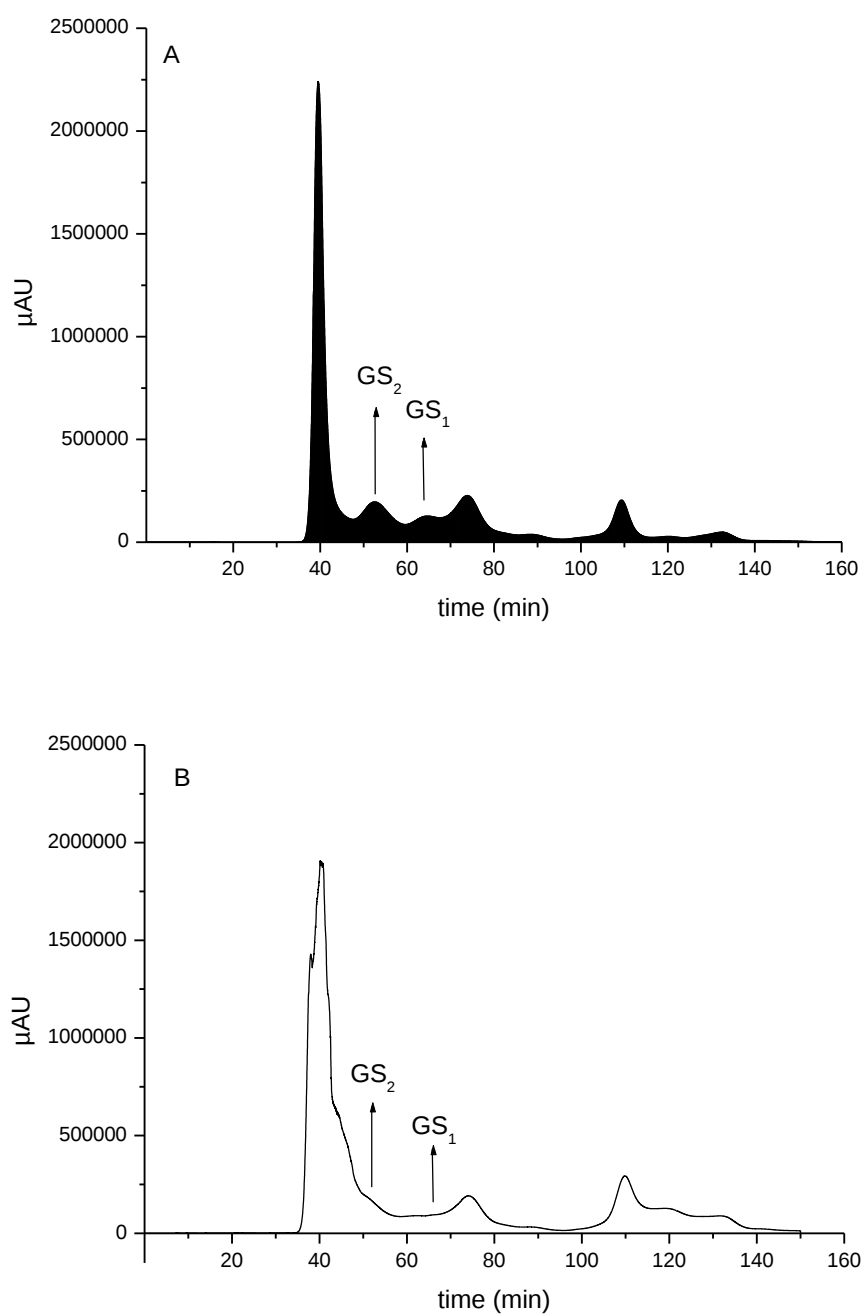


Figure 17. Purification of the GS isoforms in cells grown under high (■) (A) and low (□) (B) NO_3^- using a Sephacryl S-200 HR column. GS active fractions are shown with arrows.

Under high NO_3^- , the relative and specific activities of GS_1 were higher than GS_2 activity (Fig. 18A and 18B). On the opposite, both relative and specific activities of GS_2 were higher than GS_1 activity under low NO_3^- (Fig. 18A and 18B). After application on the HiTrap Blue

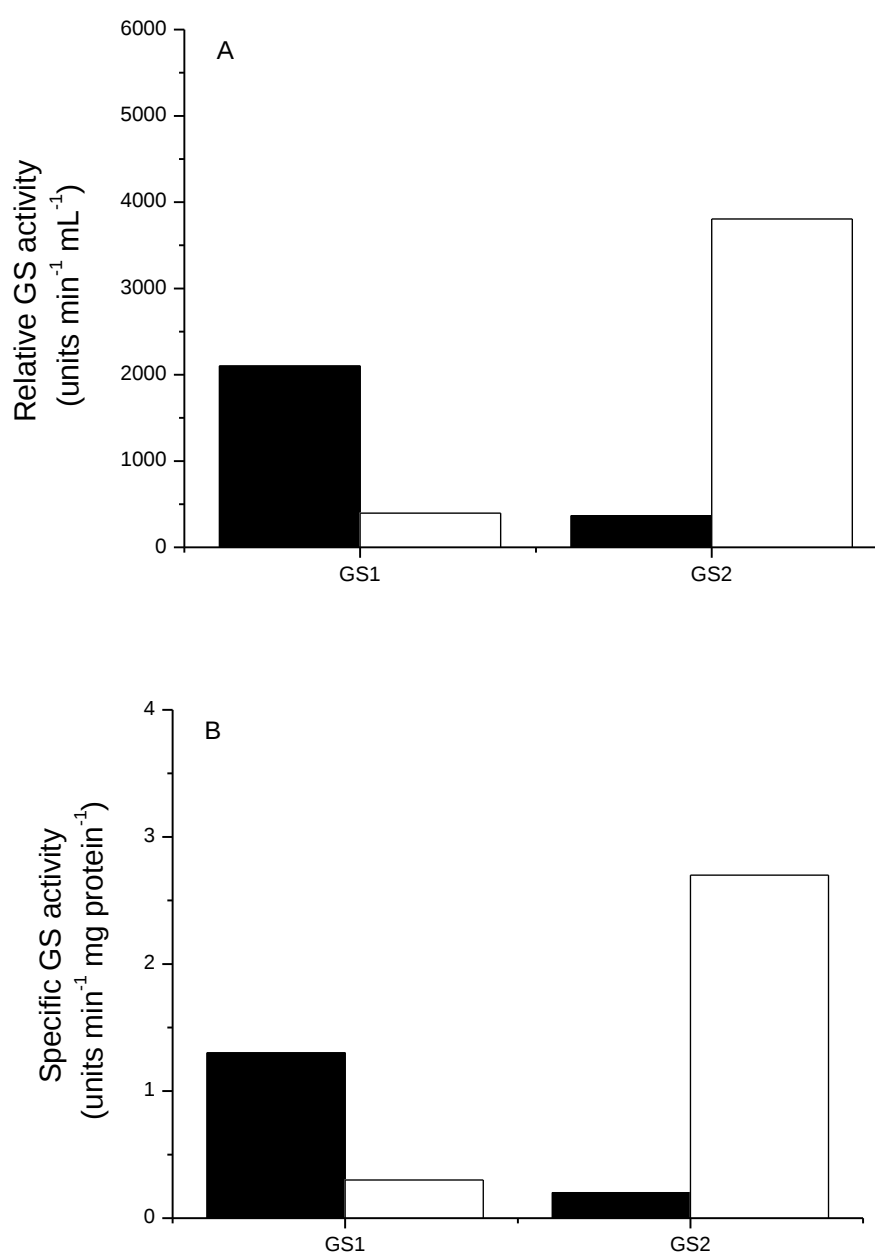


Figure 18. Relative (A) and specific (B) activities ($\times 5$) of the GS_1 and GS_2 isoforms in high (■) and low (□) NO_3^- -grown cells after separation on the Sephacryl S-200 HR column. GS_1 : cytosolic isoform; GS_2 : chloroplastic isoform.

HP column, GS₁ was eluted after 1-2 min and GS₂ after 7-8 min in both NO₃⁻ treatments (data not shown). After application of the GS active fractions on the MonoQ column, GS₁ was eluted after 4 min and GS₂ after 15-16 min in both NO₃⁻ treatments (Fig. 19A, 19B, 20A and 20B). GS₁ in low NO₃⁻-grown cells could not be further purified on the MonoQ column.

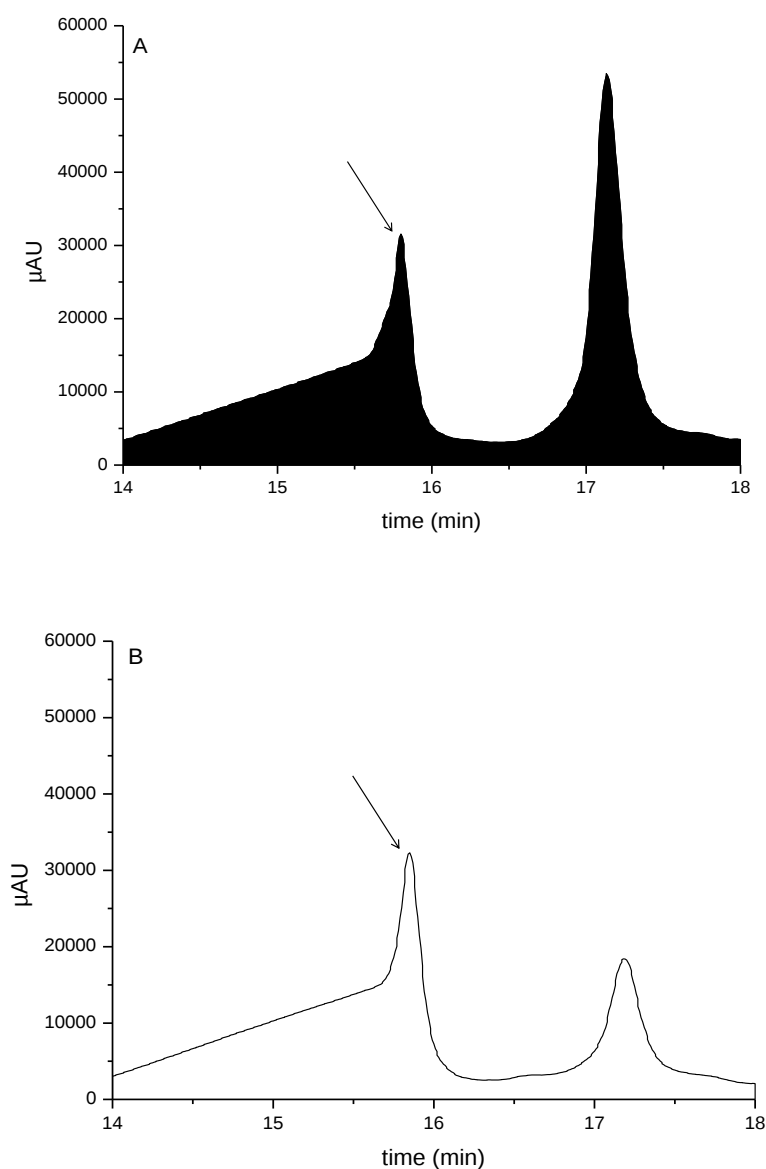


Figure 19. Purification of the GS₂ isoform in cells grown under high (■) (A) and low (□) (B) NO₃⁻ using a MonoQ column. GS active fractions are shown with arrows.

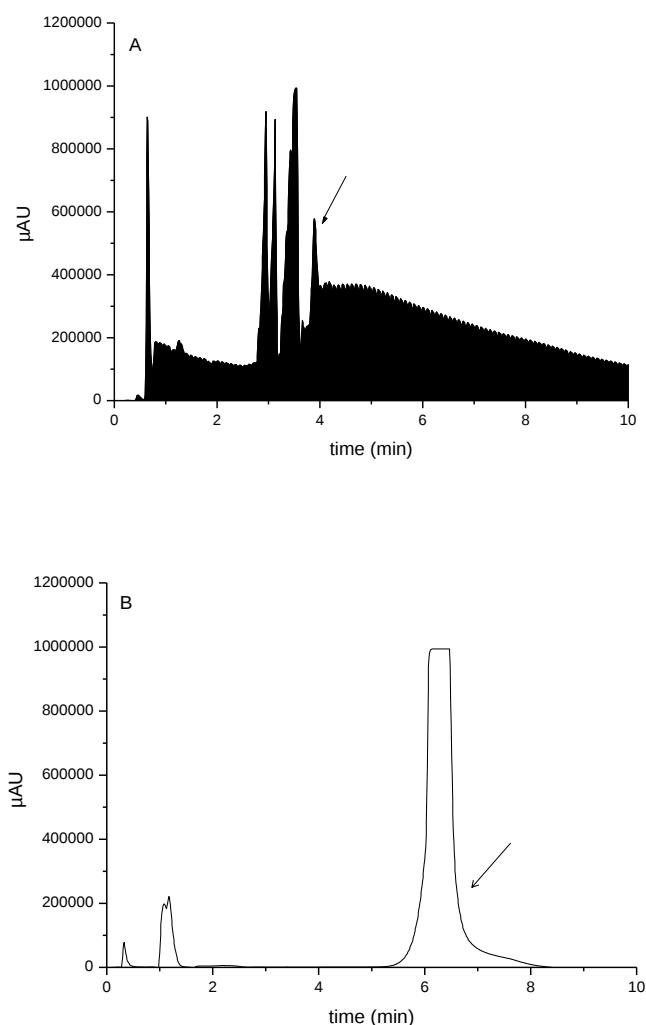


Figure 20. Purification of the GS₁ isoform in cells grown under high (■) (A) and low (□) (B) NO₃⁻ using a MonoQ column. GS active fractions are shown with arrows. The GS₁ isoform could not be further purified in the MonoQ column under low NO₃⁻.

The relative GS₁ and GS₂ activities from the fractions of the HiTrap Blue HP and MonoQ column are shown in Tab. VII. An aliquot of the GS active fraction of high and low NO₃⁻ - grown cells (after the MonoQ 5/50 GL column) was applied on a 10% SDS-PAGE gel and a protein band at ca. 45 kDa was detected (Fig. 21). Due to the low protein amount of the

Table VII. Purification of GS isoforms in *E. huxleyi* cells grown under high (H) and low (L) NO_3^- (GS₁: cytosolic GS isoform; GS₂: chloroplastic GS isoform).

GS activity was measured according to the GS synthetase assay. Relative GS activity is given in nmol L-Glu γ -monohydroxamate $\text{min}^{-1} \text{mL}^{-1}$. Specific GS activity is given in nmol L-Glu γ -monohydroxamate $\text{min}^{-1} \text{mg protein}^{-1}$.

NO_3^- treatment	Purification step	GS isoforms	Relative GS activity	Specific GS activity	Purification factor
H		Crude extract	951	0.3	1
	Sephacryl S-200 HR	GS ₁	2100	1.3	4.3
		GS ₂	395	0.3	1
	HiTrap Blue HP	GS ₁	913	3	10
		GS ₂	213	-	-
	MonoQ 5/50 GL	GS ₁	89	102	340
		GS ₂	24	-	-
L		Crude extract	1918	0.55	1
	Sephacryl S-200 HR	GS ₁	365.5	0.2	-
		GS ₂	3805	2.7	4.9
	HiTrap Blue HP	GS ₁	304.5	-	-
		GS ₂	2039.5	6.8	12.3
	MonoQ 5/50 GL	GS ₁	6	-	-
		GS ₂	113	-	-

purified GS isoforms, only the GS₁ isoform of high NO₃⁻-grown cells was clearly visualized after silver staining. This gives some evidence that GS₁ protein expression is increased under high NO₃⁻.

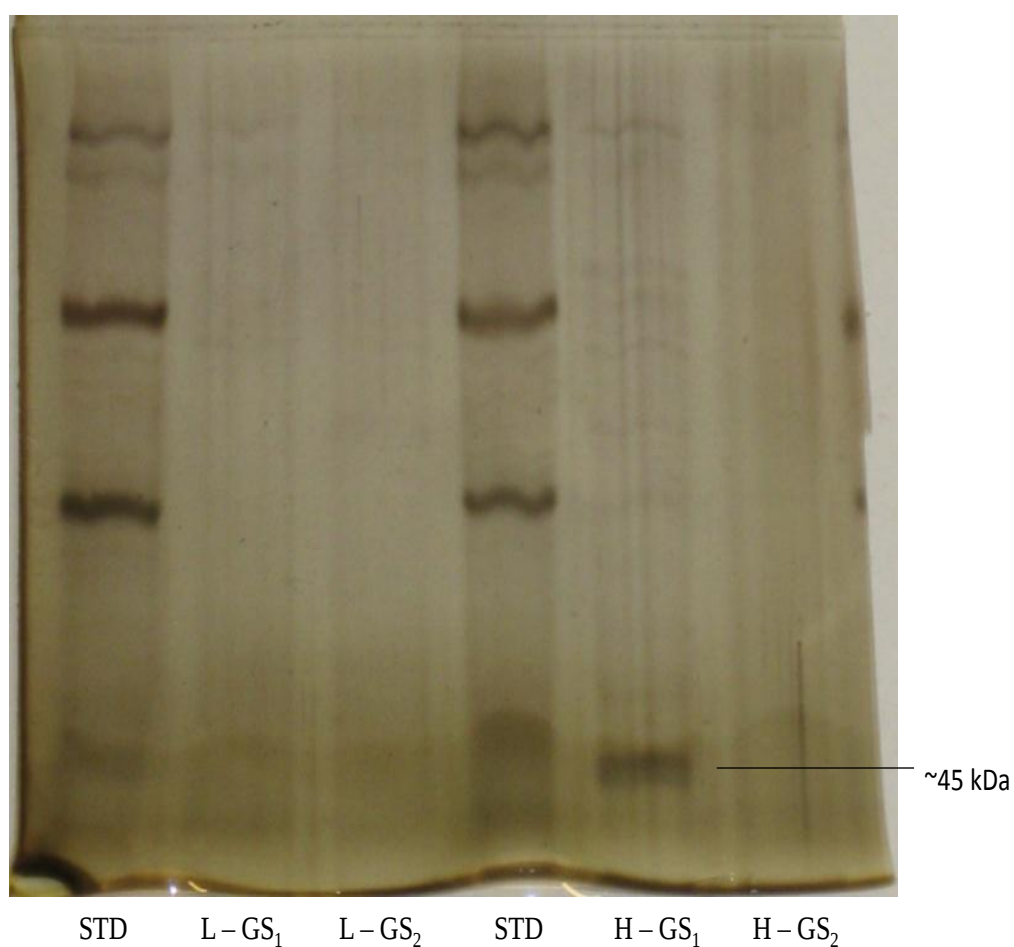


Figure 21. An aliquot of the glutamine synthetase (GS) active fractions after separation on a MonoQ 5/50 GL column was applied on a 10 % SDS-PAGE gel. A band at approximately 45 kDa was visualized after silver staining (Std: protein standard – ovalalbumin: 45 kDa).

3.6. Photosynthesis and respiration

The maximum rate of net photosynthesis ($V_{\max}$) expressed on a per cell basis was about two-fold lower in low NO_3^- -grown cells, with a value of $6.5 \pm 0.8 \text{ fmol O}_2 \text{ min}^{-1} \text{ cell}^{-1}$ compared to $12.9 \pm 0.7 \text{ fmol O}_2 \text{ min}^{-1} \text{ cell}^{-1}$ under high NO_3^- (Tab. VIII; $t = -3.063$, $p = 0.022$; Fig. 22). When net photosynthesis was expressed on a per mg chl α basis, no difference of its rate was observed between high and low NO_3^- -grown cells. The amount of Chl α was about two-fold lower in cells under low NO_3^- (data not shown). However, the $K_{1/2}$ of photosynthesis for CO_2 and DIC was lower in low NO_3^- -grown cells (Fig. 22A and 22B). The $K_{1/2}$ (CO_2) of photosynthesis was $4.1 \pm 0.6 \text{ }\mu\text{M}$ in high and $2.5 \pm 0.6 \text{ }\mu\text{M}$ in low NO_3^- -grown cells (Tab. VIII; $t = -3.158$, $p = 0.019$), and the $K_{1/2}$ (DIC) was $443 \pm 79 \text{ }\mu\text{M}$ in high and $263 \pm 57 \text{ }\mu\text{M}$ in low NO_3^- -grown cells (Tab. VIII; $t = -2.639$, $p = 0.038$). The light curve of net photosynthesis revealed that the α value was similar between the two NO_3^- treatments (1.47 ± 0.21 for high and 1.39 ± 0.06 for low NO_3^- -grown cells; Tab. VIII; Fig. 23). The rate of respiration expressed on a per cell basis was similar between high and low NO_3^- -grown cells (ca. $2 \text{ fmol O}_2 \text{ min}^{-1} \text{ cell}^{-1}$) (Tab. VIII; $t = -1.59$, $p = 0.163$). When respiration was expressed on mg Chl α basis, its rate was higher under low NO_3^- .

Table VIII. Kinetics $V_{\max}$ and $K_{1/2}$ of net photosynthesis, apparent Chl α specific efficiency of light harvesting (α), kinetics of gross CO_2 uptake, net CO_2 uptake and HCO_3^- uptake, as well as respiration in cells grown under high (H) and low (L) NO_3^- .

For net photosynthesis, gross and net CO_2 uptake, HCO_3^- uptake and respiration in cells grown under high (H) NO_3^- (n=5). All other values represent the means of triplicates ($\pm$ SD).

NO_3^- treatment			H	L
Net photosynthesis	$V_{\max}$	$\text{fmol O}_2 \text{ min}^{-1} \text{ cell}^{-1}$	12.9 ± 0.7	6.5 ± 0.8
	$K_{1/2} (\text{CO}_2)$	$\mu\text{mol L}^{-1}$	4.1 ± 0.6	2.5 ± 0.6
	$K_{1/2} (\text{DIC})$	$\mu\text{mol L}^{-1}$	443 ± 79	263 ± 57
α		$\mu\text{mol O}_2 (\text{mg Chl } \alpha)^{-1} \text{ h}^{-1}$ $(\mu\text{mol photon m}^{-2} \text{ s}^{-1})^{-1}$	1.5 ± 0.2	1.4 ± 0.1
Gross CO_2 uptake	$V_{\max}$	$\text{fmol CO}_2 \text{ min}^{-1} \text{ cell}^{-1}$	6.1 ± 1.9	4.5 ± 1.2
	$K_{1/2} (\text{CO}_2)$	$\mu\text{mol L}^{-1}$	3.5 ± 0.5	2.2 ± 0.3
Net CO_2 uptake	$V_{\max}$	$\text{fmol CO}_2 \text{ min}^{-1} \text{ cell}^{-1}$	2 ± 0.4	2.8 ± 0.5
	$K_{1/2} (\text{CO}_2)$	$\mu\text{mol L}^{-1}$	5 ± 1.6	4.4 ± 1.7
HCO_3^- uptake	$V_{\max}$	$\text{fmol HCO}_3^- \text{ min}^{-1} \text{ cell}^{-1}$	8.2 ± 0.6	2.4 ± 0.3
	$K_{1/2} (\text{HCO}_3^-)$	$\mu\text{mol L}^{-1}$	502 ± 133	192 ± 21
Respiration	$V_{\max}$	$\text{fmol O}_2 \text{ min}^{-1} \text{ cell}^{-1}$	2.3 ± 0.4	1.9 ± 0.1

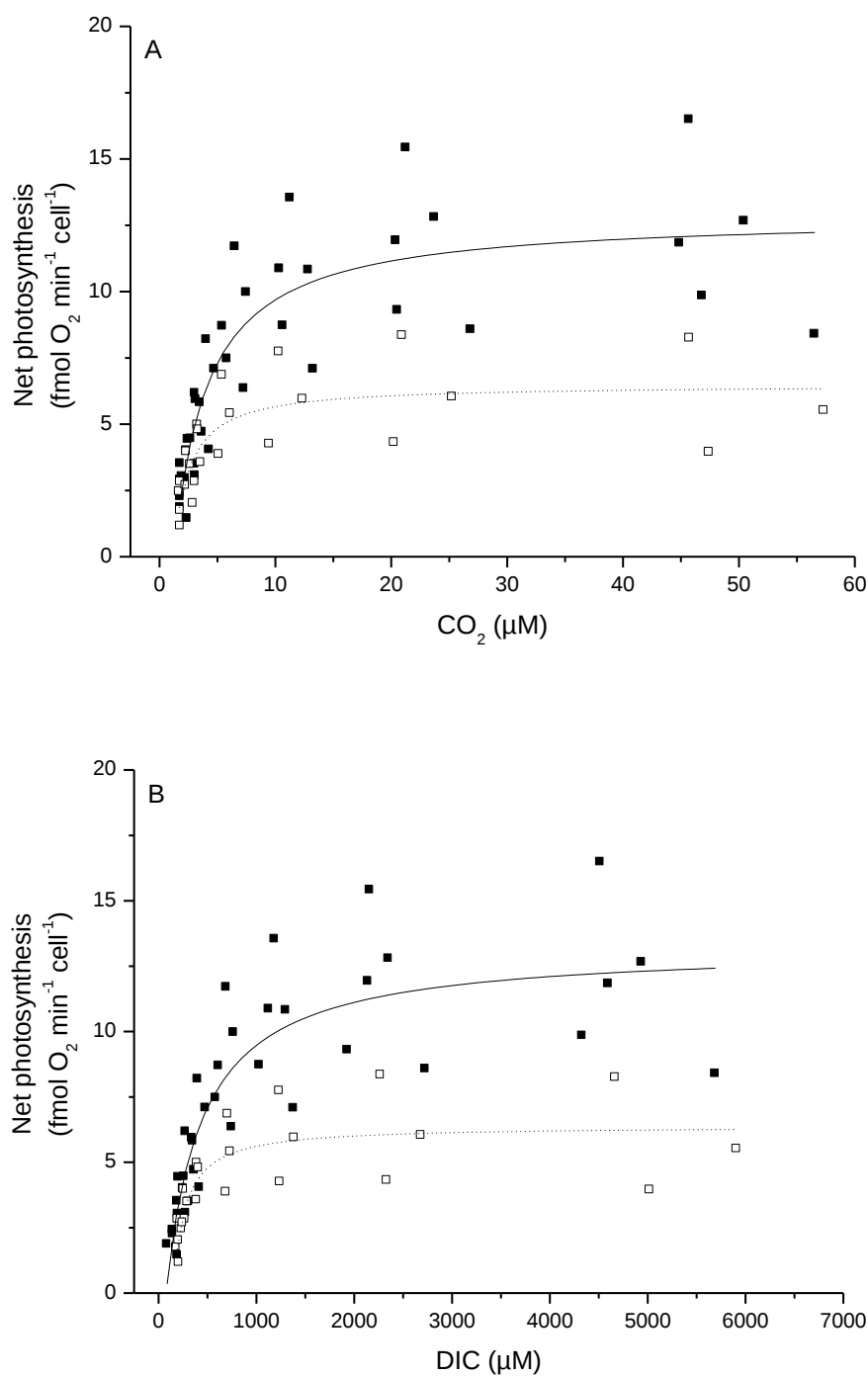


Figure 22. Kinetic curve of net photosynthesis for CO₂ (A) and DIC (B) in cells grown under high (■) and low (□) NO₃⁻. For net photosynthesis in cells grown under high (H) NO₃⁻ (n=5). All other values represent the means of triplicates.

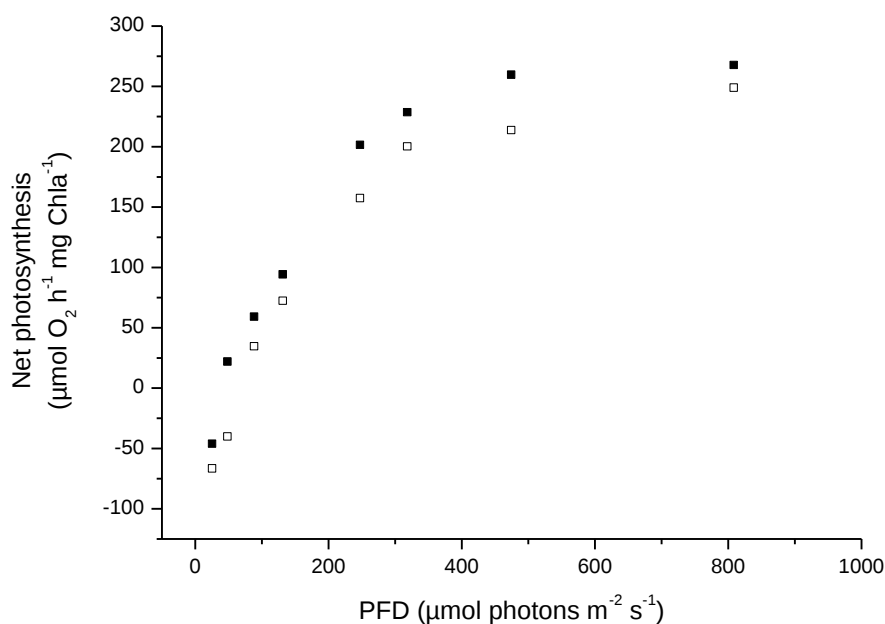


Figure 23. The effect of various light intensities (PFD) on net photosynthesis in cells grown under high (■) and low (□) NO_3^- . All values represent the means of triplicates.

3.7. Inorganic carbon fluxes

The effects of N availability were stronger on the uptake of HCO_3^- (Fig. 224). The rate of HCO_3^- uptake was approximately 4-fold lower under low NO_3^- with a value of $2.4 \pm 0.3 \text{ fmol HCO}_3^- \text{ min}^{-1} \text{ cell}^{-1}$ compared to $8.2 \pm 0.6 \text{ fmol HCO}_3^- \text{ min}^{-1} \text{ cell}^{-1}$ in high NO_3^- -grown cells (Tab. VIII; $t = -5.425$, $p = 0.001$). While the rate of HCO_3^- uptake decreased, the affinity of HCO_3^- uptake increased under low NO_3^- . Low NO_3^- -grown cells had a $K_{1/2}$ for HCO_3^- of $192 \pm 21 \text{ } \mu\text{M}$ compared to $502 \pm 133 \text{ } \mu\text{M}$ in high NO_3^- -grown cells (Tab. VIII, $t = -3.043$, $p = 0.022$). Both, maximum gross and net CO_2 uptake rates (V_{max}) per cell were similar between high and low NO_3^- -grown cells (Fig. 25A and 25B). The V_{max} of gross CO_2 uptake was ca. 4.5 to 6 $\text{fmol CO}_2 \text{ min}^{-1} \text{ cell}^{-1}$ in both high and low NO_3^- -grown cells (Tab. VIII). Net CO_2 uptake V_{max} was approximately 2 to 3 $\text{fmol CO}_2 \text{ min}^{-1} \text{ cell}^{-1}$ in all cells (Tab. VIII). Nevertheless, low

NO_3^- -grown cells had a lower $K_{1/2}$ of gross CO_2 uptake for CO_2 compared to cells under high NO_3^- . The $K_{1/2}$ of gross CO_2 uptake for CO_2 was $3.5 \pm 0.5 \mu\text{M}$ under high and $2.2 \pm 0.3 \mu\text{M}$ under low NO_3^- (Tab. VIII; $t = -3.198$, $p = 0.018$). The $K_{1/2}$ of net CO_2 uptake for CO_2 was about 4 to 5 μM for both high and low NO_3^- -grown cells (Tab. VIII).

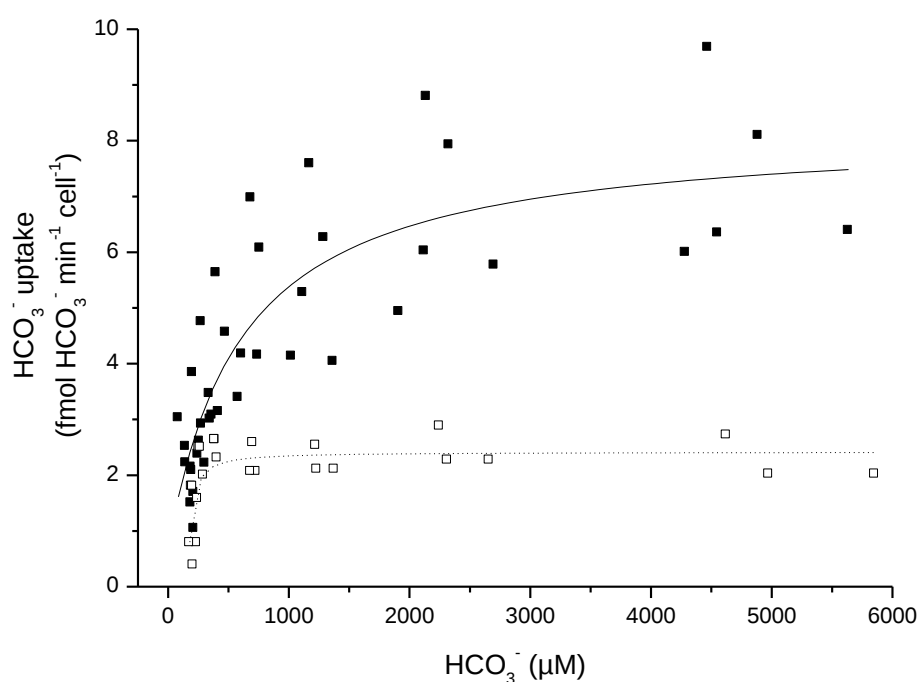


Figure 24. Kinetics of HCO_3^- uptake in cells grown under high (■) and low (□) NO_3^- . For HCO_3^- uptake in cells grown under high (H) NO_3^- ($n=5$). All other values represent the means of triplicates.

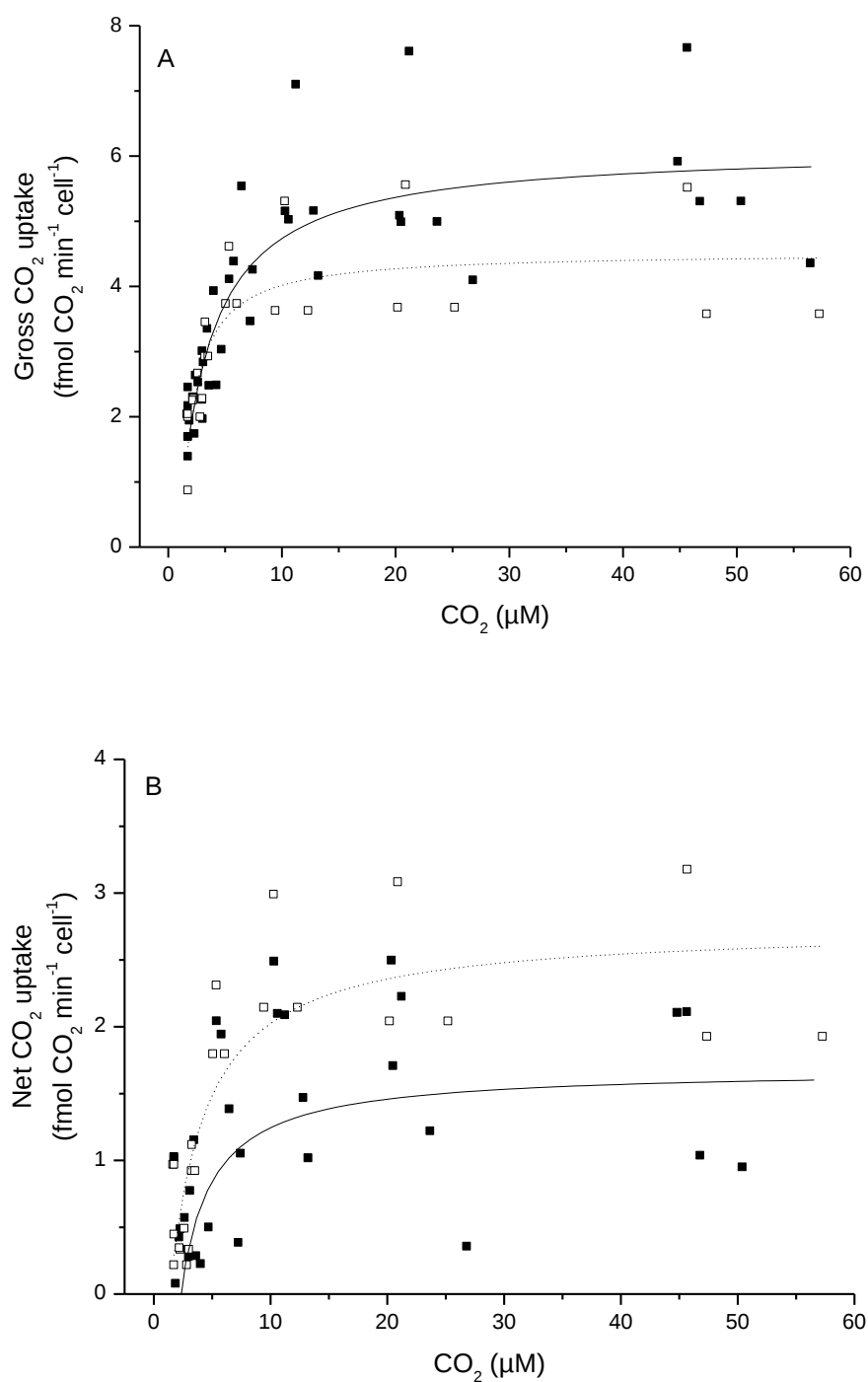


Figure 25. Kinetics of gross CO₂ (A) and net CO₂ (B) uptake in cells grown under high (■) and low (□) NO₃⁻. For gross and net CO₂ uptake in cells grown under high (H) NO₃⁻ (n=5). All other values represent the means of triplicates.

4. Discussion

The aim of the present study was to investigate how cells regulate C and N fluxes and to determine the relative composition of macromolecular pools in response to NO_3^- availability in *E. huxleyi*. Despite of similar growth rates and C to N ratios the allocation of fixed C and N changes with NO_3^- supply.

4.1. Growth parameters

The response of cells to different N availability generally depends on the degree of limitation. The aim of the present study was to impose moderate N limitation in cells of *E. huxleyi* and as a result, growth rates did not change much in response to NO_3^- availability (Tab. IV). However, low NO_3^- -grown cells reduced C as well as N quotas (Tab. IV). As a consequence, C and N production were lower under low NO_3^- (Fig. 9a and 9b). N limitation was also indicated by decreasing volume and protein content per cell under low NO_3^- (Tab. IV). Indeed, microalgae are able to maintain high growth rates during the onset of N limitation (Giordano et al., 2001) and are able to respond to the large variation of N cell quota (Parslow et al., 1984; Borchardt et al., 1996) similar to cells in our study. Maintaining high growth rates during the onset of N limitation may also be among other factors attributing to the capacity of intracellular NO_3^- storage, which occurs in various phytoplankton cells (Dortch 1982; Stolte and Riegmann, 1995; McGlathery et al., 1996). This storage mechanism has not been studied yet for *E. huxleyi* and thus, we cannot make any suggestions about a possible intracellular NO_3^- storage in cells of our study.

The number of visible coccoliths per cell did not change and the percentage of incomplete coccoliths, which contain less calcite than normal coccoliths strongly increased due to N

limitation (Fig. 10; Fig. 11). Thus, we infer that the reduced cellular inorganic C content and inorganic C production were likely caused by a reduced inorganic C content per coccolith. However, a diminished calcite content of coccoliths produced under N limitation was observed in a chemostat-experiment using *E. huxleyi* (Paasche, 1998). The author also noted that coccoliths produced under N limitation were smaller than coccoliths produced under N repletion and he described the morphology of the former as “undercalcified”. These observations are in line with our findings. Paasche (1998) also reported a lower Ca^{2+} content in coccoliths of N limited cells, but unfortunately we did not measure the Ca^{2+} content in coccoliths of cells in our study.

4.2. Macromolecular pools

Nitrate availability had also an impact on the C and N macromolecular pools in *E. huxleyi*. Low NO_3^- -grown cells had a lower ratio of proteins to lipids whereas the ratios of proteins to carbohydrates and lipids to carbohydrates increased compared to cells under high NO_3^- (Fig. 12). Similar results in the ratios of macromolecular pools have been also obtained for N limited cells of the marine diatom *Chaetocheros muellerii* by Giordano et al. (2001).

Giordano et al. (2001) reported that proteins belong to the primary cell compounds, which are reduced due to low NO_3^- availability and ongoing N limitation. Such changes can be explained by the constant growth rates (continuous cell divisions) in combination with the low NO_3^- availability and the decline in cellular components by new biomass production (McGlathery et al., 1996). Among proteins, RubisCO is the most abundant in photosynthetic organisms and the reduction in the amount of RubisCO as well as of its activity is well established in algae under N limitation (Falkowski et al., 1989; Beardall, 1991; Young and Beardall, 2005). Moreover, N limitation was found to cause a rearrangement of intracellular

pools (McGlathery et al., 1996) driving carbon away from protein synthesis and relocating it into lipids (Giormano et al., 2001).

4.3. Nitrate assimilation

Growing either under high or low NO_3^- , cells are able to maintain constant C to N ratio by regulating their N assimilation according to the N availability. Under low NO_3^- , cells had lower NR and NiR $V_{\max}$ compared to those grown under high NO_3^- conditions (Tab. V). This effect is attributed to the decrease of cell volume and cell protein under low NO_3^- because NR and NiR $V_{\max}$ are quite similar in the two NO_3^- treatments when enzyme activities are expressed on a per cell volume or cell protein basis. Since the genes coding for NR and NiR are located in the same gene cluster as has been previously reported for *C. reinhardtii* (Quesada et al, 1998), low NO_3^- should lead to lower expression of NR and NiR mRNA in *E. huxleyi* cells thereby decreasing the NR and NiR protein contents and activities. We did not observe any significant modification in the K_m (NADH) of NR for this substrate under high or low NO_3^- (Tab. V). The findings make sense because NADH is not expected to become a limiting factor at high light intensities under which cells of both NO_3^- treatments were acclimated. Nevertheless, the affinities of NR for NO_3^- and NiR for NO_2^- are higher under low NO_3^- (Tab. V) indicating a more efficient reduction of NO_3^- to NH_4^+ in these cells. The cellular mechanism(s) responsible for the increase of the catalytic efficiency of NR and NiR under low NO_3^- is (are) not known. 14-3-3 proteins may be involved in his type of regulation, since they were found to play a role in regulating protein activity and turnover rate (Korkhout et al., 1994; Oeckling et al., 1997; Weiner and Kaiser, 1999). Whether 14-3-3 proteins are involved in regulating the catalytic efficiencies of NR and NiR in *E. huxleyi* and whether they

are responsible for higher affinities of NO_3^- and NO_2^- reduction under low NO_3^- or not have still to be tested.

Maintaining constant C to N ratio under high and low NO_3^- also includes the regulation of the enzymes in the GS/GOGAT cycle where C and N assimilation meet. Irrespective of the type of normalization, GS $V_{\max}$ was found to be higher in low NO_3^- -grown cells (Tab. VI). This increase in GS $V_{\max}$ in cells under low NO_3^- might be due to GS₂ activity, which was found to be higher than GS₁ activity in these cells (Fig. 18b). The amount of the GS₂ isoform was not affected by NO_3^- availability in cells of *E. huxleyi* despite the difference in GS₂ specific activity between the different NO_3^- treatments (Fig. 19; Tab. VII). This finding agrees with the study of Chen and Silflow (1996), who reported that the level of GS₂ transcripts was unaffected by NO_3^- availability in *C. reinhardtii*. The same authors also showed that the level of GS₁ transcripts in *C. reinhardtii* is induced by NO_3^- . This may explain the lower amount and activity of the GS₁ isoform in cells at 10 μM NO_3^- compared with cells grown at 280 μM NO_3^- (Fig. 20, Fig. 21, Tab. VII). The increase of GS activity in microalgae has been attributed to light activation due to reduction of the enzyme by reduced thioredoxin but also to de novo biosynthesis of GS protein during the light period (Tischner and Hüttermann, 1980; Schmidt, 1981; Tischner and Schmidt, 1982; Florencio et al., 1993). Besides, the increase in GS $V_{\max}$, GS affinity for Glu was higher in low NO_3^- -grown cells whereas enzyme affinities for ATP and NH_2OH remained similar in cells of the two NO_3^- treatments (Tab. VI). This difference in the affinity for Glu may be related to the proportion and the affinity for Glu of each of the GS isoform in the tested NO_3^- concentrations. Unfortunately we could not conduct a kinetic analysis on the two isoforms due to low cell densities in order to evaluate the affinity of GS₁ and GS₂ for Glu. 14-4-3 proteins could also play a role in the regulation of GS

catalytic activity (Moorhead et al., 1999; Finnemann and Schjoerring, 2000), but this type of regulation has not been tested yet in *E. huxleyi*.

As expected, GOGAT activity showed more or less the same response with GS activity to NO_3^- availability since both enzymes are part of the same cycle (Tab. VI). As in the case of total GS, the higher activity of GOGAT at $10 \mu\text{M NO}_3^-$ may be due to regulation in the protein level by thioredoxin (Tischner and Schmidt, 1982). The lower PEPCK activity observed at $10 \mu\text{M NO}_3^-$ (Tab. VI) likely reflects the need of these cells to match the flow of C into amino acid synthesis with the flow of N under these conditions.

4.4. Photosynthesis and respiration

Net photosynthesis per cell and Chl α decreased in cells under low NO_3^- (Tab. VIII). When expressed per cell chlorophyll, photosynthetic rates were similar in the two NO_3^- treatments. While low NO_3^- availability decreases the amount of chlorophyll and the rate of photosynthesis per cell, higher affinities of photosynthesis for CO_2 and DIC were measured in low NO_3^- -grown cells (Tab. VIII). Lower rates of photosynthesis may be related primarily to the composition of PSI and PSII in terms of integral proteins and pigments, and secondly to the abundance and regulation of proteins, which receive most of the reductant (NADPH) and ATP produced by photosynthesis such as Rubisco. Indeed, N limitation leads to increased ratios of carotenoids to Chl α , causes a reduction in thylakoid stacking and absorptivity and decreases the amount of Rubisco per cell in algae (Plumley et al., 1989; Falkowski et al., 1989; Beardall et al., 1991). Furthermore, despite of the onset of N limitation the apparent chlorophyll-specific efficiency of light harvesting (α) in low NO_3^- -grown cells was similar with the one under high NO_3^- (Tab. VIII). Previous experiments showed that N limitation may

increase, decrease or maintain α (Osborne and Geider, 1986; Kolber et al., 1988; Herzig and Falkowski, 1989) depending on the investigated species as well as the degree of N limitation.

In contrast to photosynthesis, the rate of respiration per cell was similar between high and low NO_3^- -grown cells (Tab. VIII), but was higher in low NO_3^- -grown cells when expressed on cell chlorophyll. Under low NO_3^- , respiration may compensate for the lower amount of reductant and ATP provided by photosynthesis in these cells. Weber and Turpin (1989) reported that the reductant deriving from photosynthesis is not sufficient enough to meet the demands of NO_3^- and NO_2^- reduction in N-limited cells of *S. minutum* and this deficiency in reducing power is compensated by TCA cycle reductant. The measured respiration rates in low NO_3^- -grown cells may also be related to higher rates of N assimilation in the dark compared to cells under high NO_3^- . It has been shown that respiration increases in the dark, in order to support N assimilation in the green alga *S. minutum* (Vanlerberghe et al., 1992).

4.5. Inorganic carbon uptake

Cells of *E. huxleyi* change their preference of inorganic C species with respect to N availability. At low NO_3^- , cells down regulated HCO_3^- uptake compared to cells grown at high NO_3^- (Tab. VIII), probably as a result of less HCO_3^- transporters in these cells due to the lower NO_3^- availability. The increase in the affinity of HCO_3^- transport in cells at low NO_3^- may be related to a direct interaction of 14-3-3 proteins with the HCO_3^- transporters, or to an indirect interaction of 14-3-3 proteins with the H^+ -ATPase (Korkhout et al., 1994; Oeckling et al., 1997), which is essential for proton transport across membranes. Despite the smaller size of cells at low NO_3^- , gross CO_2 uptake (uptake of CO_2 into the cell) and net CO_2 uptake (gross CO_2 uptake minus CO_2 efflux) were similar in cells of both NO_3^- treatments (Tab. VIII). This

response in CO₂ uptake, in combination to the higher affinity for CO₂ in cells at low NO₃⁻ (Tab. III), can be related to the regulation of proteins, which utilize CO₂ such as RubisCO and/or carbonic anhydrase (CA). Again, phosphorylation and interaction with 14-3-3 proteins may be the reason for higher affinities for CO₂ in cells grown at low NO₃⁻.

4.6. Intracellular concentrations of nitrate and carbon dioxide

Based on the enzyme kinetics measured for NR and the kinetics of net photosynthesis derived from the MIMS measurements, we estimated the intracellular concentrations of NO₃⁻ and CO₂ which allows for balanced growth, i.e. maintenance of a constant C to N ratio. Under these circumstances, the net fixation rate of C (C_{flux}) can thus be expressed in terms of the N assimilation rate (N_{flux}), using the C to N ratio:

$$\frac{C_{\text{flux}}}{N_{\text{flux}}} = \frac{\text{POC}}{\text{PON}} \quad (1)$$

The upper boundary for C_{flux} in balance with N assimilation is provided by the maximum NR activity (V_{max}^{NR}) and would be obtained at infinite NO₃⁻ concentration in the cytosol. Using our result for V_{max}^{NR}, the upper limit for C_{flux} that derives from equation (1) is very similar to the maximum net photosynthesis (in terms of C), which we obtained from the MIMS measurements (V_{max}^{CO₂} = V_{max}^{O₂}/PQ, with PQ being the photosynthetic quotient). For balanced growth to be maintained (equation 1), the upper limit for C_{flux} must assume a value of 9.3 fmol min⁻¹ cell⁻¹ for the high NO₃⁻ condition (POC/PON = 8.5), and 5.3 fmol min⁻¹ cell⁻¹ for cells acclimated to the low NO₃⁻ concentration (POC/PON = 8.3). Given the photosynthetic quotient from a NO₃⁻-utilizing cell (PQ=1.36; Williams and Robertson, 1991), the MIMS measurements predict a V_{max}^{CO₂} of 9.5 ± 0.5 fmol min⁻¹ cell⁻¹ and 4.8 ± 0.6 fmol min⁻¹ cell⁻¹, for

the high and low NO_3^- conditions, respectively. Hence, the ratios of $V_{\max}^{\text{CO}_2}$ and $V_{\max}^{\text{NR}}$ are approximately given by the C to N ratios for the two growth NO_3^- concentrations used for this study (high NO_3^- : 1.0 POC/PON, low NO_3^- : 0.9 POC/PON). Using equation (1) and $V_{\max}^{\text{CO}_2} / V_{\max}^{\text{NR}} \approx \text{POC/PON}$ it follows that

$$\frac{C_{\text{flux}}}{V_{\max}^{\text{CO}_2}} \approx \frac{N_{\text{flux}}}{V_{\max}^{\text{NR}}} \quad (2)$$

Finally, we assumed for the C_{flux} the enzyme activity of RubisCO and for the N_{flux} the enzyme activity of NR. Then, equation (2) yields the condition for balanced growth in terms of the intracellular concentrations of NO_3^- and CO_2

$$\frac{\text{CO}_2}{K_{\text{Rub}}} \approx \frac{[\text{NO}_3^-]}{K_{\text{NR}}} \quad (3)$$

with K_{Rub} and K_{NR} being the Michaelis-Menten constant for RubisCO and NR, respectively. Generally, for an enzyme possessing the activity V_0 , the ratio of the concentration of the substrate S and the Michaelis-Menten constant K_m determines the degree of saturation of the enzyme activity ($V_0/V_{\max} = 1/(K_m/[S] + 1)$). Based on equation (3) it follows that a constant C to N ratio can be maintained for intracellular concentrations of CO_2 and NO_3^- which yield a comparable saturation of the enzyme activity for the enzymes involved in C and N assimilation. Hence, the maintenance of a constant C to N ratio requires a concerted regulation of the intracellular $[\text{CO}_2]$ and $[\text{NO}_3^-]$ to equilibrate the fluxes through the C and N assimilation pathway. It should be noticed that equation (3) is valid for all degrees of enzyme saturation ($0 < V_0/V_{\max} < 1$) in the cells acclimated to the two growth regimes. For instance, when the NR activity reaches 80% of its maximum value, NO_3^- accumulates in the cytosol, yielding concentrations of 396 μM and 296 μM for the cells acclimated to high and low NO_3^- , respectively. For balanced growth to be maintained (equation (3)), the CO_2 concentration at

the site of RubisCO must then exceed K_{Rub} by a factor of 4, for the high and low NO_3^- conditions. The values of K_{Rub} for the two growth conditions are not known. Assuming a K_{Rub} of 30 μM (Badger et al., 1998), it follows a $[\text{CO}_2]$ at RubisCO of 120 μM . We assumed steady state conditions for N assimilation. In other words, the substrates inside the cell accumulate until the rates of reactions are the same for all the enzymes involved in the N assimilation pathway (from NO_3^- to NH_4^+ assimilation by GS):

$$N_{\text{flux}} = \frac{V_{\text{max}}^{\text{NR}} [\text{NO}_3^-]}{K_{\text{NR}} + [\text{NO}_3^-]} = \frac{V_{\text{max}}^{\text{NiR}} [\text{NO}_2^-]}{K_{\text{NiR}} + [\text{NO}_2^-]} = \frac{V_{\text{max}}^{\text{GS}} [\text{NH}_4^+]}{K_{\text{GS}} + [\text{NH}_4^+]} \quad (4)$$

Then, the NiR kinetics data predicts an accumulation of high NO_2^- concentrations at the site of NiR activity. Similar K_m values of NiR for NO_2^- have been also reported for the flagellate *Prorocentrum minutum* and internal NO_2^- pools were 4 to 5 mM in the diatom *Chaetocheros* sp. (Lomas and Glibert, 2000). The requirement for a high NO_2^- concentration for *E. huxleyi* NiR is suggestive of the fact that this enzyme is confined in the proximity of the chloroplast envelope, possibly near the entry point of NO_2^- into the chloroplast. The V_{max} of NiR and GS are probably underestimated due to suboptimal conditions caused by the sample preparation and the pH optimum of the enzymes. GS activity was found to be ca. 40% higher when measured with a radioactive assay with ammonia as substrate in comparison to the colorimetric assay with NH_2OH as substrate (Listrom et al., 1997) and therefore cannot be used to estimate intracellular NH_4^+ concentrations.

4.7. Conclusions

In conclusion, NO_3^- availability seems to have a crucial impact on the regulation of C and N fluxes in the marine coccolithophore *E. huxleyi*, without necessarily changing the C to N ratios and/or growth rates of cells. With the exception of GS and GOGAT, cells generally respond to low NO_3^- availability with a down-regulation of activities of proteins involved in the acquisition and assimilation of inorganic C and N. Simultaneously, the substrate affinities of these proteins increase in response to low NO_3^- . This study indicates that the intracellular CO_2 and NO_3^- concentrations near RubisCO and NR have to be carefully regulated in order to equilibrate the fluxes through the C and N assimilation pathways, thus ensuring adequate growth under the respective NO_3^- conditions. Decreasing NO_3^- availability in the future ocean due to increased stratification of the upper water masses will decrease marine primary production and microalgae cells, such as the coccolithophore *E. huxleyi*, by reduction of individual size. Nevertheless, NO_3^- availability does not affect the PIC to POC ratio in *E. huxleyi* cells, as long as growth rates and C to N ratios remain constant, and have putatively no significant effect in the biologically mediated ocean-atmosphere CO_2 exchange.

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Own contributions

The PhD thesis with the title was conducted within three years from the 1st of February 2007 till the 31th of January 2010. In the first year I did several test-studies in order to identify the appropriate growth conditions for the phytoplankton species *Emiliana huxleyi* as well as the substrate concentrations needed for the kinetic measurements of enzyme and uptake kinetics. The actual experiment was run throughout the whole second year. During this period I took the samples and I conducted all the measurements stated in this thesis. The third year was dedicated into data processing and writing the manuscript for publishing, which has been submitted in the journal Plant Physiology.

I measured the concentrations of dissolved inorganic carbon and alkalinity in the alkalinity labour of the Biogeosciences group in the Alfred-Wegener-Institute. I took and prepared the samples for the elemental composition of *E. huxleyi* and they were measured in the Alfred-Wegener-Institute by the chemical engineer responsible for the mass spectrometer Mr. Klaus-Uwe Richter. I measured the protein amounts of algae samples with the Bradford method in the Tecan labour of the Cell biology and Toxicology group in the Alfred-Wegener-Institute. I prepared the samples for the determination of the macromolecular pools and they were sent to Italy to be measured with Fourier Transform Infrared Spectroscopy by Dr. Alessandra Norici from the University of Ancona. I prepared the samples for the assessment of coccolith morphology and Dr. Gerald Langer made the pictures with a scanning electron

microscope. I conducted the measurements for the nitrate assimilating enzymes in the Alfred-Wegener-Institute under the guidance of Prof. Dr. Angela Koehler. These measurements were done in the Tecan labour of the Cell biology and Toxicology group. Prof. Dr. Mario Giordano from the University of Ancona provided some standard protocols for enzyme measurements and helped in discussing the results of the experiment. The purification of the glutamine synthetase isoforms was conducted by high performance liquid chromatography with the help of Mrs. Ellen Lichte from the Ecological Chemistry group in the Alfred-Wegener-Institute. I made the measurements of photosynthesis, respiration and inorganic carbon acquisition in the Alfred-Wegener-Institute under the guidance of Dr. Scarlett Trimborn and Dr. Bjoern Rost. These measurements were done in the membrane inlet mass spectrometer labour of the Biogeosciences group. Dr. Silke Thoms is the project leader, and she contributed significant in this work with her help in discussing the data and writing the manuscript.

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