

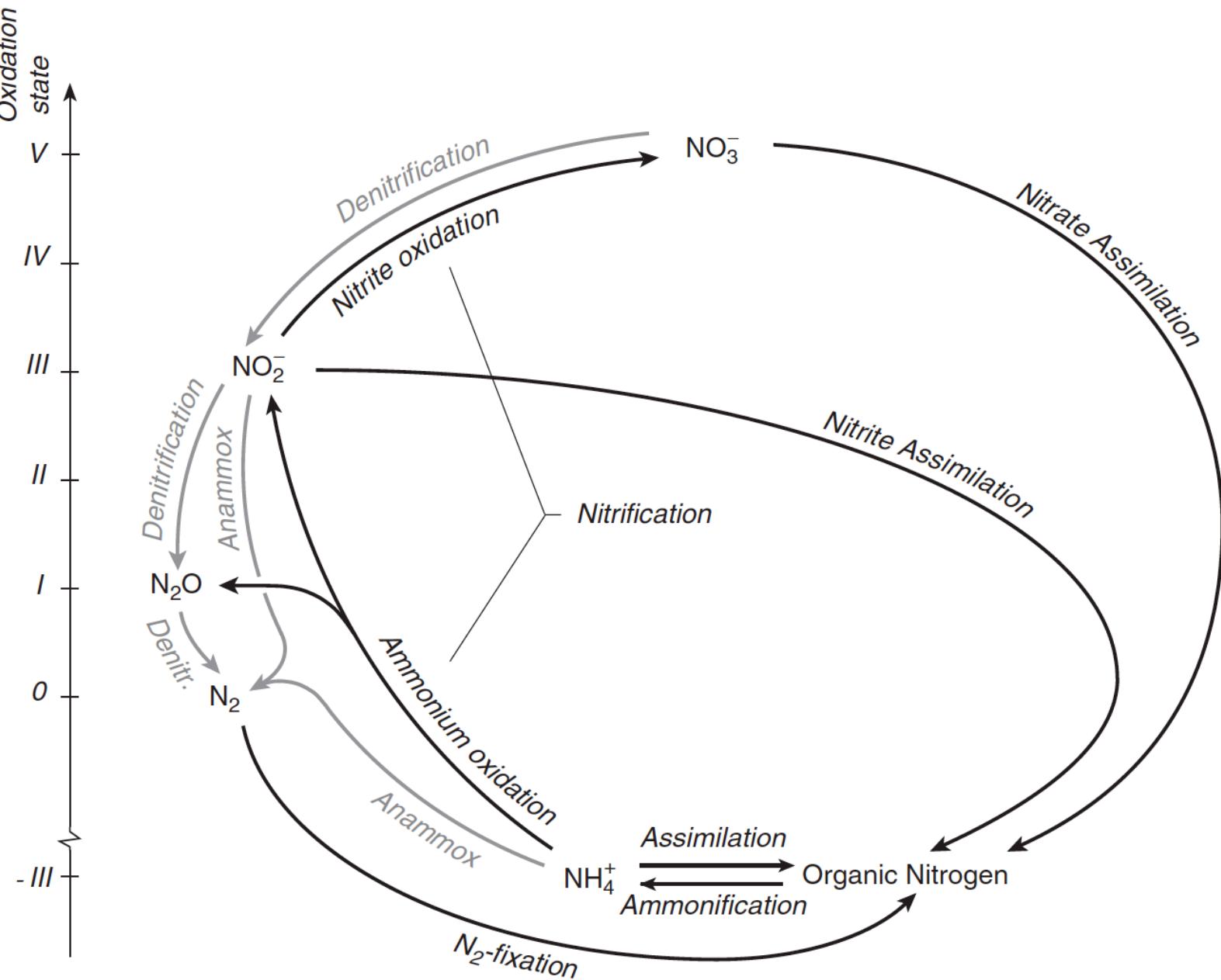
The Microbial Marine Nitrogen Cycle

Alyson Santoro

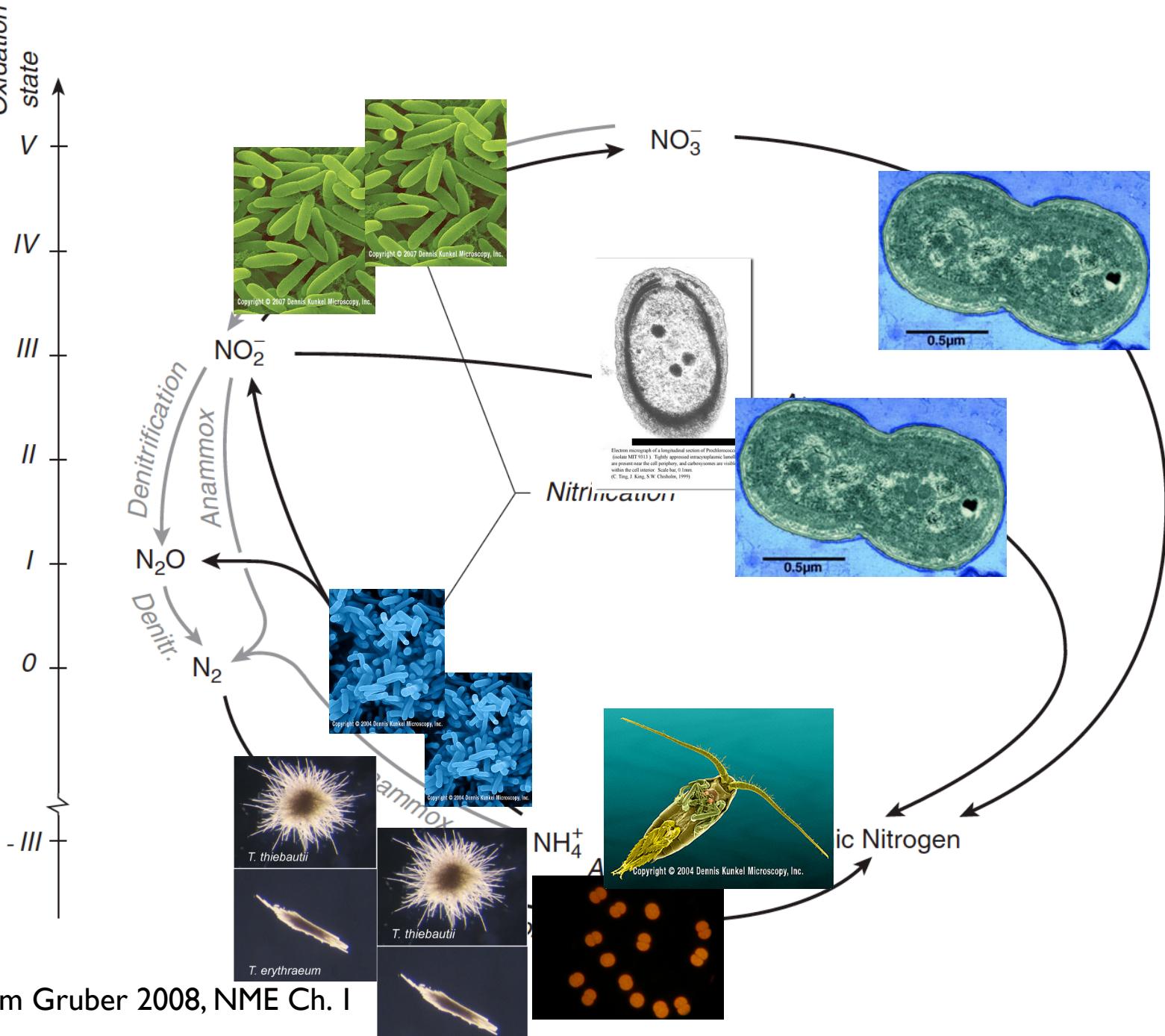
Horn Point Laboratory

University of Maryland Center for Environmental
Science

C-MORE Microbial Oceanography Course 2014



from Gruber 2008, NME Ch. I



from Gruber 2008, NME Ch. I

Outline

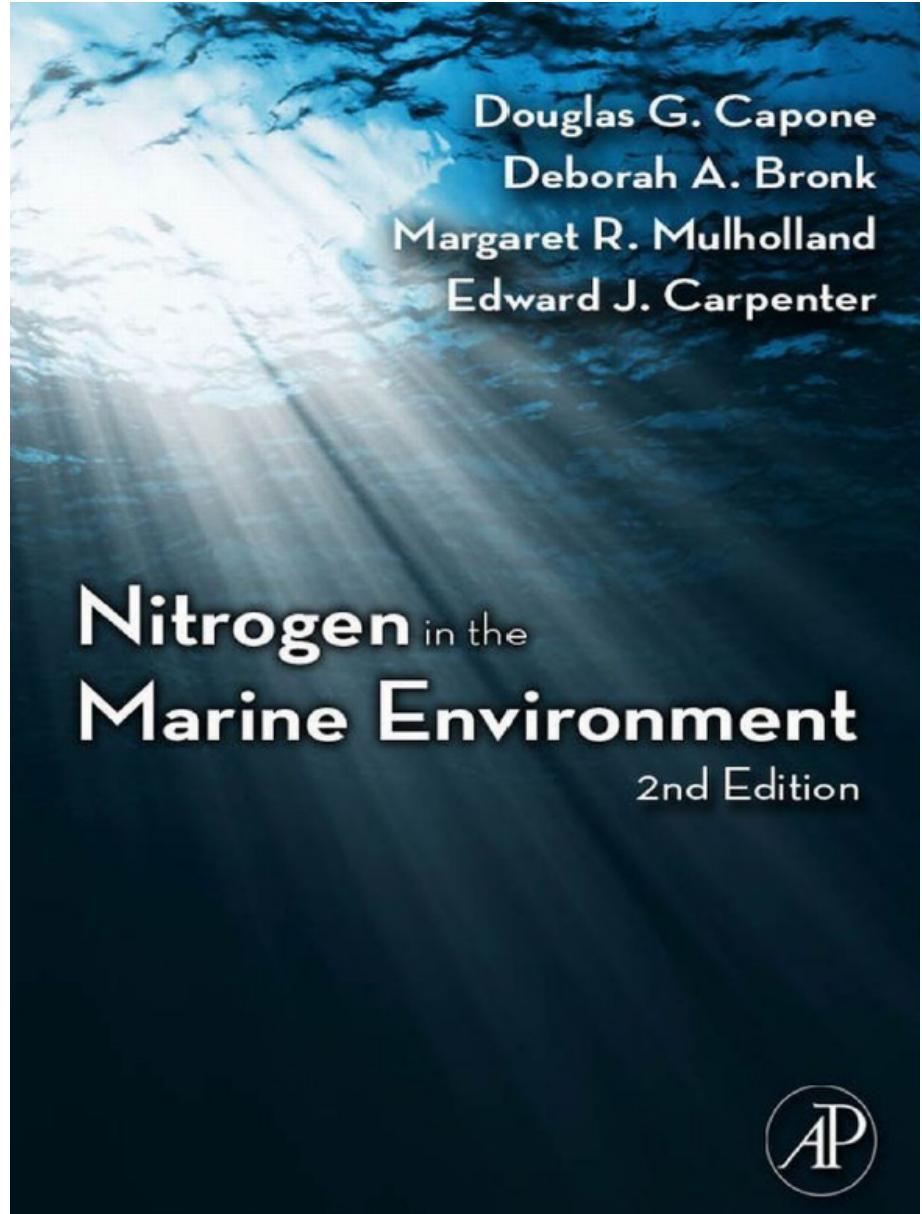
- Overview of the marine nitrogen cycle - pools, depth distributions, and fluxes
- The cast of characters - select functional groups within the microbial nitrogen cycle
- Synthetic studies of the marine microbial nitrogen cycle *in situ* - case studies of oxygen minimum zones and the primary nitrite maximum

Key Ideas:

- Nitrogen cycle is complex because N has so many redox states.
- Marine nitrogen cycle is mainly mediated by biological reactions (microbes) with unique and varying ecologies and evolutionary histories.
- The nitrogen cycle has been a showcase for integrating biogeochemistry and molecular biology, greatly aided by the suite of available ‘functional genes’ and the existence of stable N isotopes.

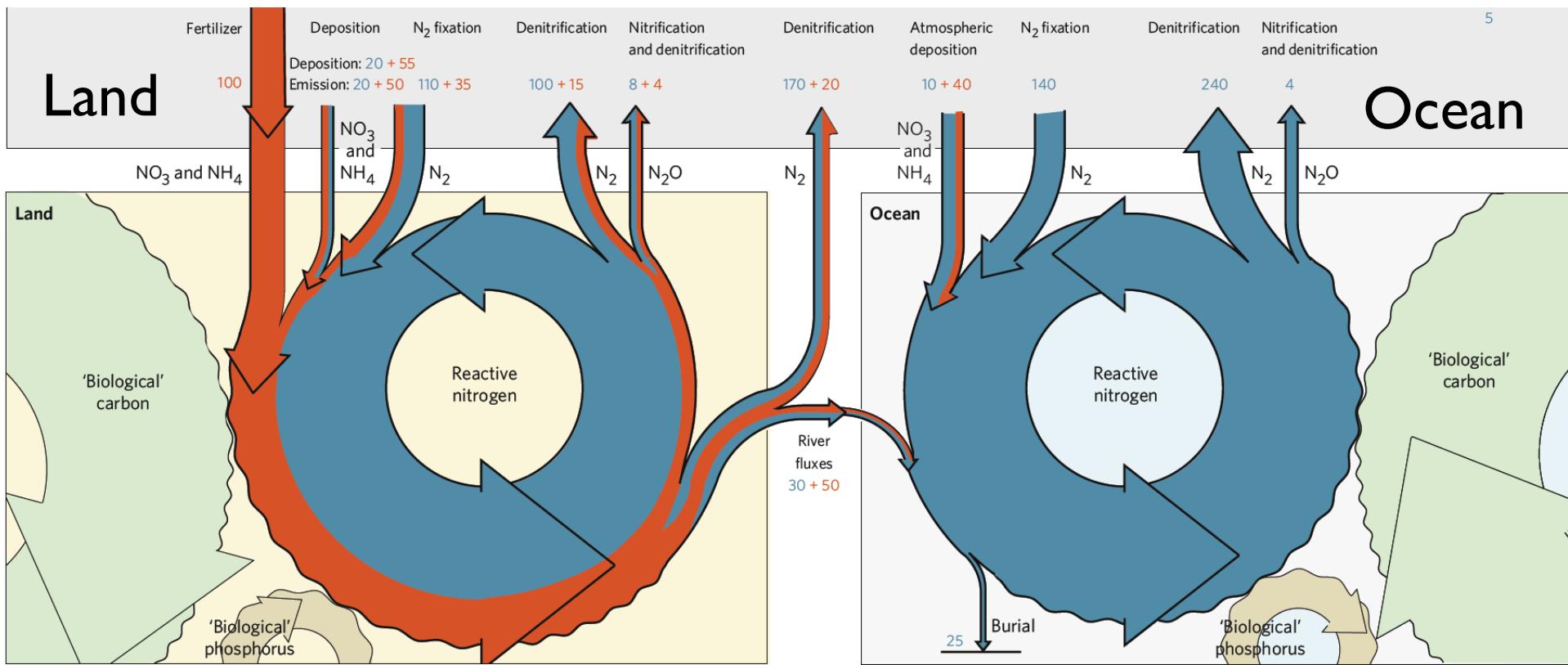
Some bedtime reading. . .

Everything you
ever wanted to
know about the
marine nitrogen
cycle, plus 36 more
chapters.



We are altering the global N cycle

5



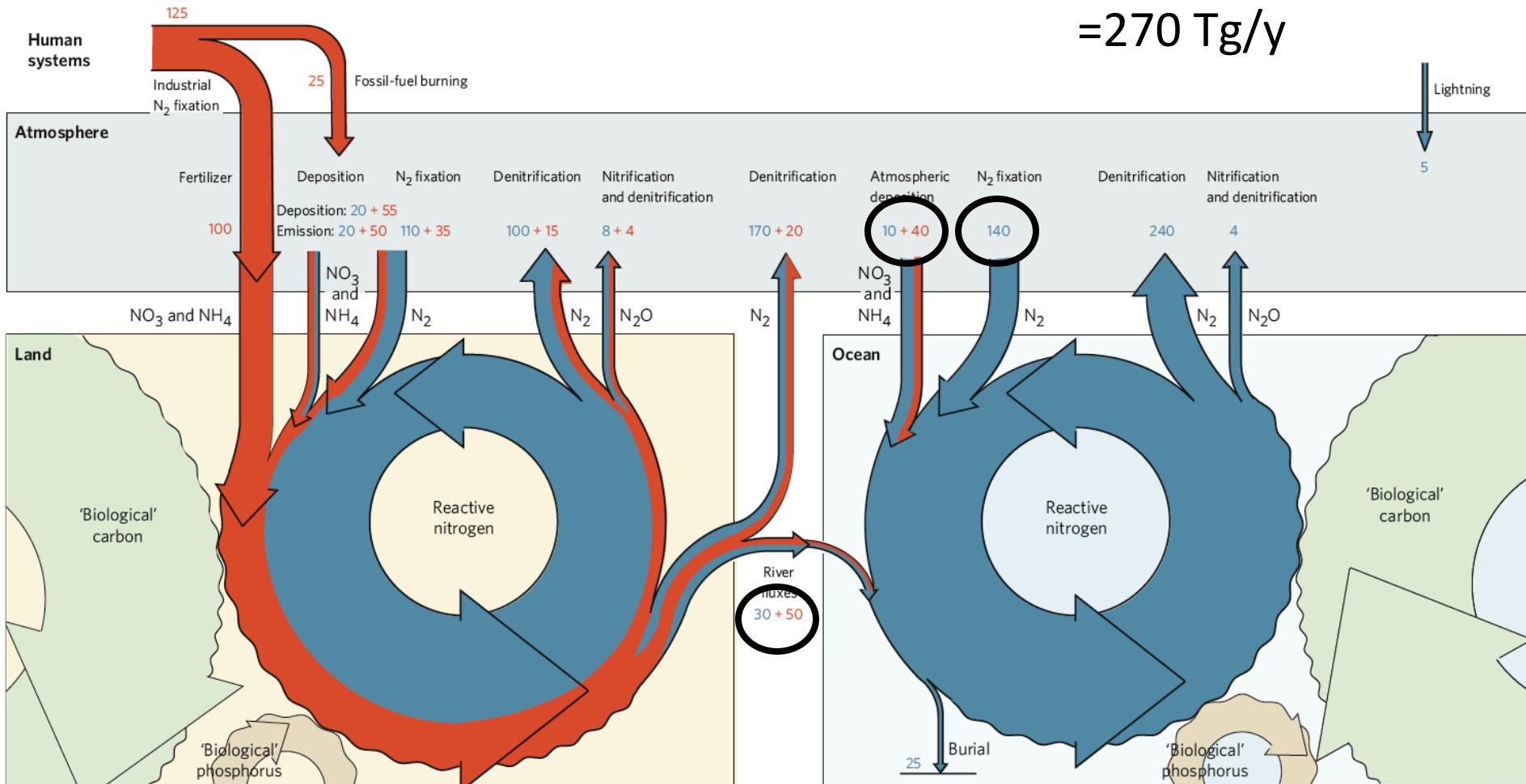
“Few of these flux estimates are known to better than 20%, and many have uncertainties of 50% or larger.”

Gruber and Galloway 2008

Input terms (Tg/yr)

Biological N ₂ fixation	140
Atm. Dep.	50
Runoff	80

$$=270 \text{ Tg/y}$$



Gruber and Galloway, 2008

Output terms (Tg/y)

Denitrification: N₂ loss

240

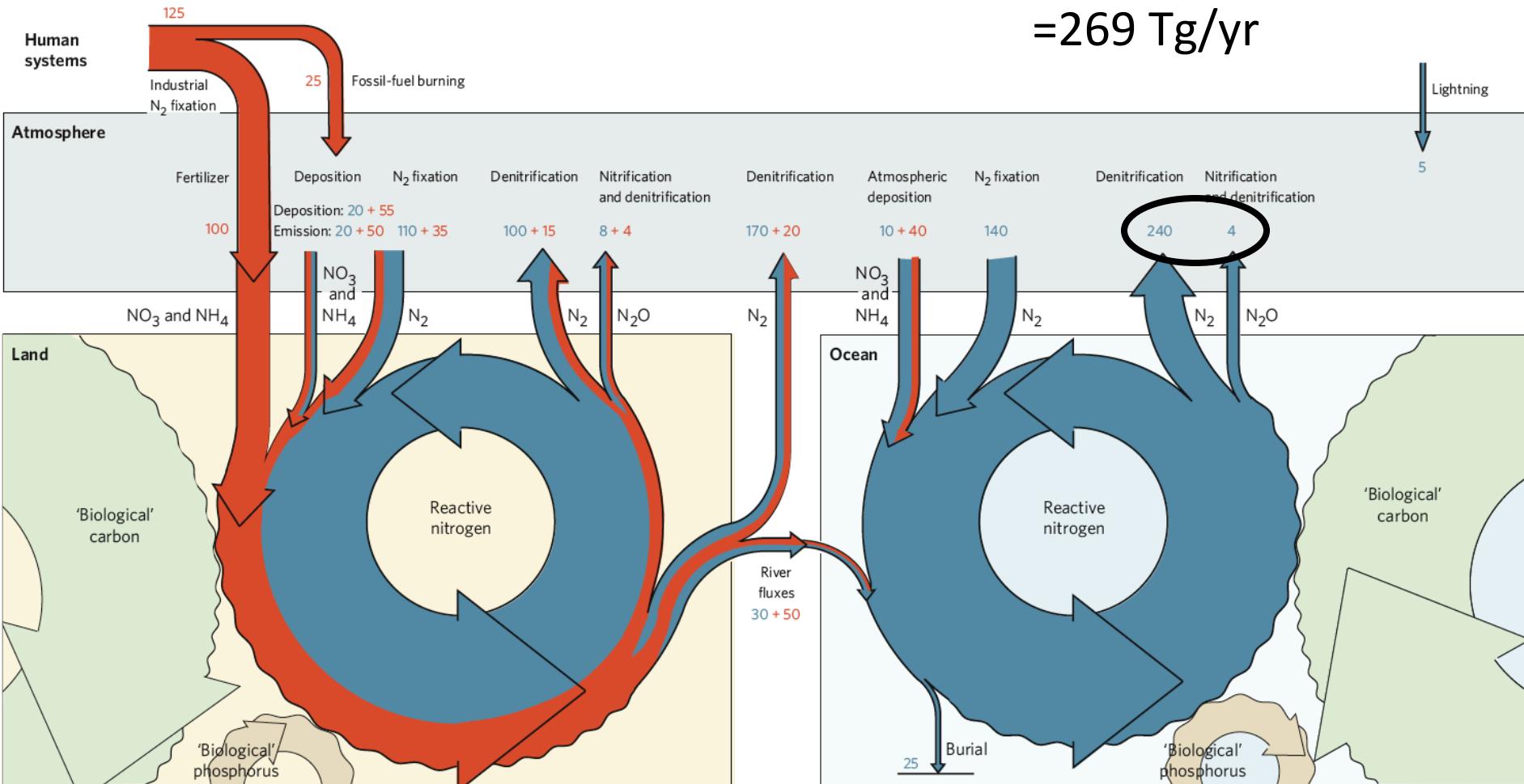
N₂O loss

4

Sediment Burial

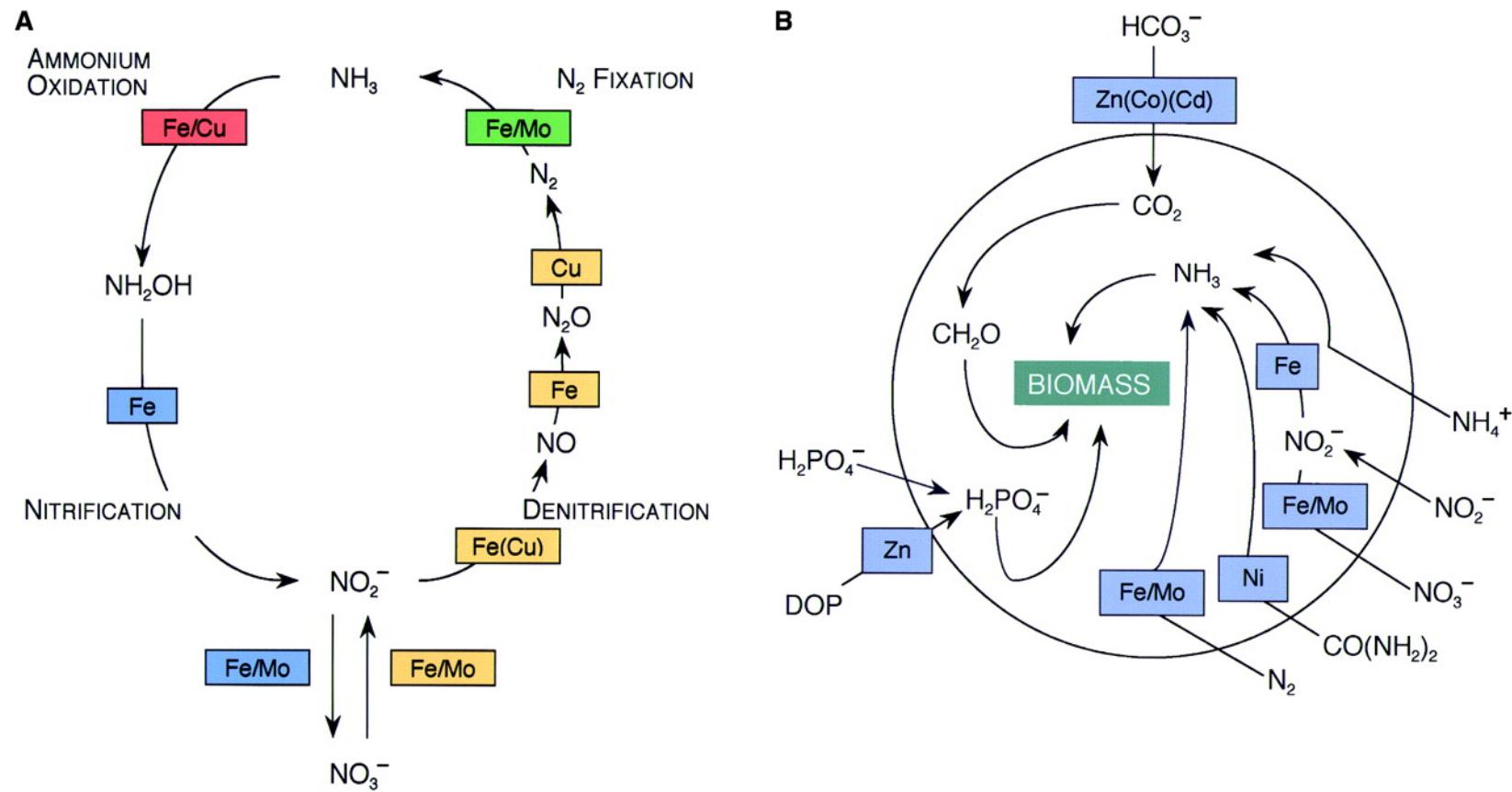
25

$$=269 \text{ Tg/yr}$$



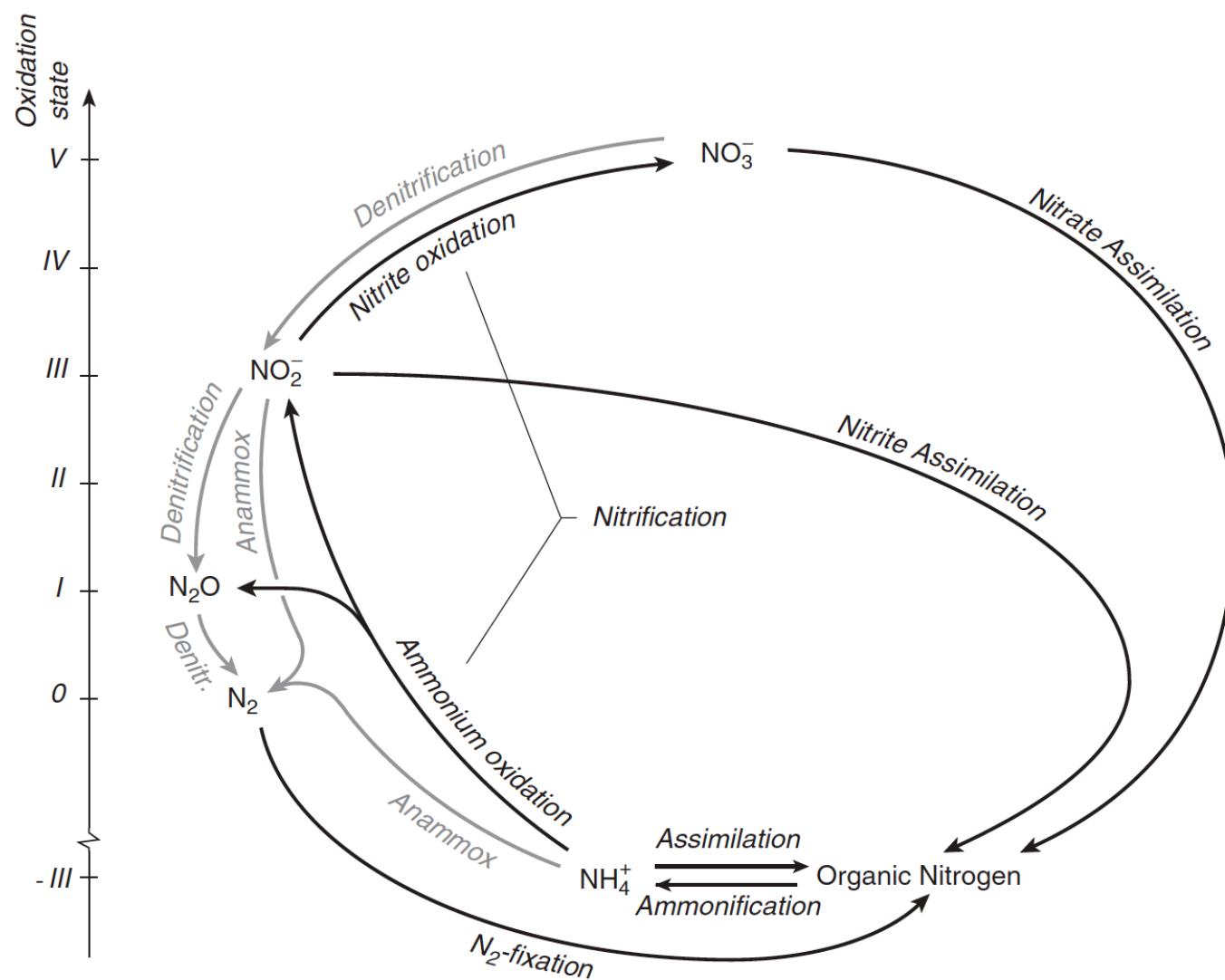
Gruber and Galloway, 2008

The nitrogen cycle is also inextricably linked to trace metal cycling



Morel and Price 2003, *Science*

Nitrogen exists in multiple redox states

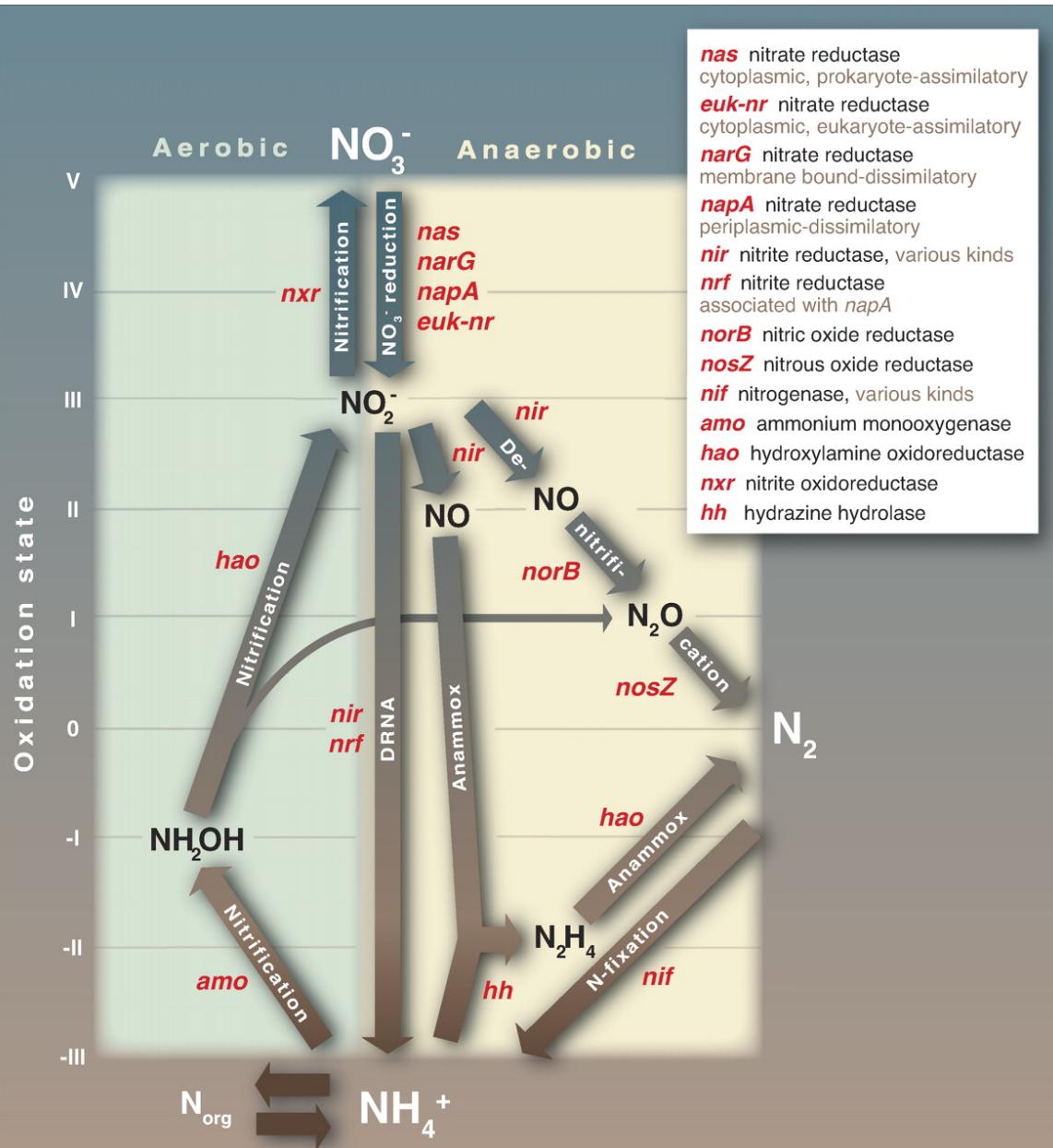


from Gruber 2008

gray = reactions in low/no oxygen environments, black = oxic

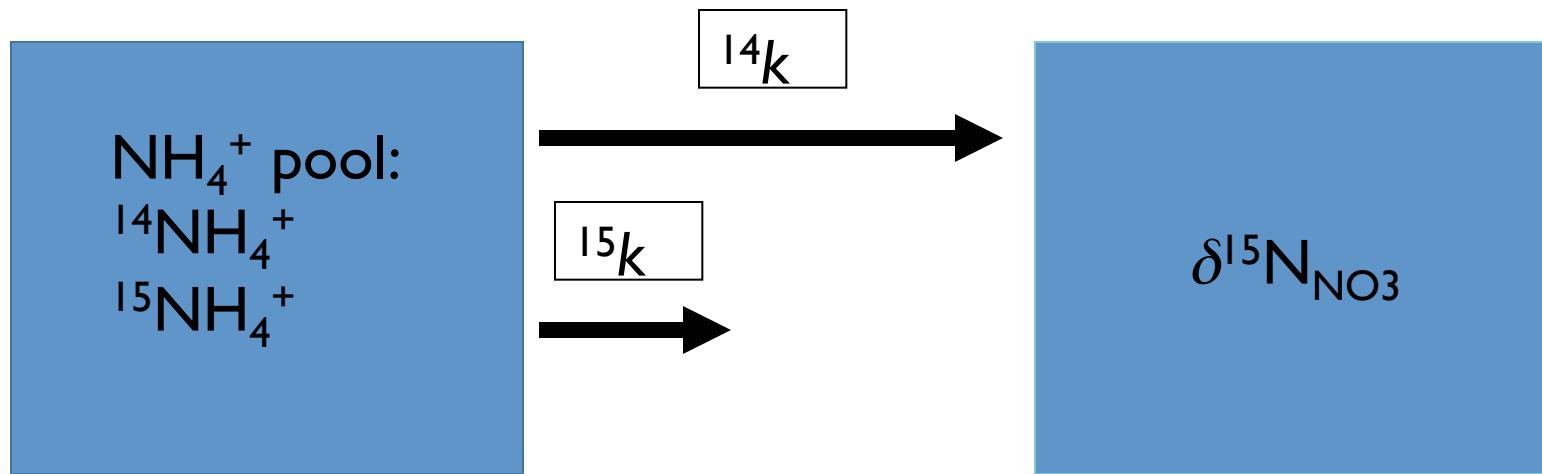
Many ‘functional genes’ available to study the nitrogen cycle

But, always remember that
genes come from cells.



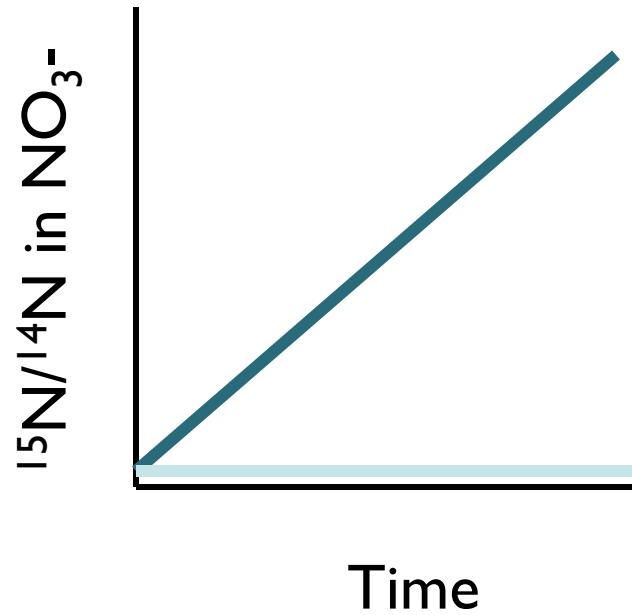
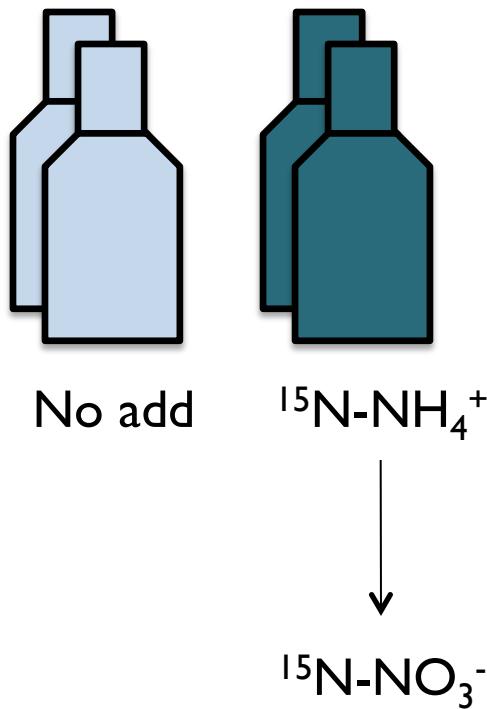
Another great thing about nitrogen: stable isotopes

Small differences in reaction rates leave an isotopic “signature” on compounds in the environment



Kinetic isotope effect:
 $\varepsilon = (^{14}k / ^{15}k - 1) \times 1000$

Stable isotopes can also be experimentally added to environmental systems as a tracer



See also Dugdale and Goering 1967; Lipschultz 2008

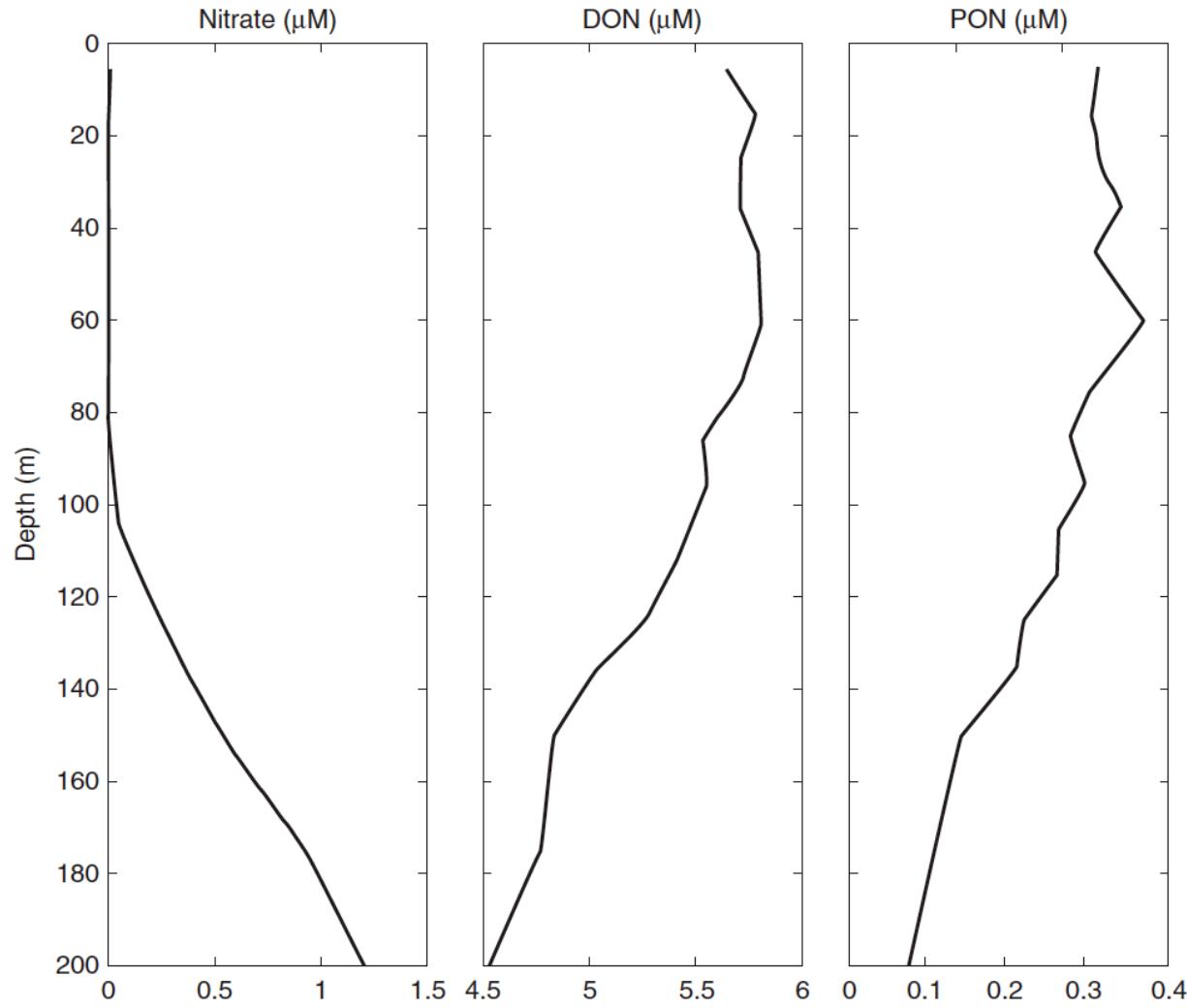
Vast majority of N in the ocean is N₂

Species	Mean conc. euphotic zone (mmol m ⁻³)	Mean conc. aphotic zone (mmol m ⁻³)	Oceanic inventory (Tg N)	Turnover rate ⁱ (Tg N yr ⁻¹)	Turnover time (years)
Nitrate, NO ₃ ^{-a}	7	31	5.8×10^5	1,570	370
Nitrite, NO ₂ ^{-b}	0.1	0.006	160		
Ammonium, NH ₄ ⁺ ^c	0.3	0.01	340	7,000	0.05
Dissolved Organic N, DON ^d	6	4	7.7×10^4	3,400	20
Particulate Organic N, PON ^e	0.4	0.01	400	8,580	0.05
Nitrous oxide, N ₂ O ^f	0.01	0.04	750	6	125
Fixed Nitrogen ^g			6.6×10^5	200	3,300
Nitrogen gas, N ₂ ^h	450	575	1×10^7	200	54,000

Gruber 2008

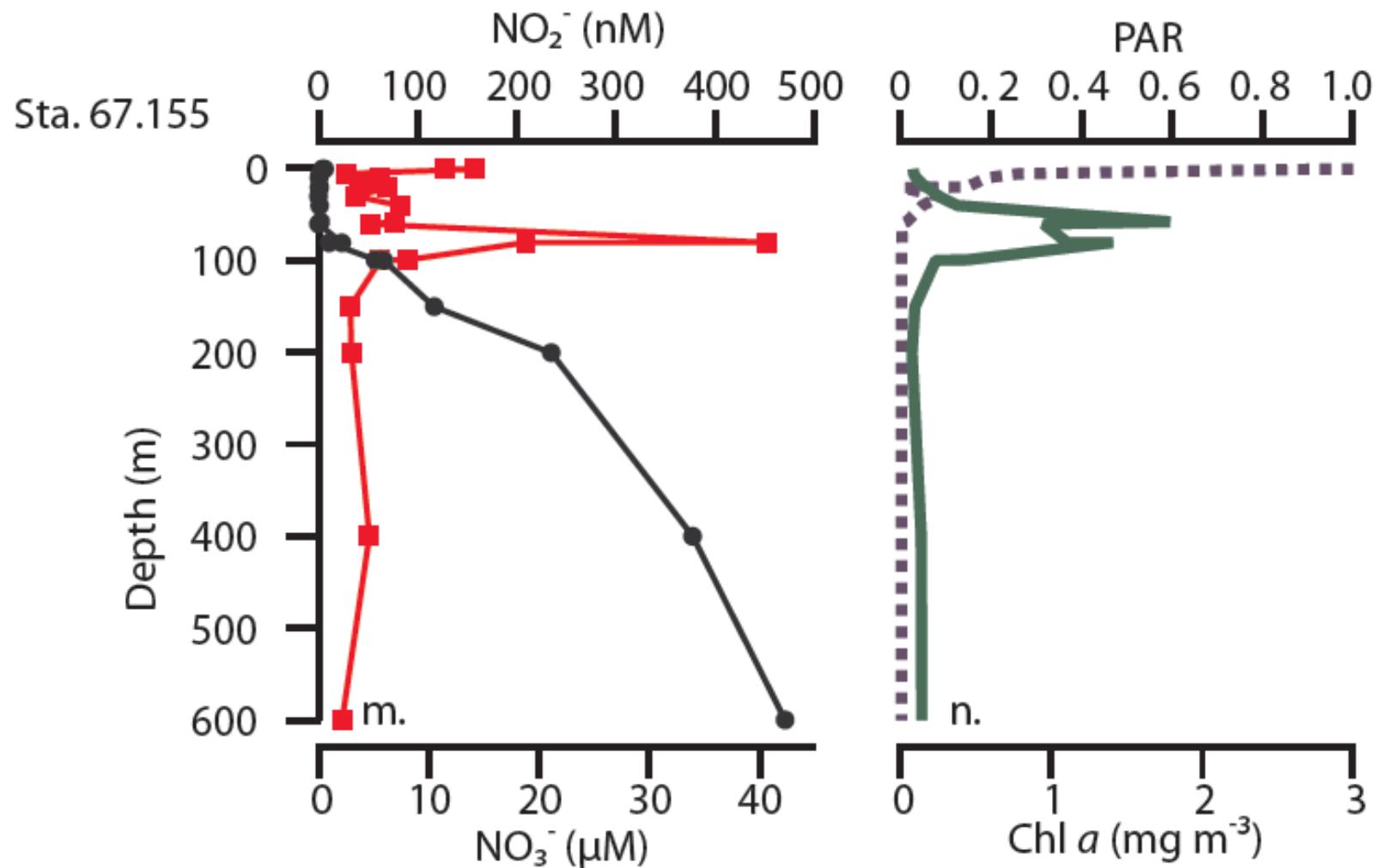
Nitrogen in the Marine Environment

DIN increases with depth, DON is highest at the surface

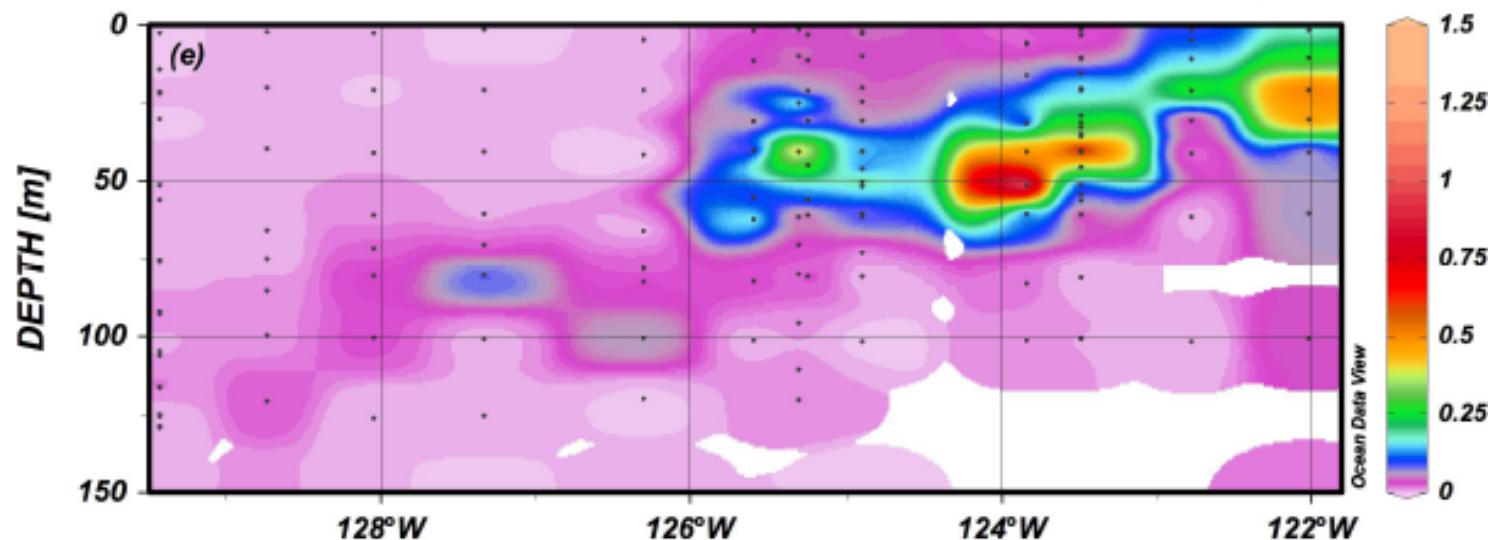


Karl et al. 2008

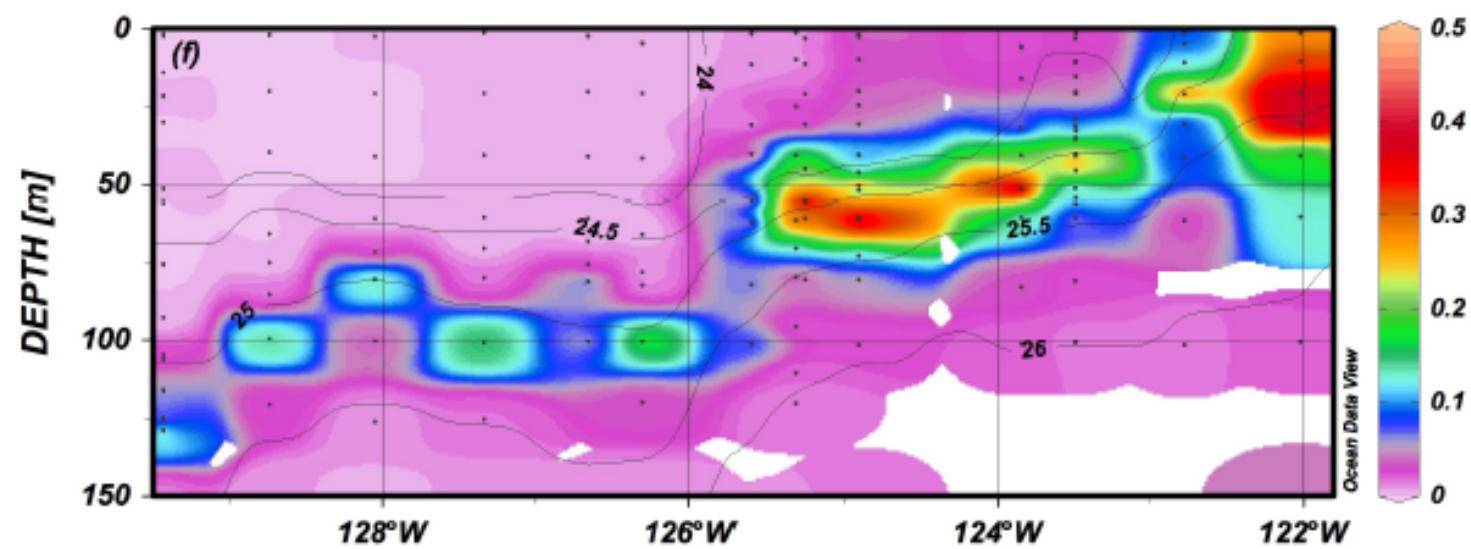
The primary nitrite maximum is a ubiquitous feature



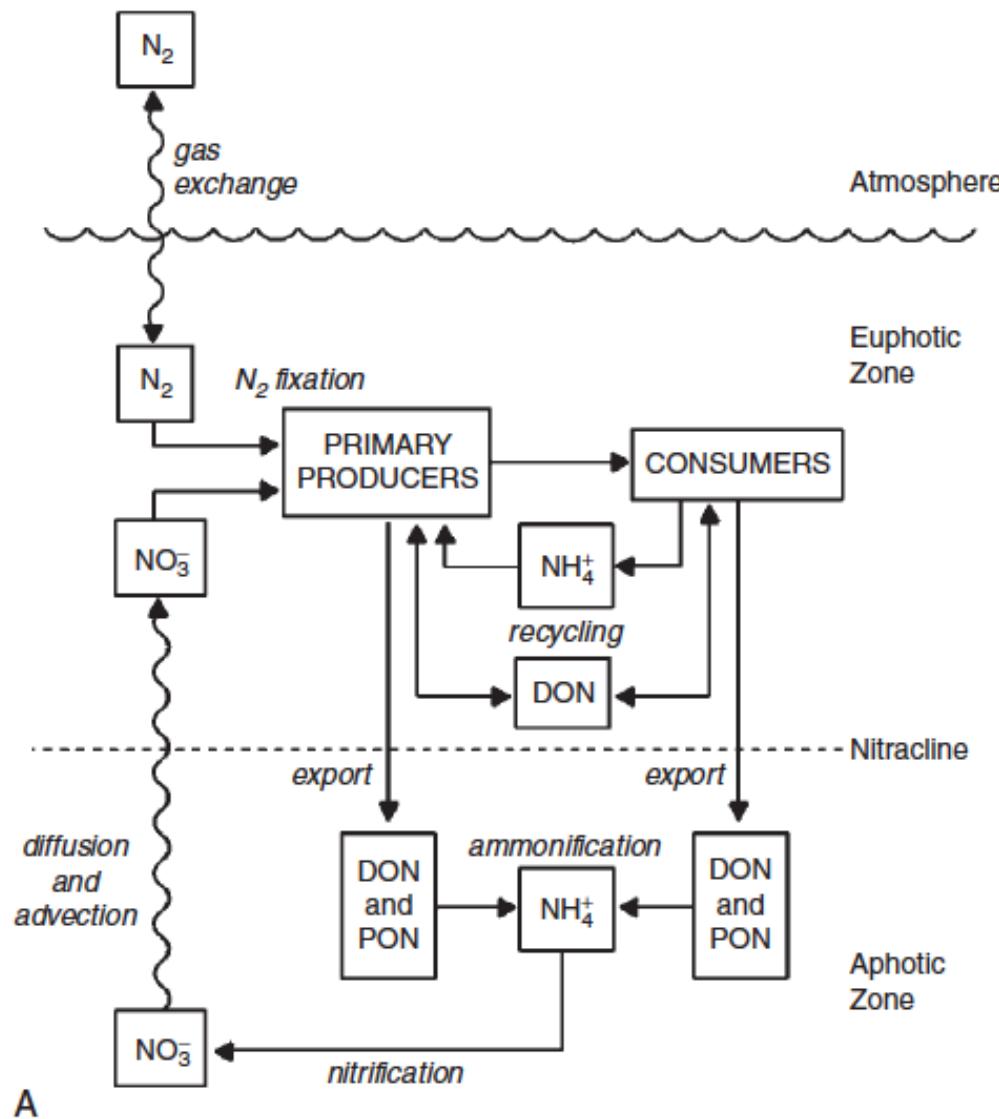
$\text{NH}_4^+ [\mu\text{M}]$



$\text{NO}_2^- [\mu\text{M}]$



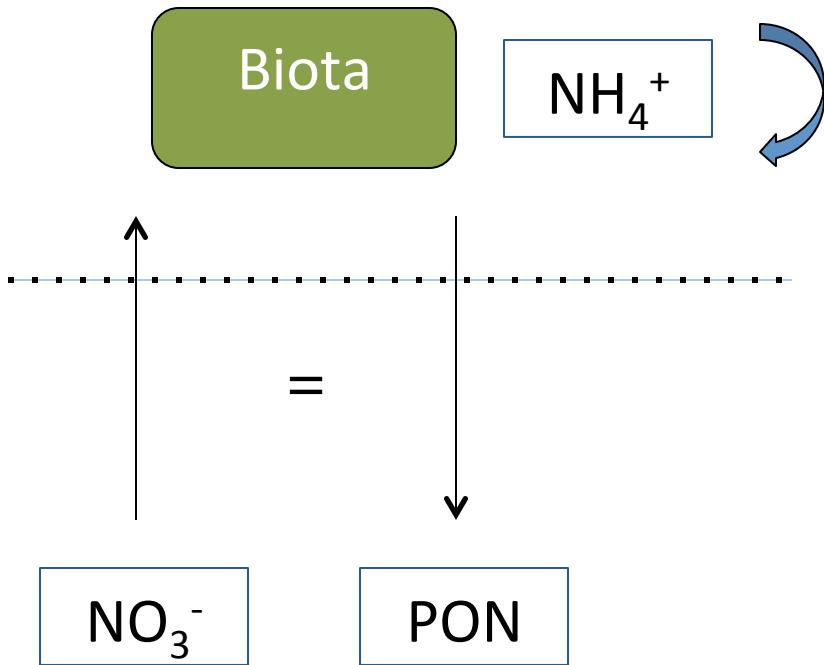
Spatial distribution of the nitrogen cycle in the marine water column



A

Karl 2008 NME

Estimating export from N uptake: The new production paradigm

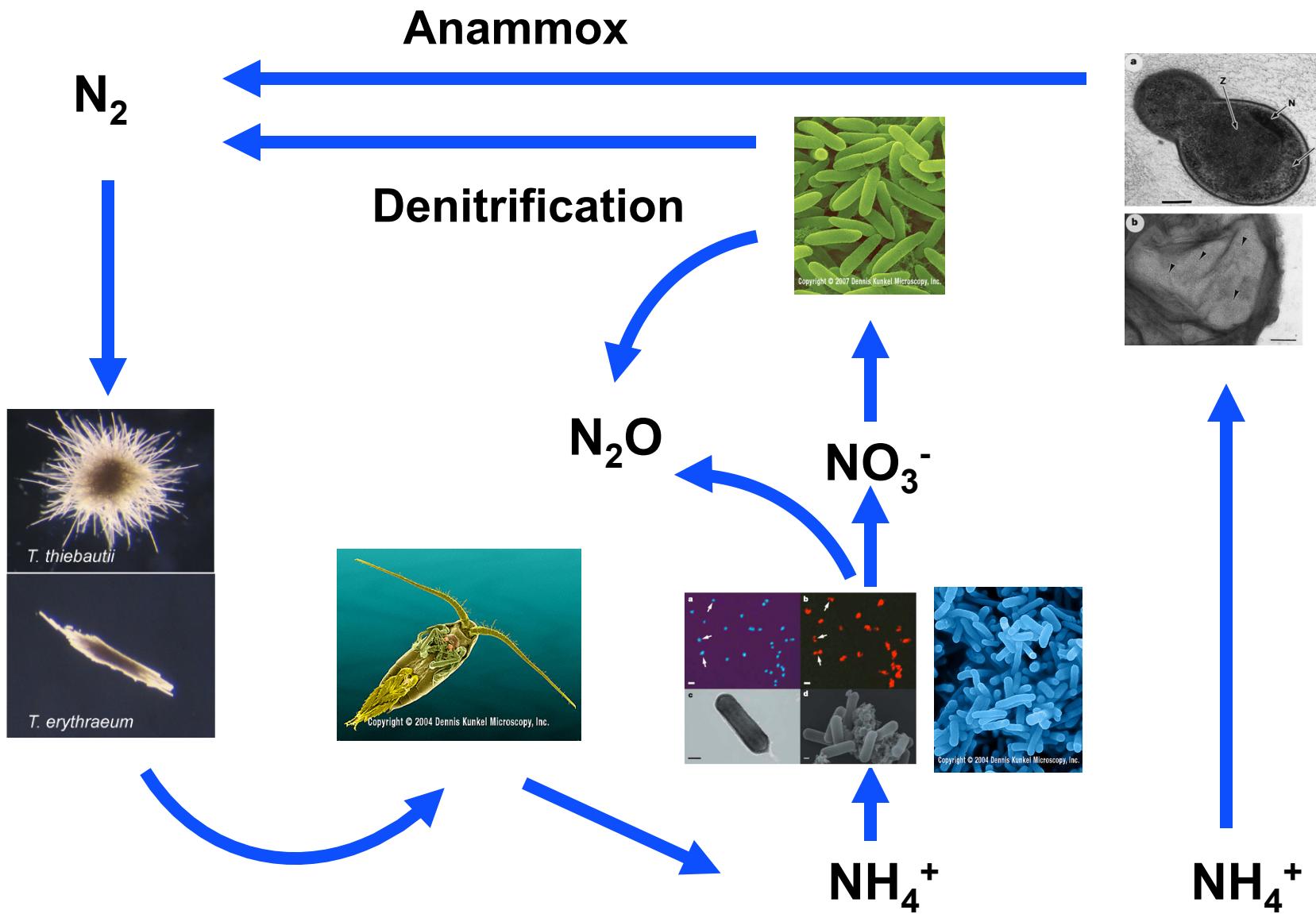


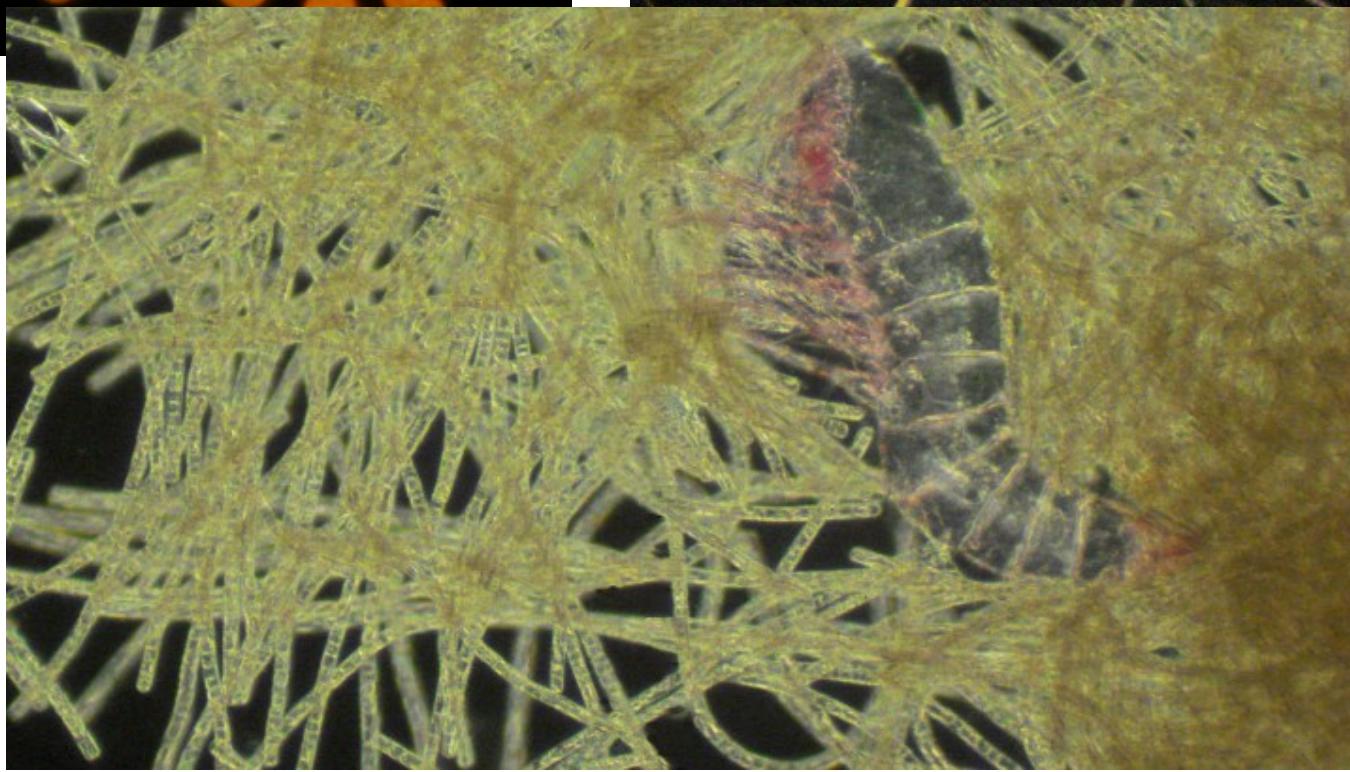
A net increase in phytoplankton biomass can only be supported by new NO_3^- entering the euphotic zone.

Complications:

- **N_2 fixation:** a source of non- NO_3^- 'new' nitrogen
- **Nitrification:** a non-new source of NO_3^-
- **DON:** release as DON is not true export

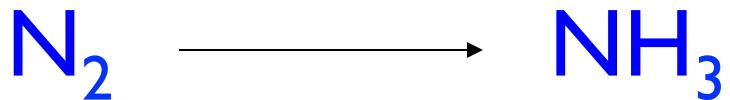
A condensed N cycle for today





Images by Angel White, downloaded from C-MORE website.

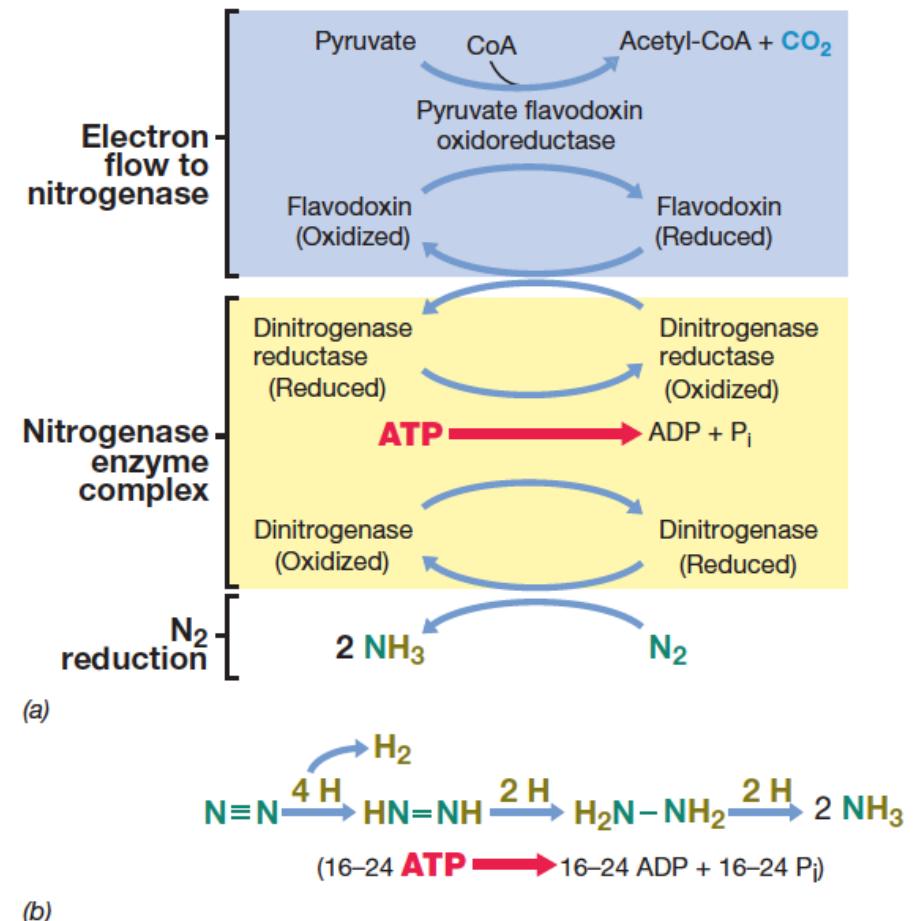
Nitrogen fixation: Acquiring N in an N-limited ocean



- Though thermodynamically favorable, not coupled to energy generation, and very energy intensive (16-29 ATP per N_2)
- Widespread among prokaryotes, and occurs in oxic and anoxic environments
- Diazotroph = “two nitrogen eater” = nitrogen-fixing organism

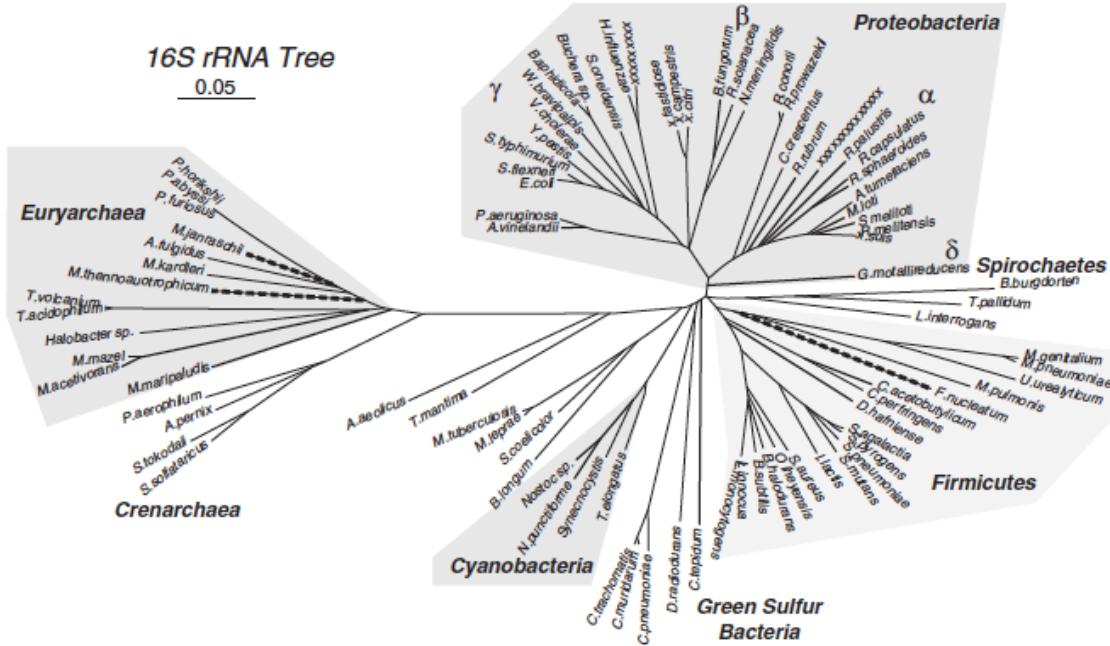
Nitrogenase: Breaking the N-N triple bond

- Nitrogenase enzyme complex = an Fe-enzyme (dinitrogenase reductase) and Fe-Mo-enzyme (dinitrogenase). Alternative forms contain Vanadium.

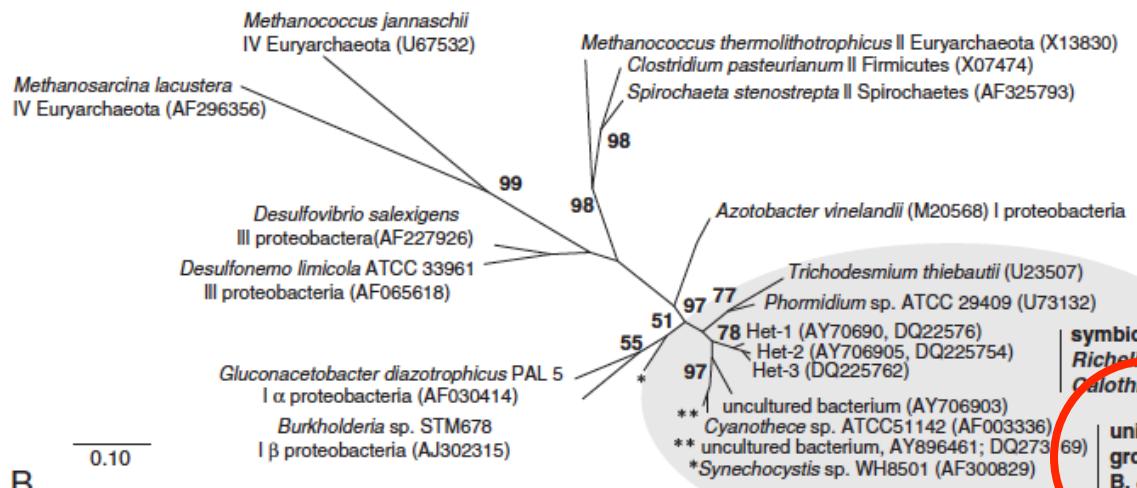


nifH and rRNA phylogenies are congruent.

Suggests that *nif* genes are ancient, inherited vertically over evolutionary time, and lost in some organisms



A



B

Great story! Zehr, Tripp, Thompson and coworkers

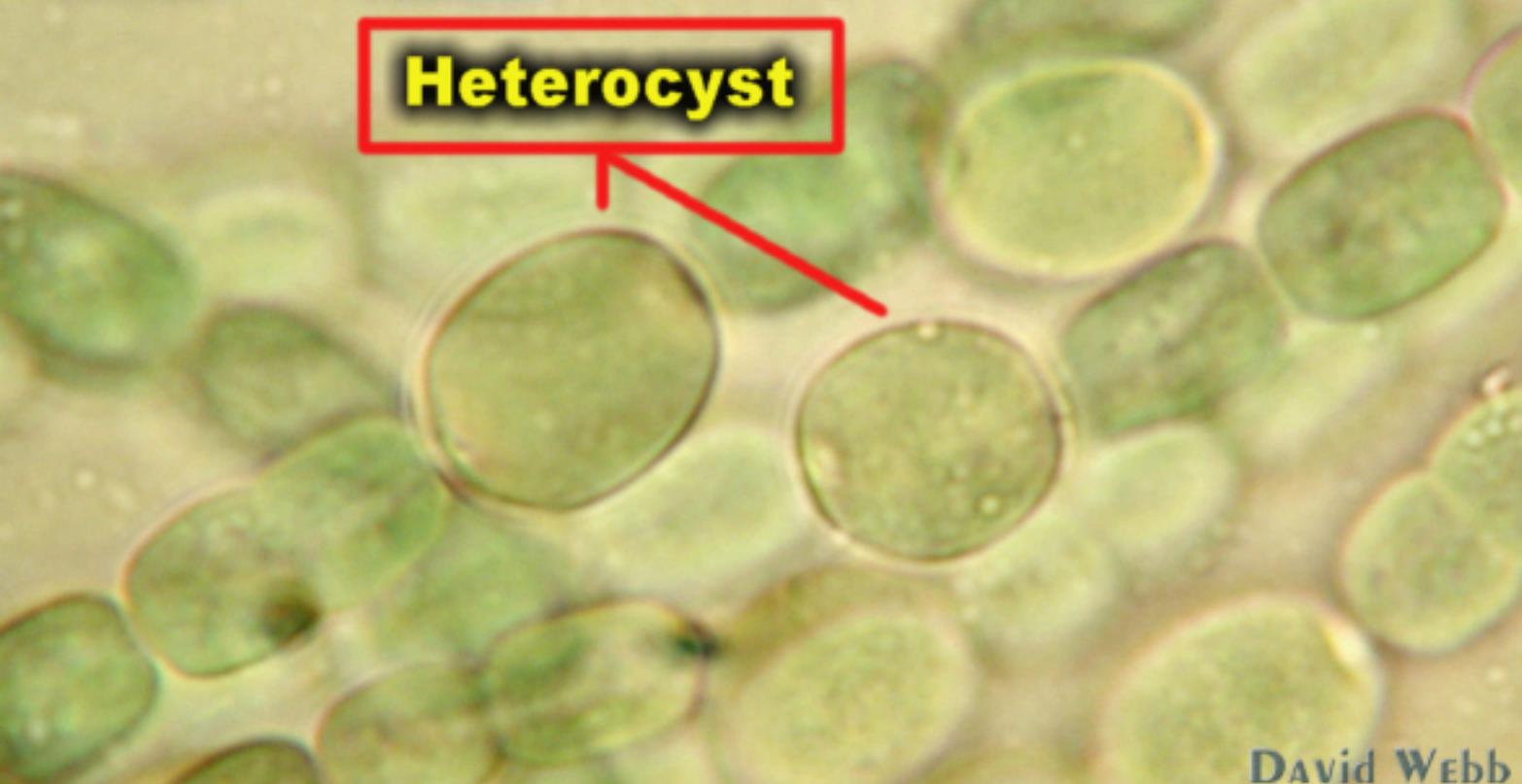
Oxygen is **bad** for N₂ fixation

O₂ is released by photosynthesis, but O₂ irreversibly inhibits nitrogenase. How can organisms that do both cope?

Dealing with O₂: Spatial segregation

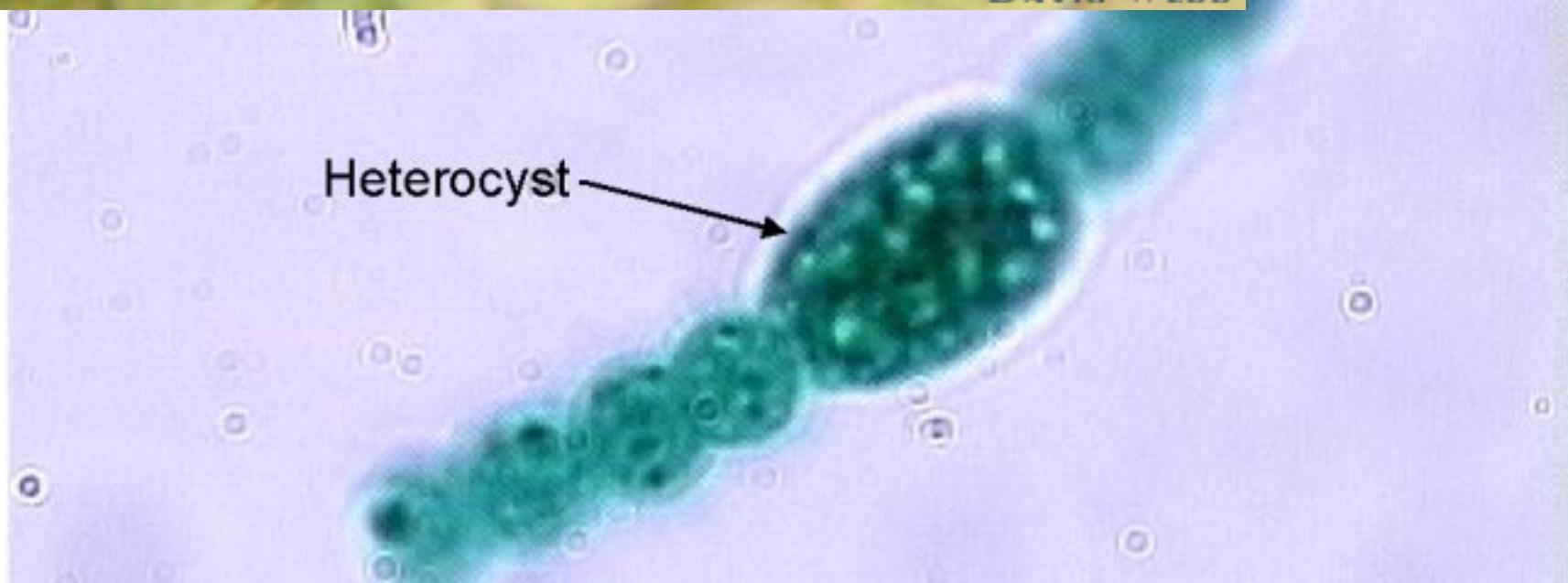
- Heterocysts: specialized cells, lack PSII. Common in freshwater cyanobacteria (*Anabena*) but relatively uncommon in marine waters, except in symbiosis with diatoms : *Richelia intracellularis*.
- Intracellular segregation: *Trichodesmium* (Finzi-Hart *et al.*, 2009); Polysaccharide capsule in aerobic heterotrophs that maintain a low internal oxygen concentration.

Heterocyst

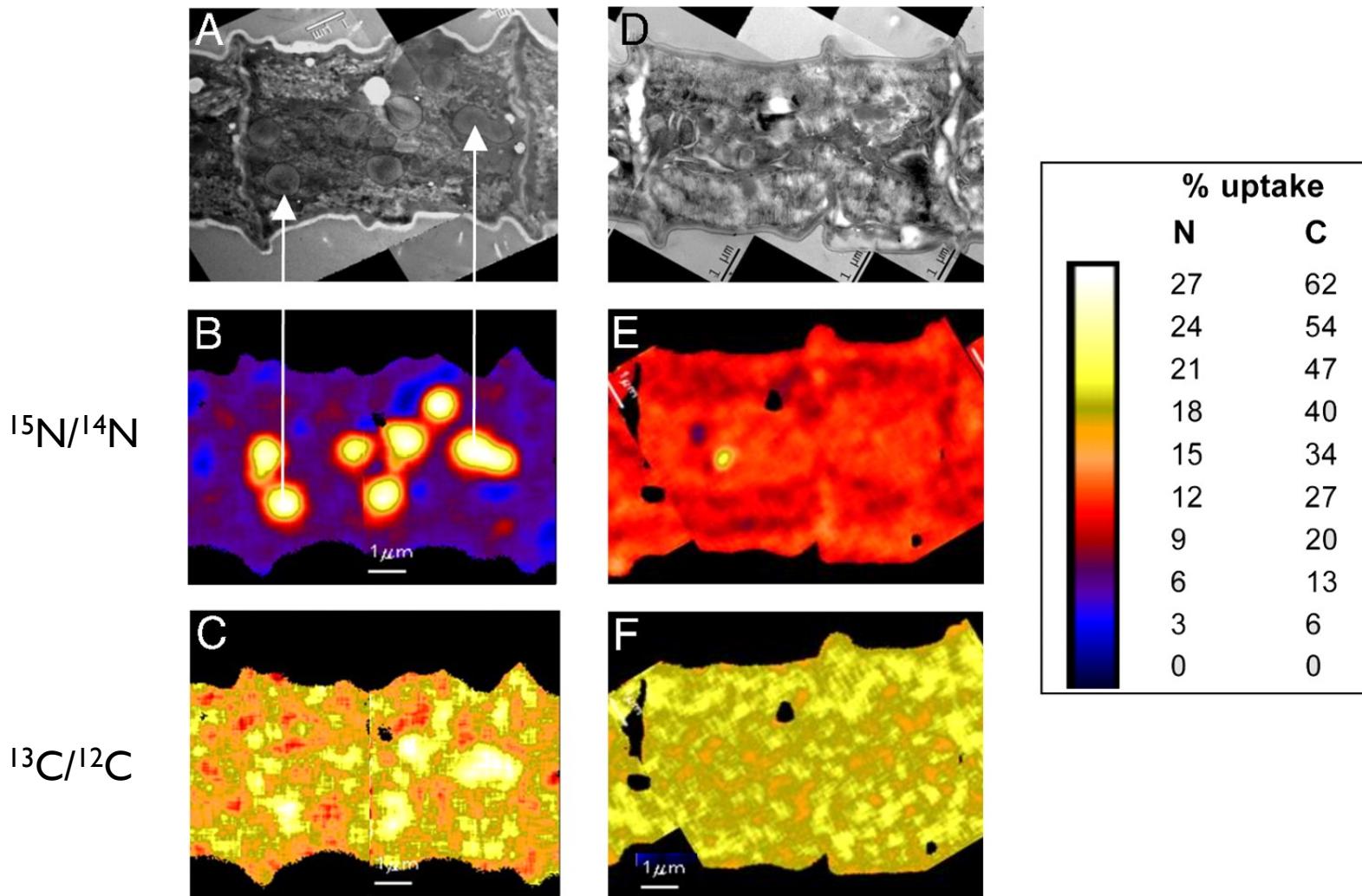


David Webb

Heterocyst



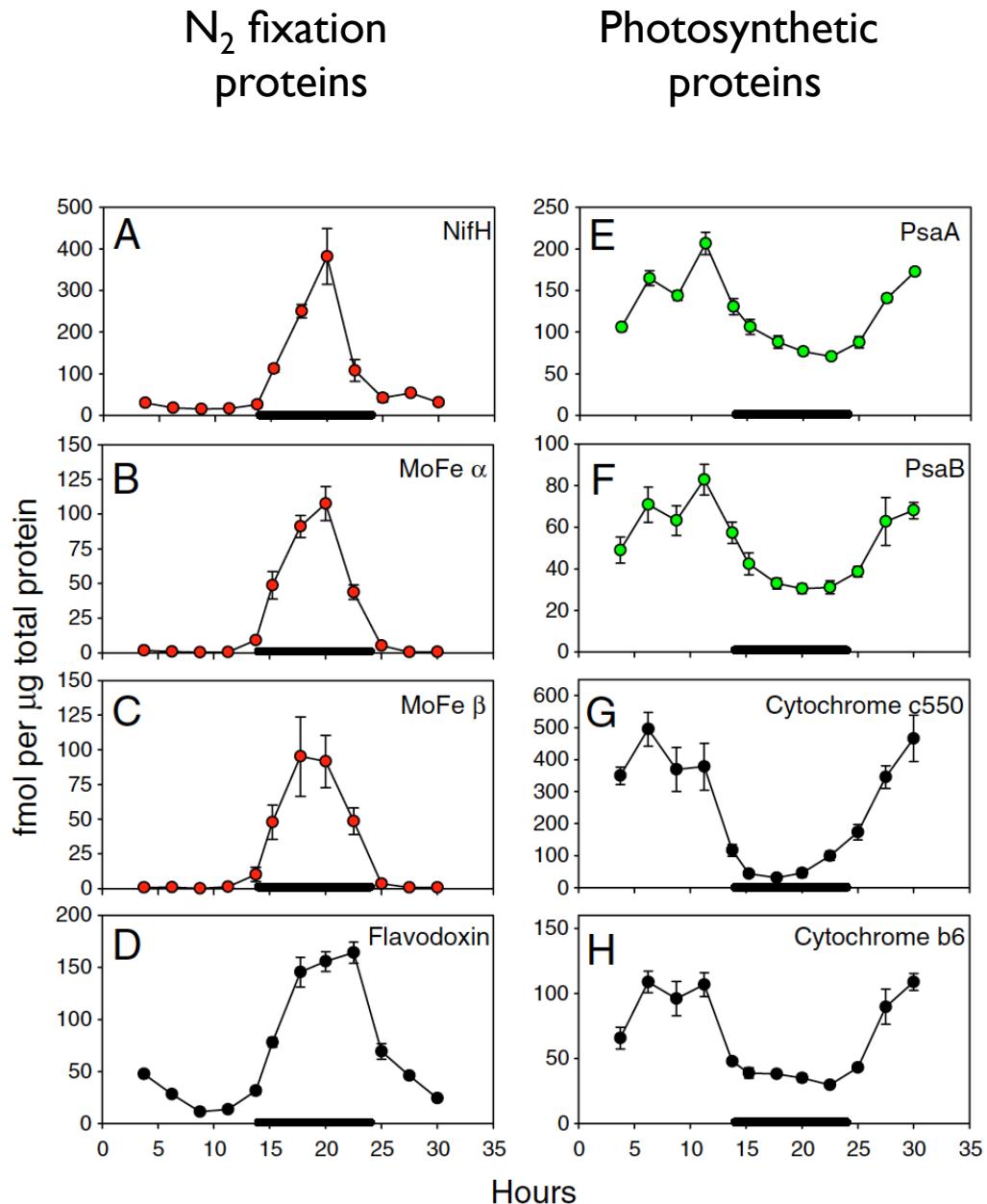
NanoSIMS used to track the site of N₂ fixation



Dealing with O₂: Temporal segregation

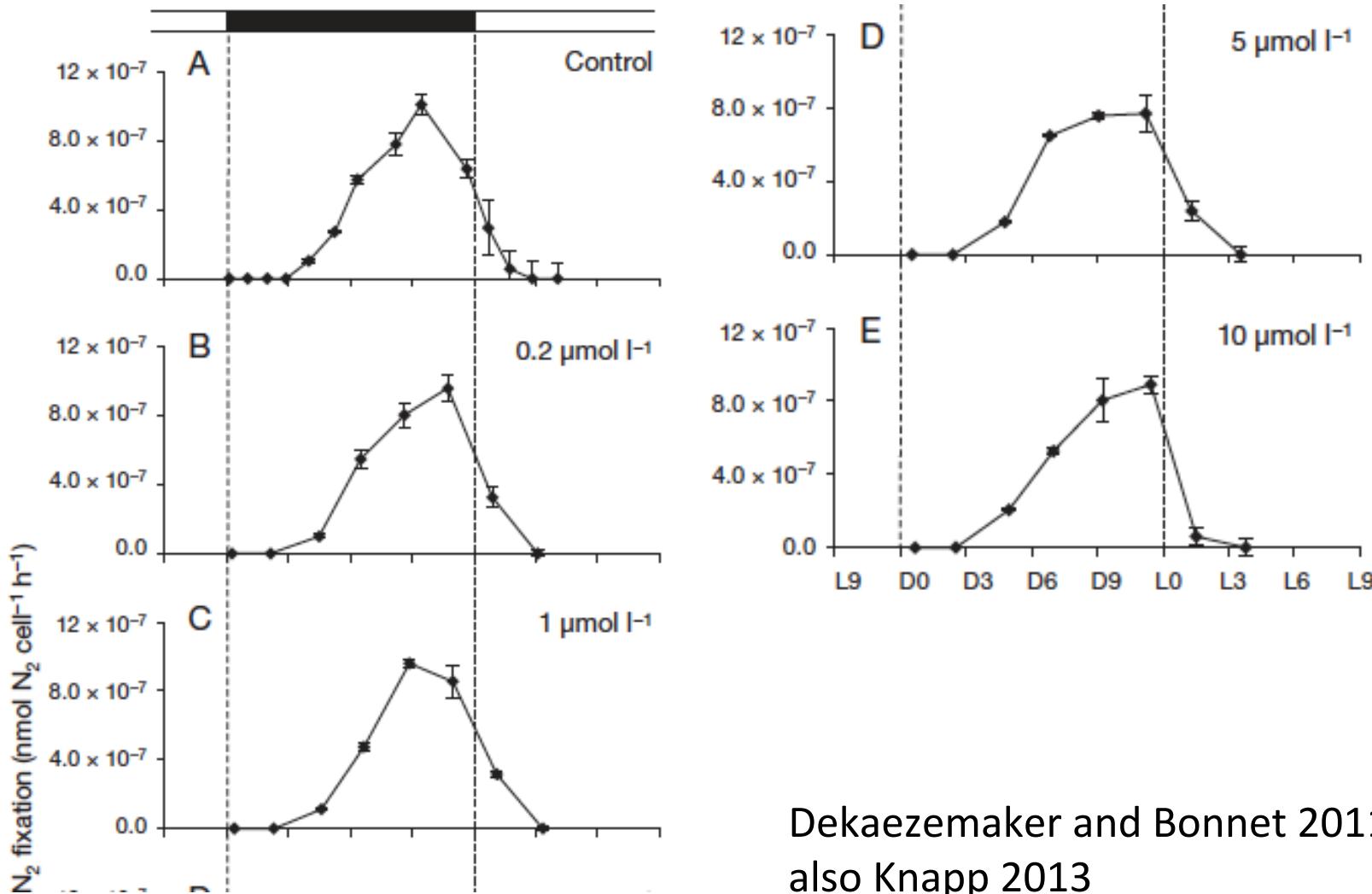
Unicellular Cyanobacteria fix nitrogen at night when there is no photosynthesis and cellular oxygen is low.

Crocospaera watsonii makes N₂ fixation proteins at night and photosynthetic proteins during the day to save Fe.



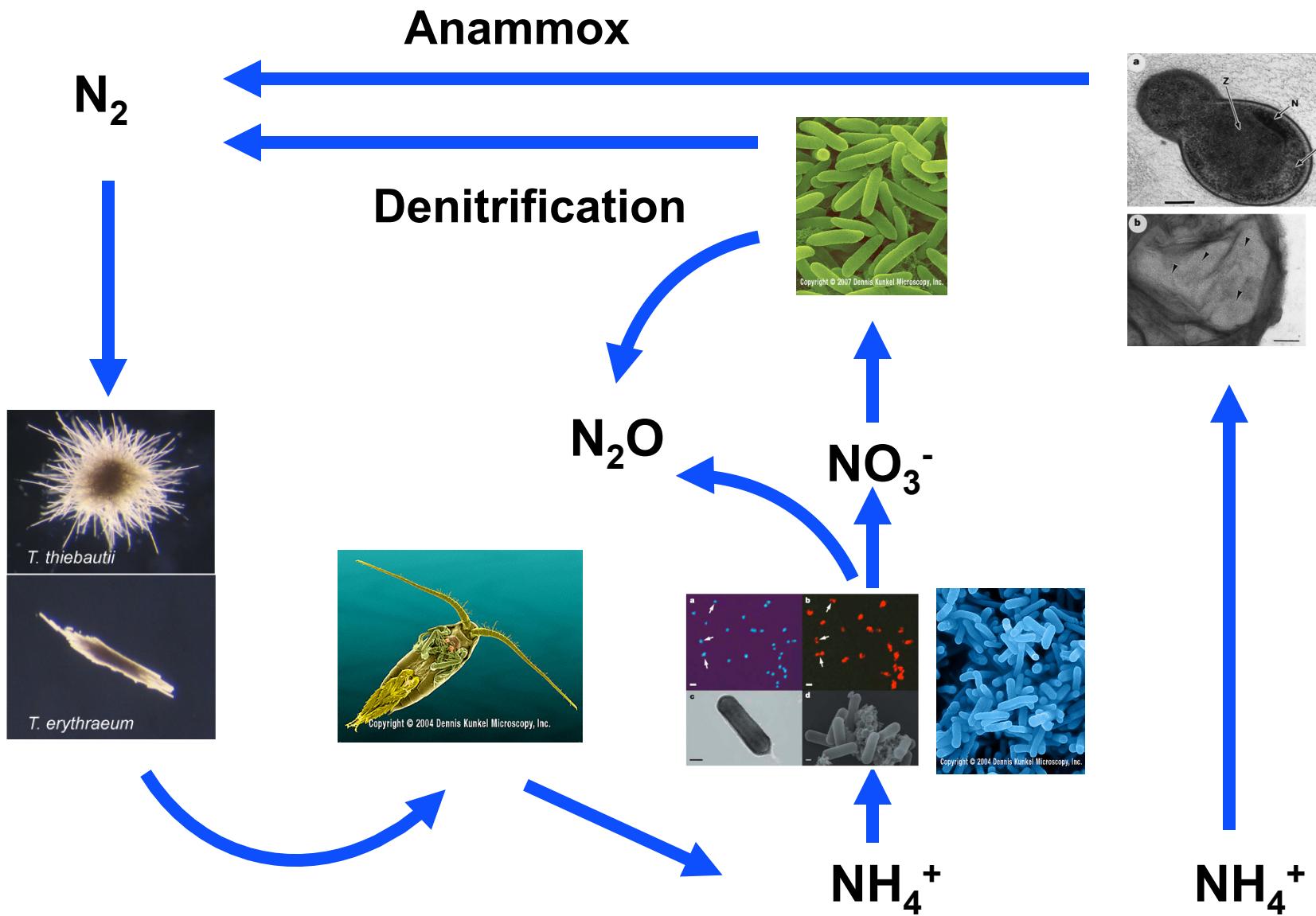
Saito et al. 2011

N_2 fixation can occur in the presence of DIN

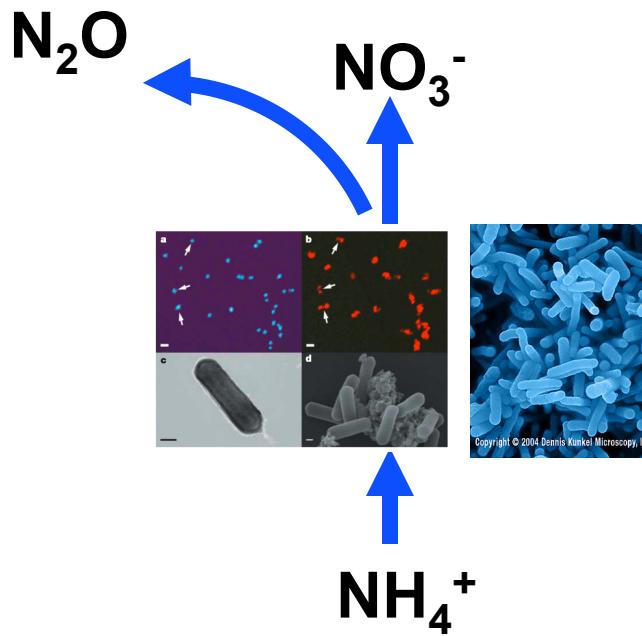


Dekaezemaker and Bonnet 2011; see
also Knapp 2013

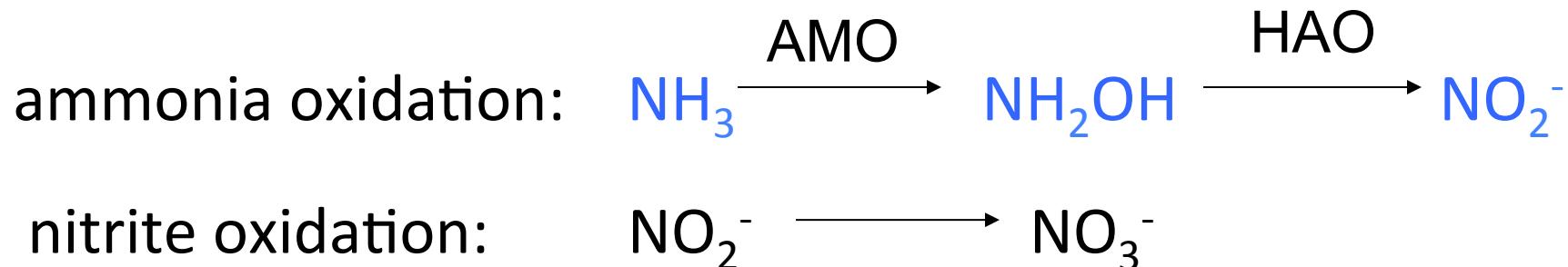
A condensed N cycle for today



A condensed N cycle for today



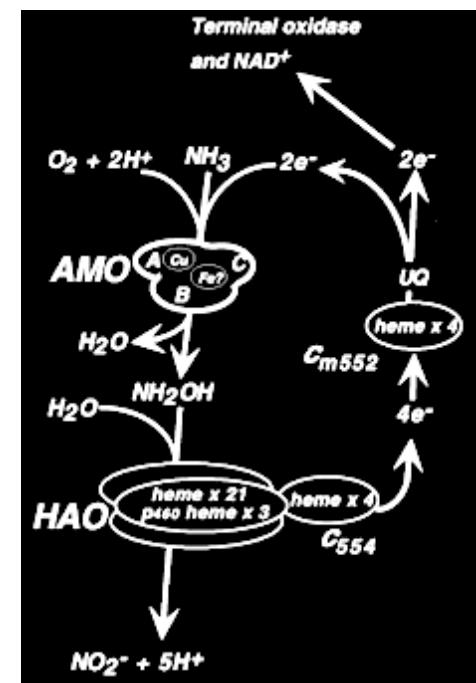
Nitrification is a two step process

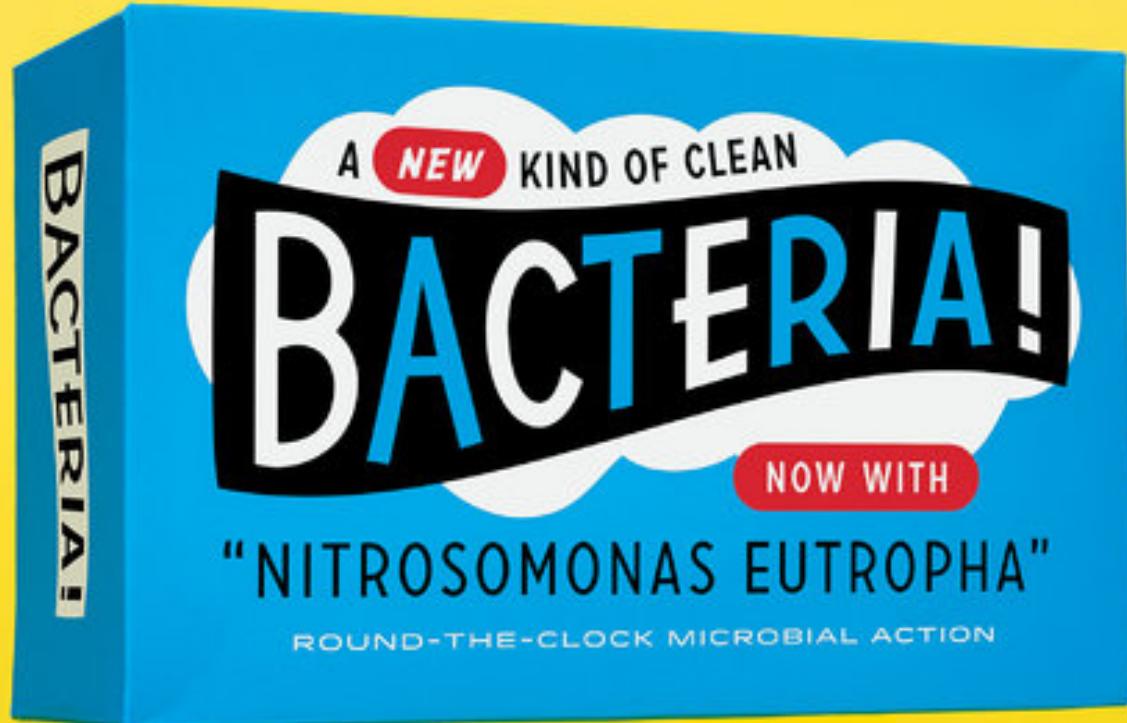


No organisms (yet?) known that can catalyze the complete oxidation of NH_3 to NO_3^- .

Some contention about whether NH_3 or NH_4^+ is transported into the cell and is the actual enzymatic substrate.

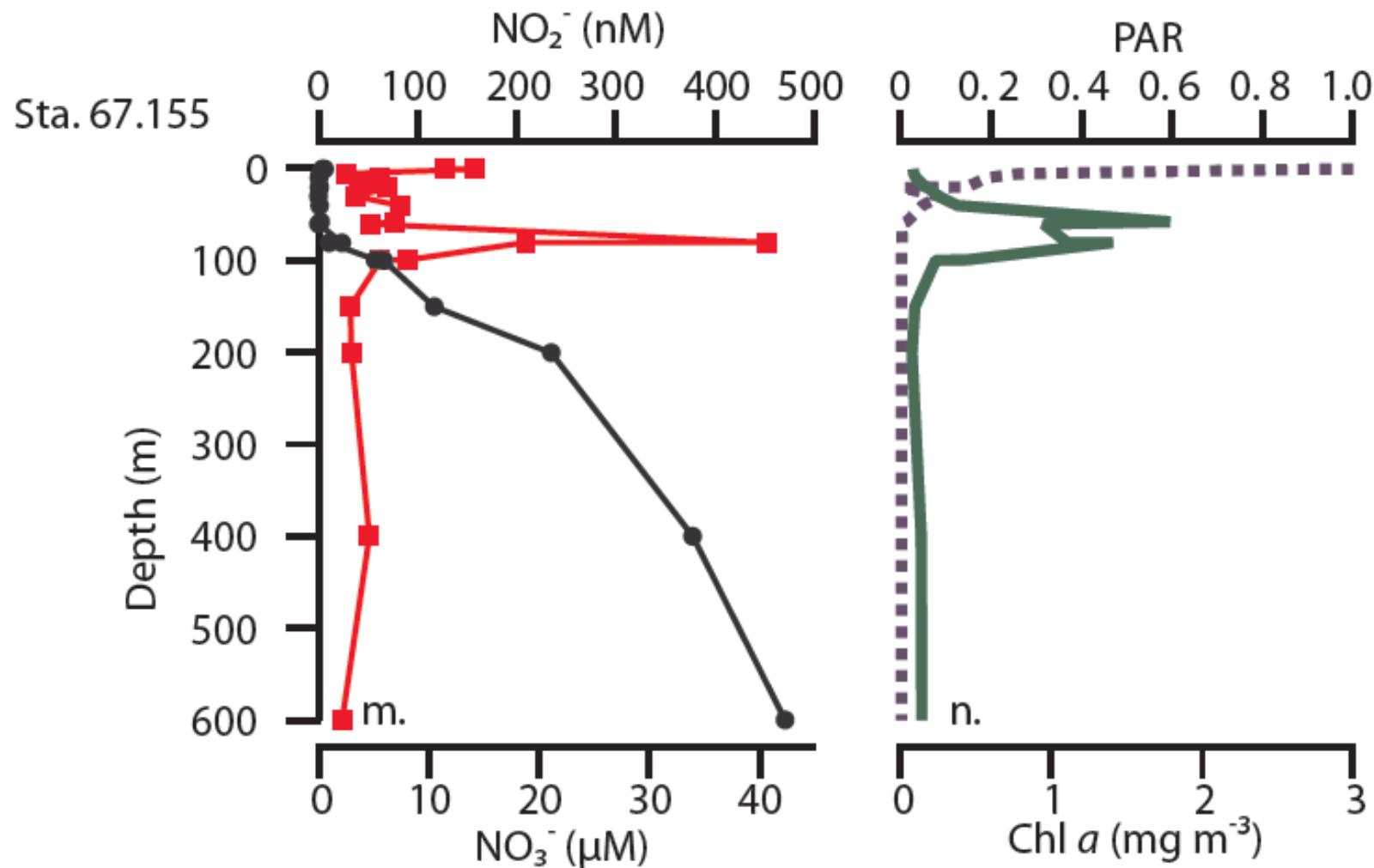
Model organisms are all autotrophic . . . fussy, slow growing, bastards.





NYT, 5/25/14

The primary nitrite maximum is a ubiquitous feature



Nitrification = nitrogen remineralization

THE EXPERIMENTAL DECOMPOSITION AND
REGENERATION OF NITROGENOUS
ORGANIC MATTER IN SEA
WATER¹

THEODOR VON BRAND, NORRIS W. RAKESTRAW AND
CHARLES E. RENN

3. The main stages in the decomposition are: dead body—ammonia—nitrite—nitrate.

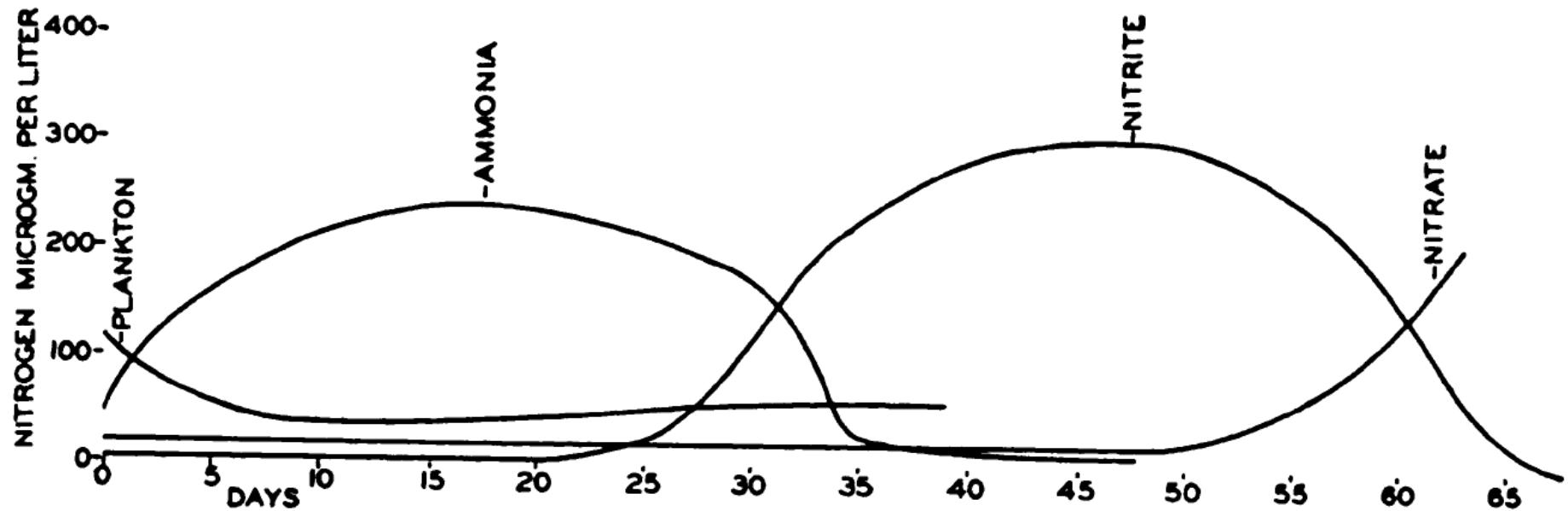
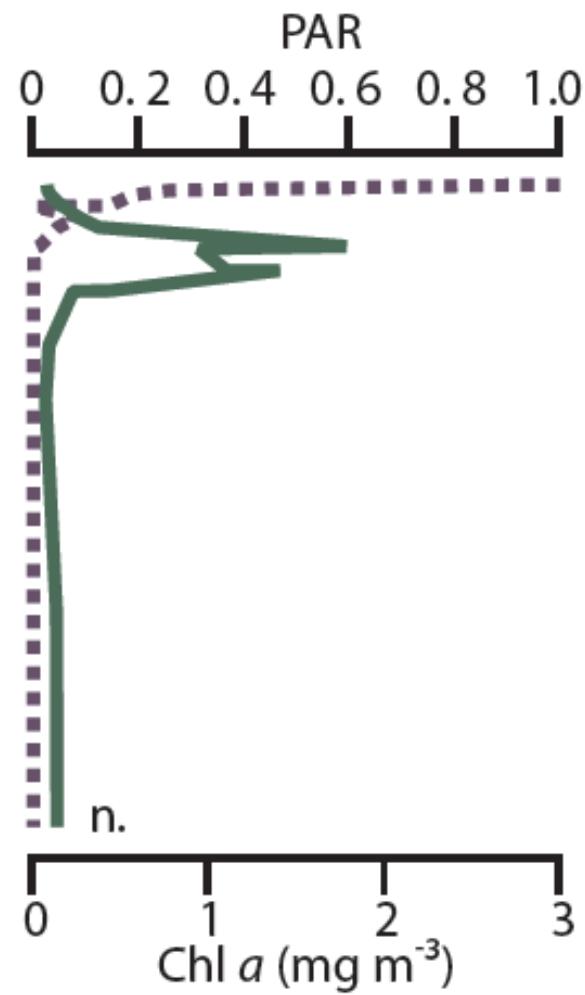
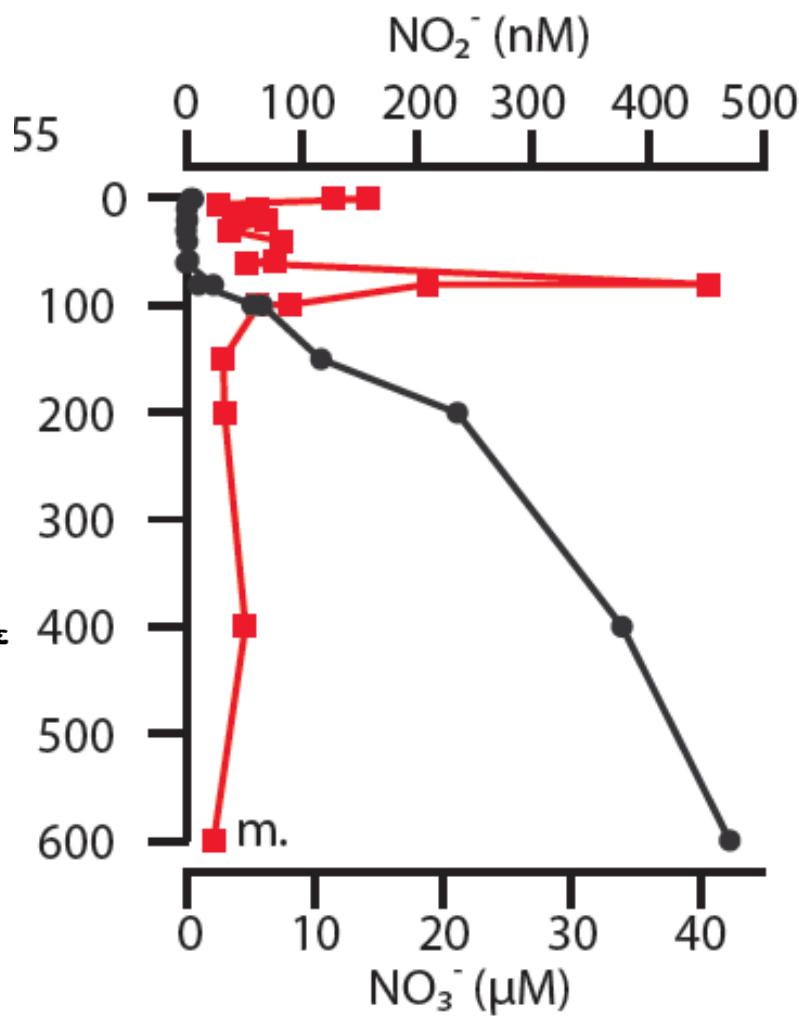
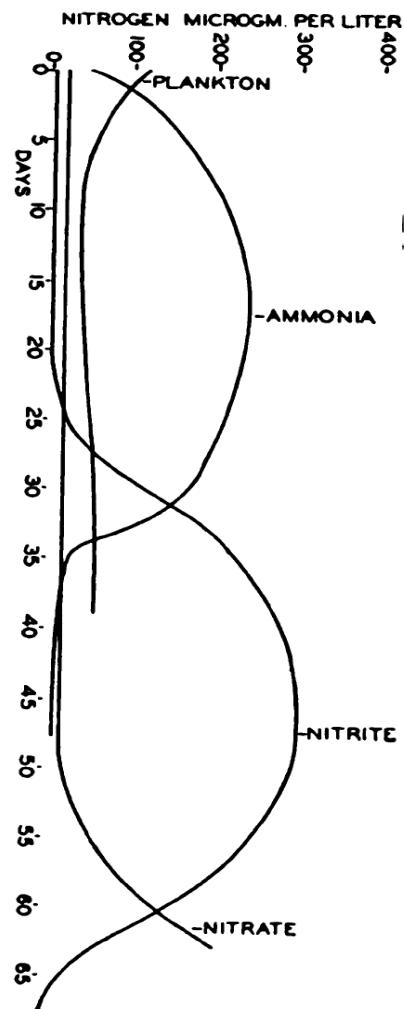


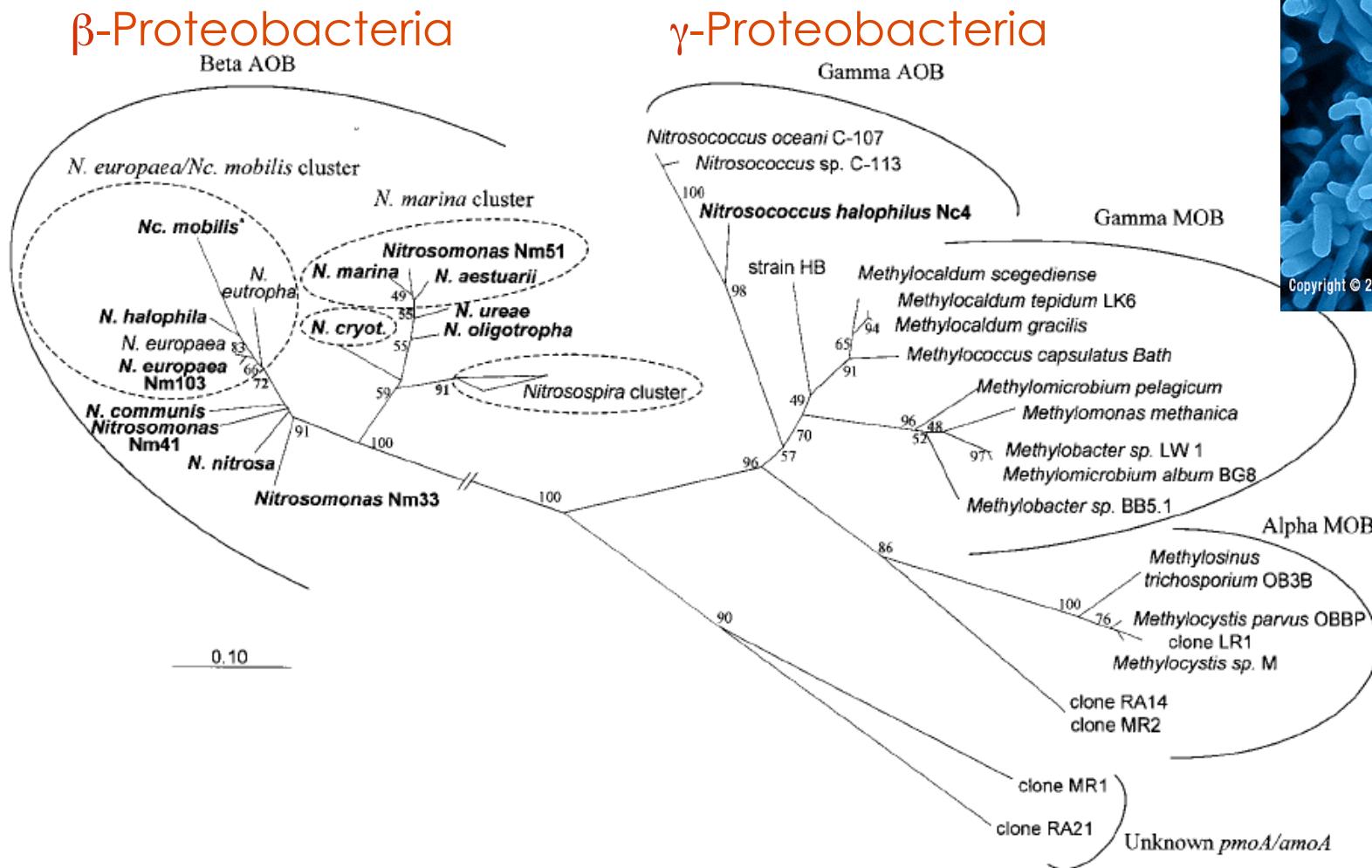
FIG. 2. Series IV. The decomposition of nitrogenous organic matter in mixed plankton, showing the appearance of soluble nitrogen compounds in the water in which it is suspended. Plankton previously filtered through No. 8 bolting silk.

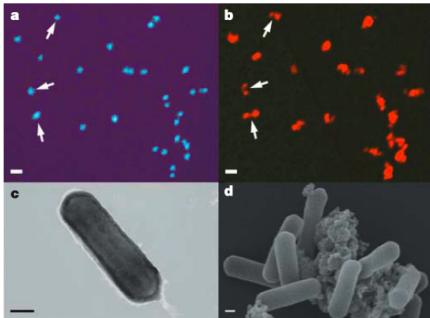


Nitrifiers perform a very specific task

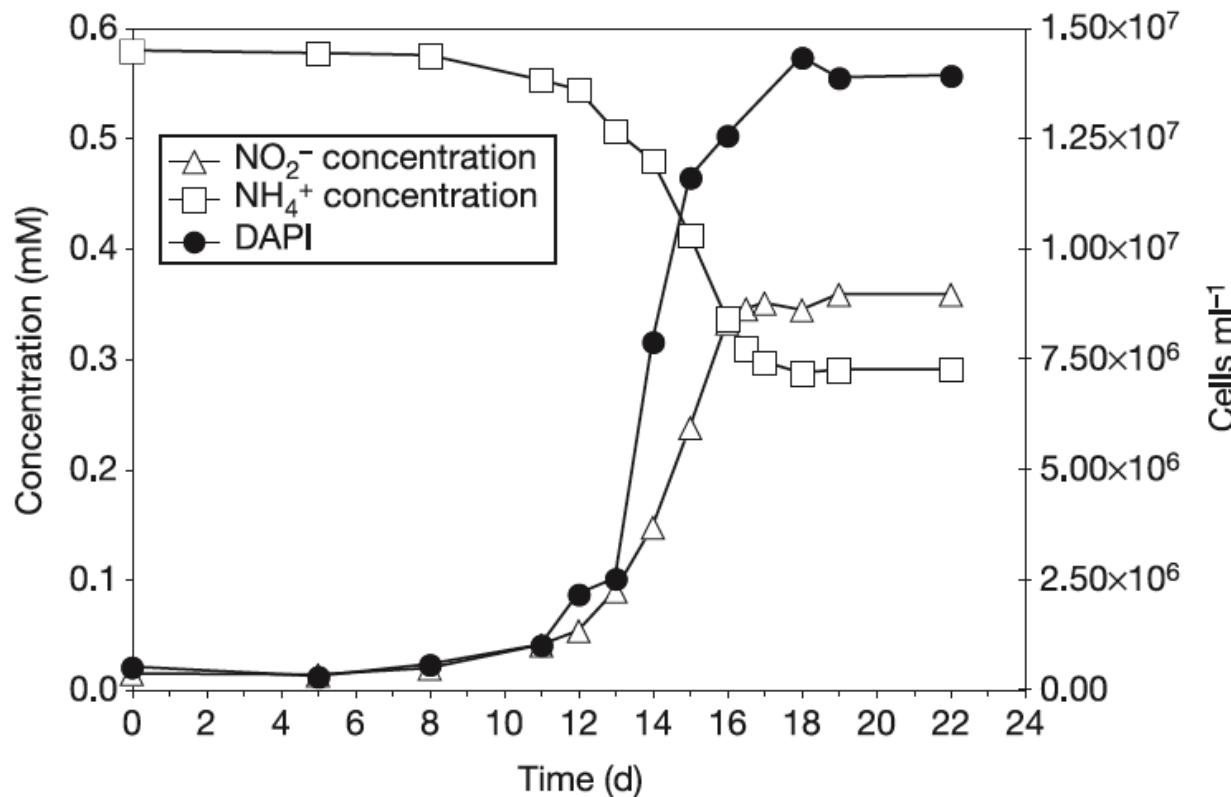
- Fix CO_2 or bicarbonate for anabolic reactions, oxidize NH_3 for energy
- Slow growth and somewhat inflexible nutritional requirements (more on this in next lecture)
- Historic view of nitrifiers in the ocean as being inhibited by light due to photobleaching of cytochrome c

amoA phylogeny of AOB is highly congruent with 16S tree





The first culture of a mesophilic marine crenarchaeon, isolated from gravel at the Shedd Aquarium.



Archaea can
oxidize NH_3

Ecology of ammonia oxidizers

- *amoA* phylogeny is congruent with 16S rRNA phylogeny. Not much lateral gene transfer.
- At low O₂ tension ammonia oxidizers produce N₂O. Some ammonia oxidizers contain *nirK* genes (dissimilatory nitrite reductase)
- Potentially inhibited by light (photobleaching of cytochrome C). Hypothesis that NOB are more inhibited than AOB.

AOA and AOB have different life history strategies

AOB

Larger cells

Easier to culture

High K_m for NH_3

Faster growth

Big genomes

Fe-based electron transport

Very light sensitive

AOA

Small cells

Difficult to culture

Low K_m for NH_3

Slow growth

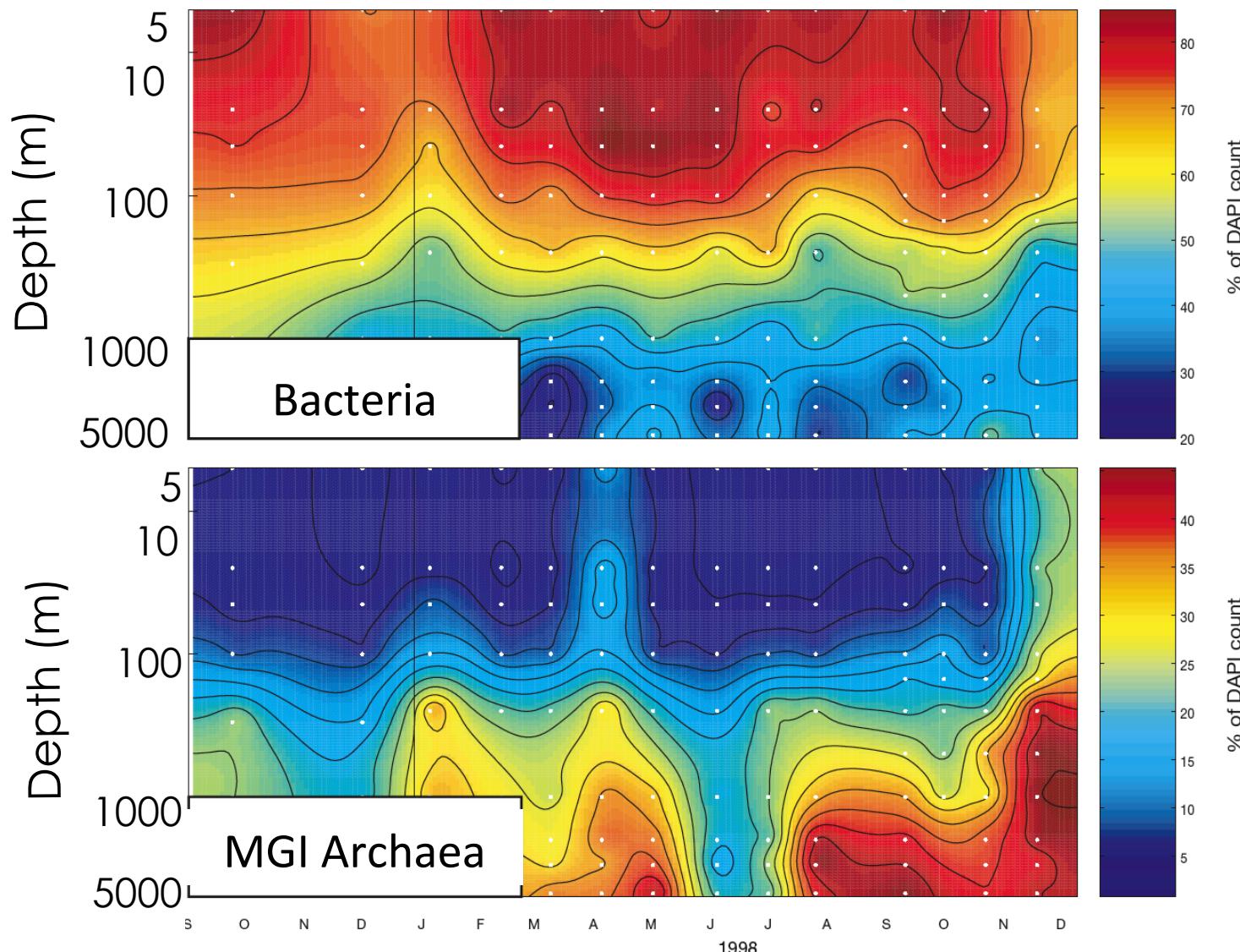
Reduced genomes

More tolerant of low O_2

Cu-based electron transport

Highly efficient C fixation pathway

Putative ammonia oxidizers are among the most abundant cells in the deep ocean.

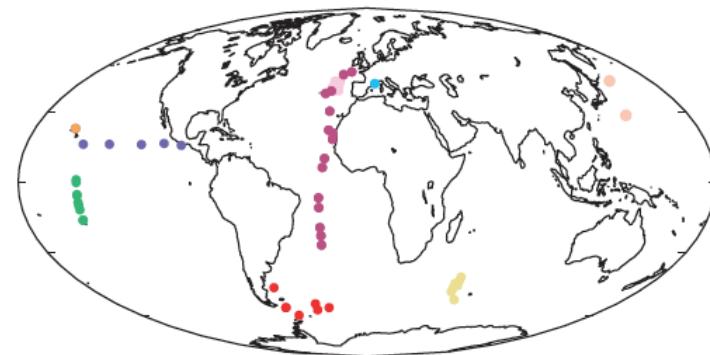


modified from Karner et al., Nature, 2001

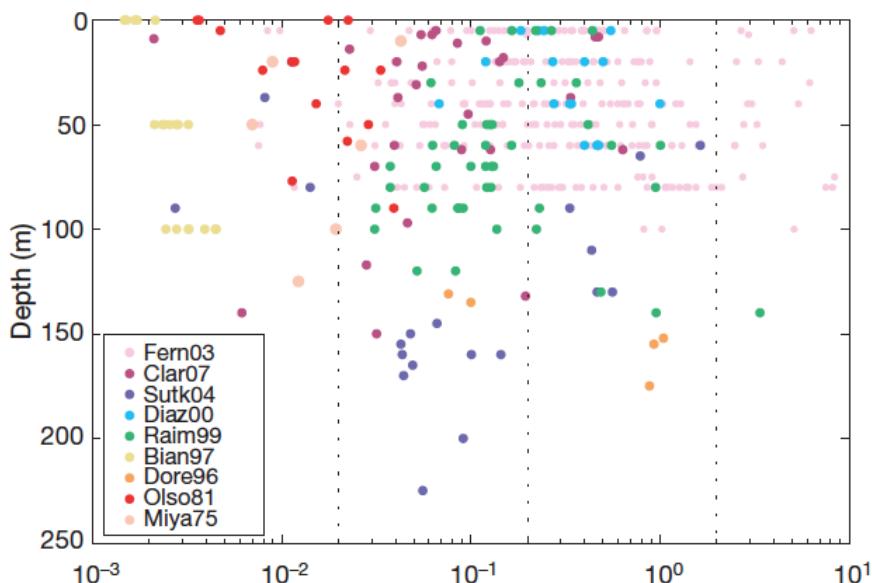
The significance of nitrification for oceanic new production

Andrew Yool¹, Adrian P. Martin¹, Camila Fernández^{2,3} & Darren R. Clark⁴

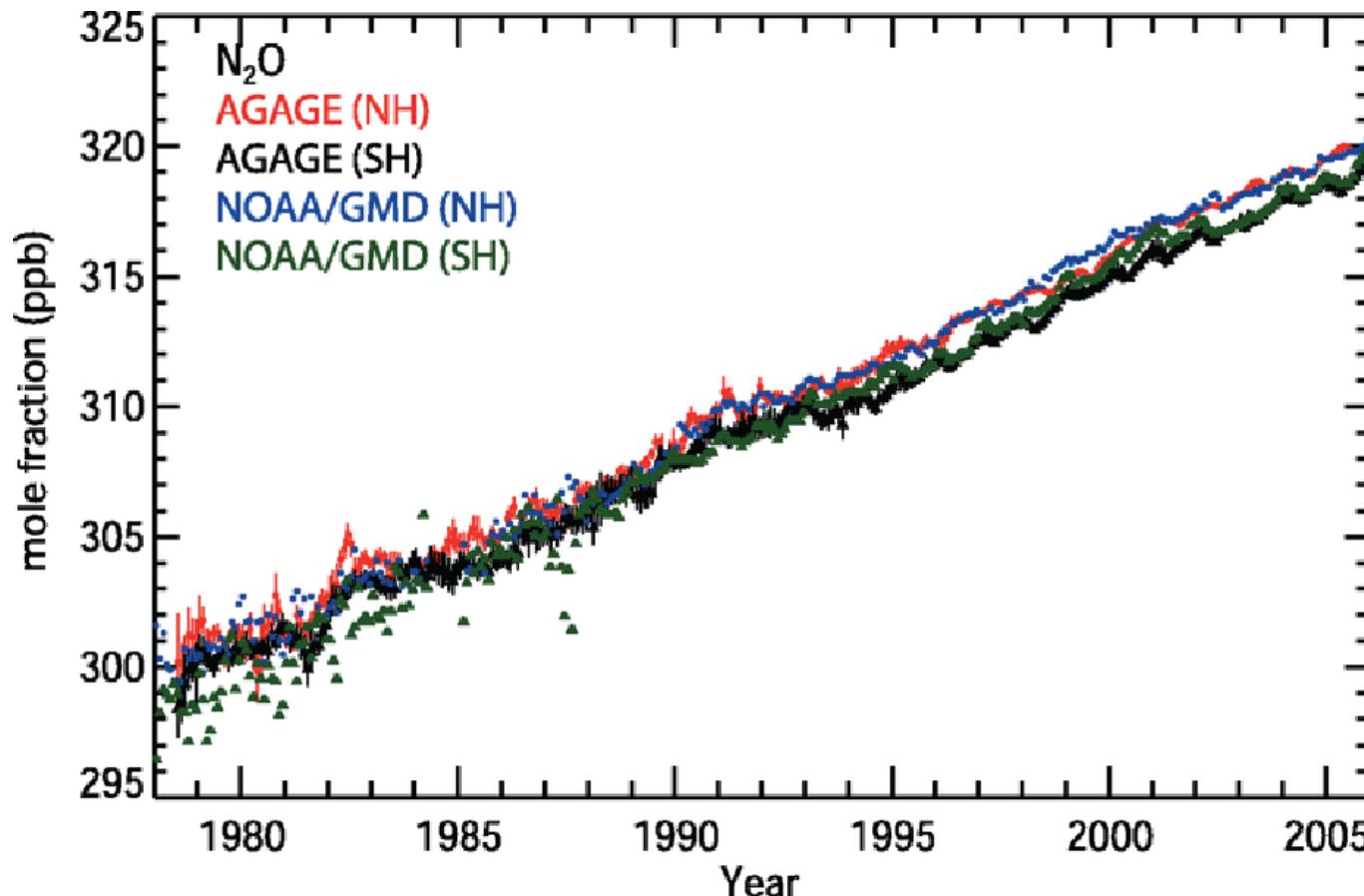
a



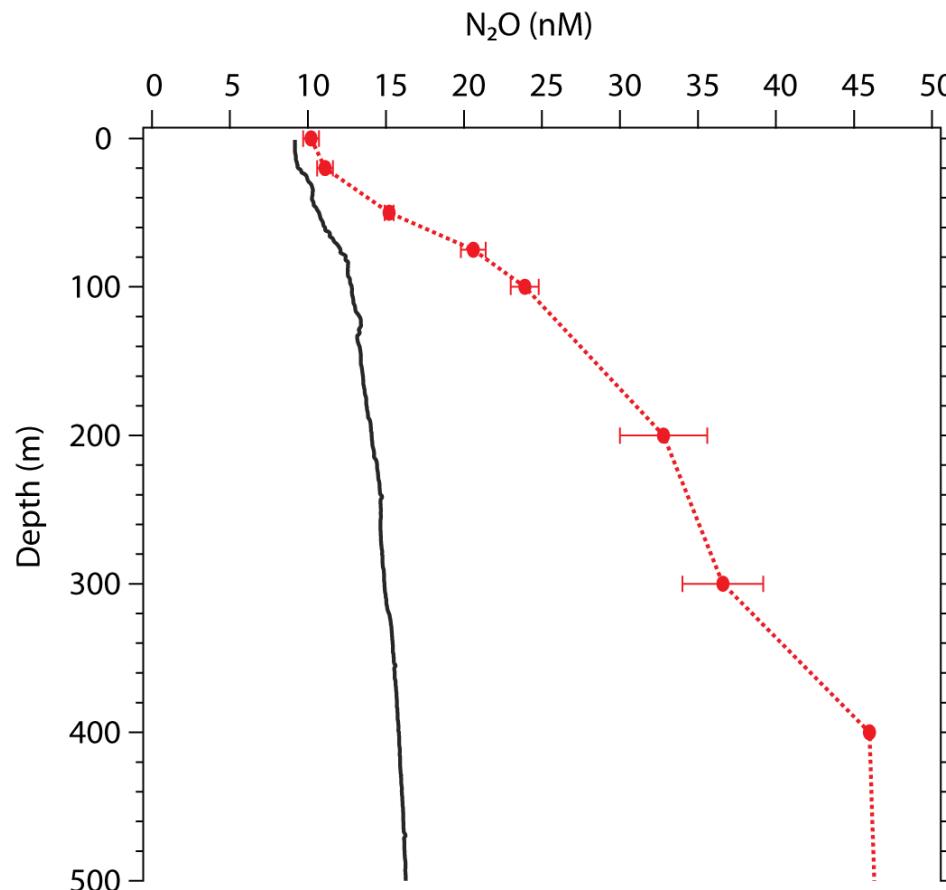
b



N_2O is a greenhouse gas whose concentration in the atmosphere is increasing

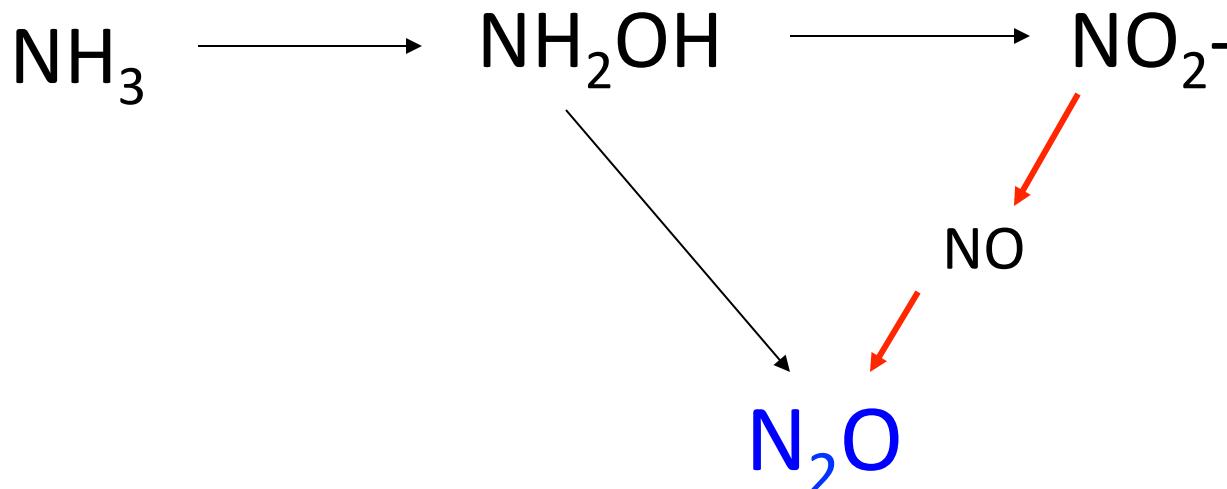


N_2O is supersaturated in most of the ocean

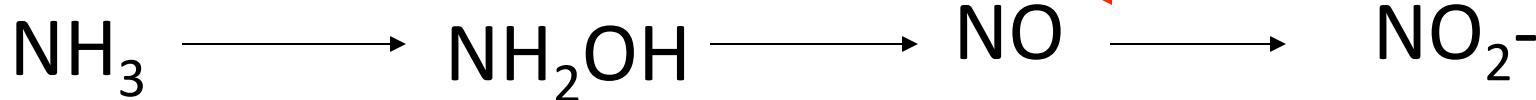


Ammonia oxidation is a potential source of marine N₂O

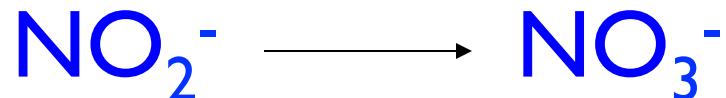
AOB:



AOA:

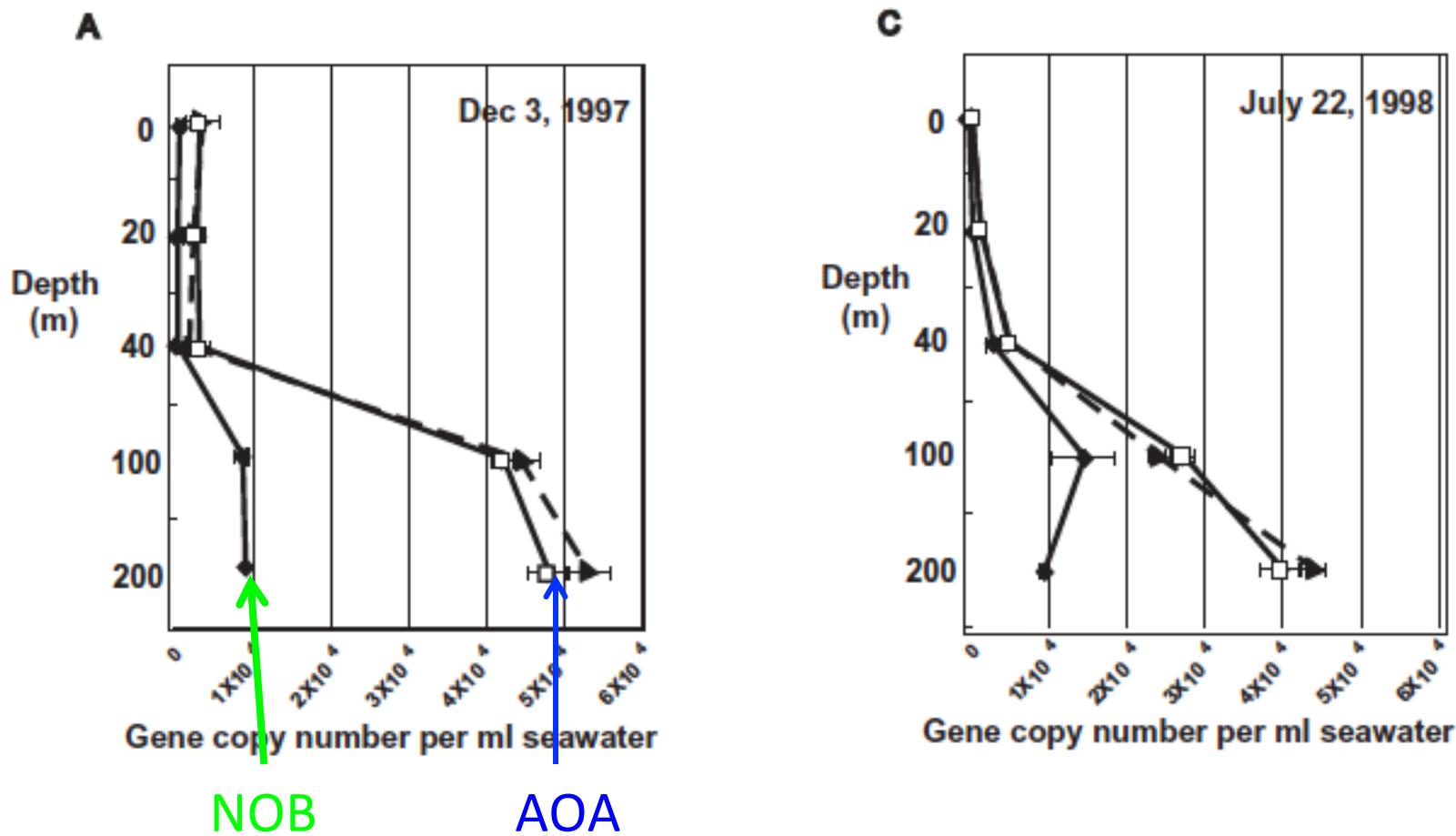


Nitrite oxidizers carry out the second step of nitrification



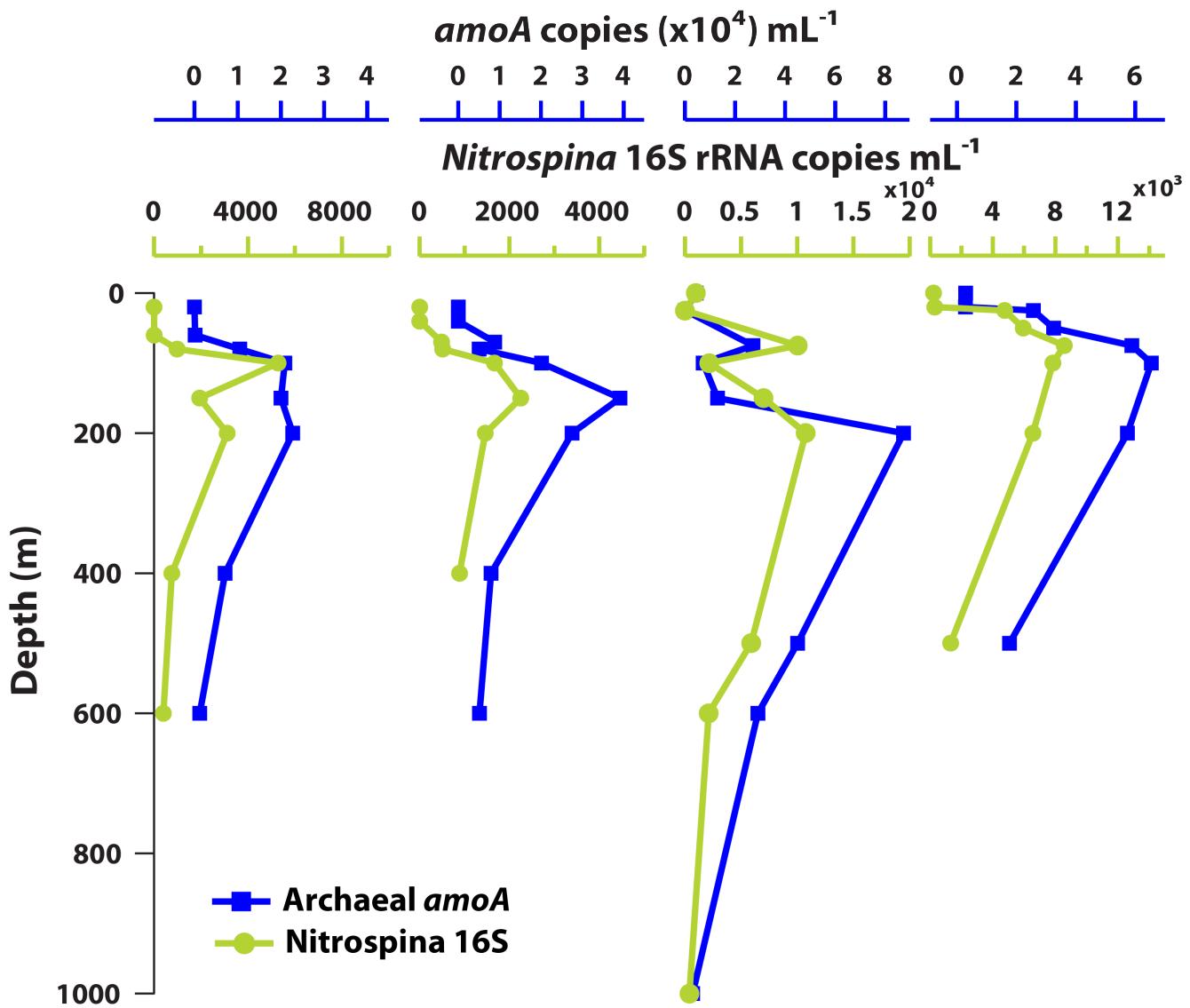
- Limited to only a few genera (*Nitrospira*, *Nitrospina*, *Nitrobacter*)
- Mostly autotrophs, but some capacity for mixotrophy
- Growing as autotrophs, they are living near the threshold for life ($\Delta G^0' = -74 \text{ kJ/mol}$)

Covariation of ammonia oxidizers (AOA) and nitrite oxidizers (NOB)



Mincer et al. 2007

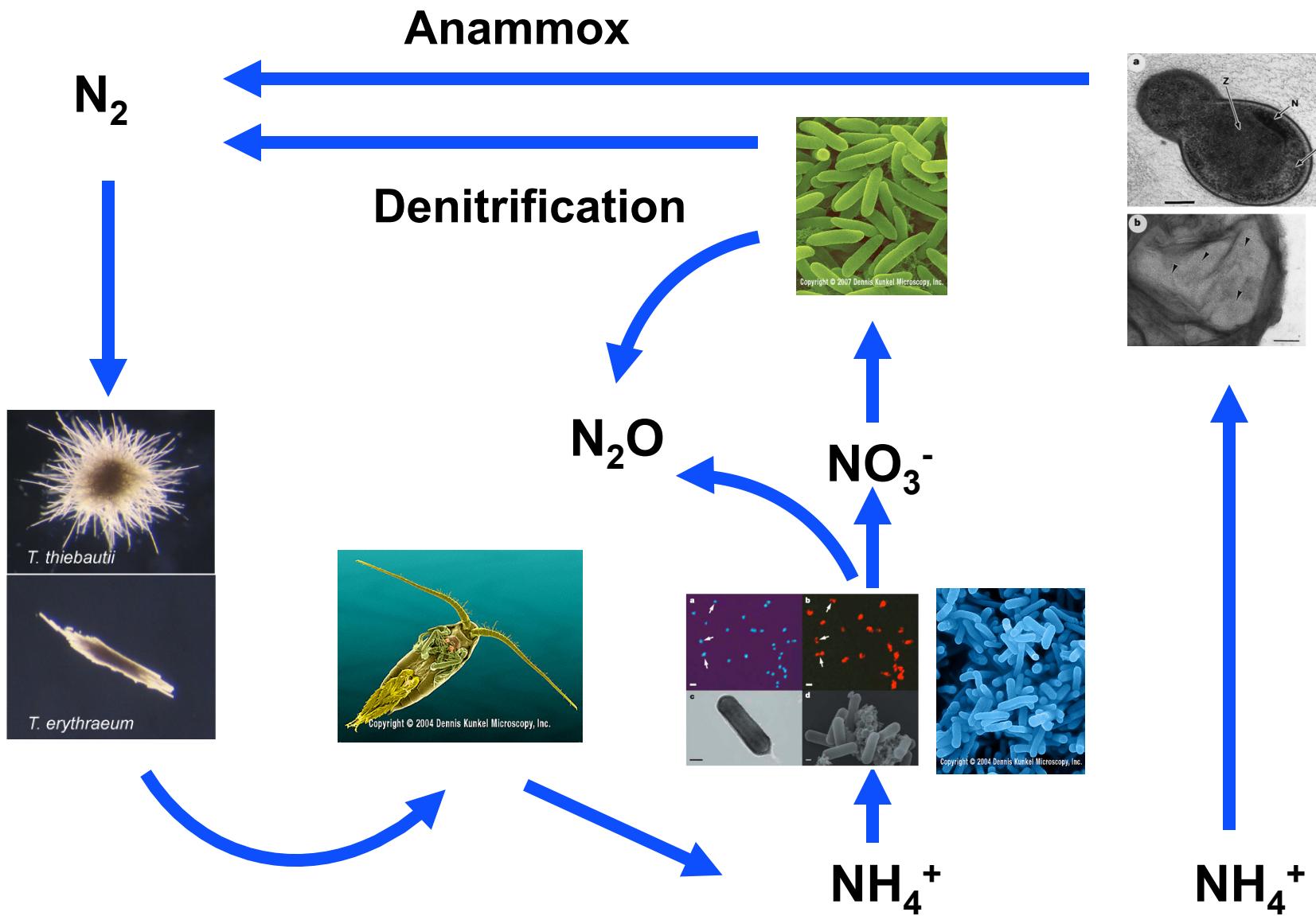
Covariation of AOA and NOB



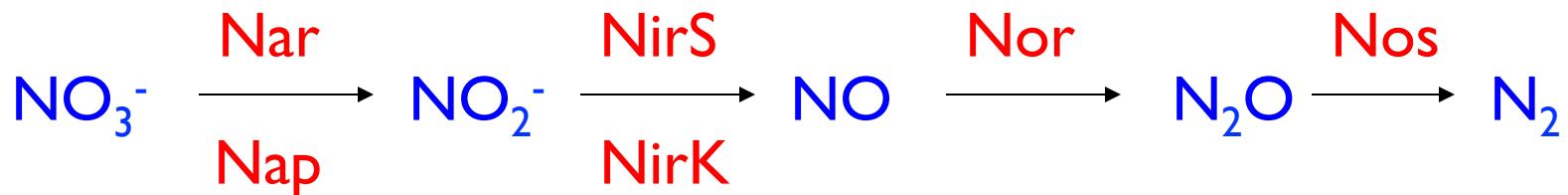
data from
CA current

Santoro et al. 2010

A condensed N cycle for today



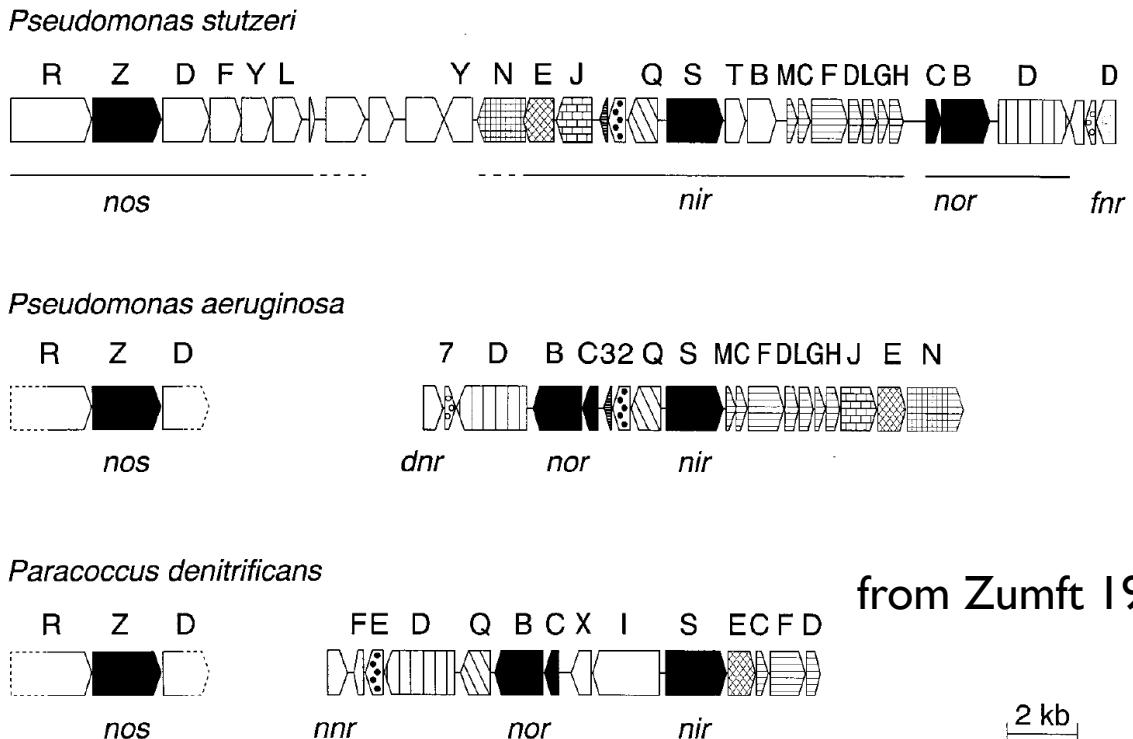
Denitrification is a multi-step process



- The dissimilatory reduction of nitrogen oxides to gaseous products under low-oxygen conditions
- A series of one electron transfers, not necessarily carried out by a single organism
- Frequently found in organisms capable of using other electron acceptors, including oxygen

Denitrification genes are a “necklace”

Genes are not congruent with 16S rRNA phylogeny (laterally transferred)



from Zumft 1997

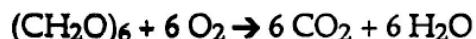
Maybe it's more of a charm bracelet . .

TABLE 11. The metabolic diversity of archaeal and bacterial genera harboring denitrifying species

Archaea	Bacteria (gram-negative)
Organotrophic	Diazotrophic
Halophilic	<i>Aquaspirillum</i>
<i>Halocarcula</i>	<i>Azospirillum</i>
<i>Halobacterium</i>	<i>Azoarcus</i>
<i>Haloferox</i>	<i>Bacillus</i>
Hyperthermophilic	<i>Bradyrhizobium</i>
<i>Pyrobaculum</i>	<i>Pseudomonas</i>
	<i>Rhodobacter</i>
Bacteria (gram-positive)	<i>Rhodopseudomonas</i>
Organotrophic	<i>Sinorhizobium</i>
Spore forming	Thermophilic
<i>Bacillus</i>	<i>Aquifex</i>
Nonspore forming	<i>Bacillus</i>
<i>Jonesia</i>	<i>Thermothrix</i>
Bacteria (gram-negative)	Psychrophilic
Phototrophic	<i>Aquaspirillum</i>
<i>Rhodobacter</i>	<i>Halomonas</i>
<i>Rhodopseudomonas</i>	Halophilic
<i>Rhodoplanes</i>	<i>Halomonas</i>
Lithotrophic	<i>Bacillus</i>
S oxidizing	Pigment-forming
<i>Beggiatoa</i>	<i>Chromobacterium</i>
<i>Thiobacillus</i>	<i>Flavobacterium</i>
<i>Thioploca</i>	<i>Pseudomonas</i>
H ₂ oxidizing	Budding
<i>Ralstonia</i>	<i>Blastobacter</i>
<i>Paracoccus</i>	<i>Hyphomicrobium</i>
<i>Pseudomonas</i>	Gliding
NO ₂ ⁻ or NH ₄ ⁺ oxidizing	<i>Cytophaga</i>
<i>Nitrobacter</i>	<i>Flexibacter</i>
<i>Nitrosomonas</i>	Magnetotactic
Organotrophic	<i>Magnetospirillum</i>
Carboxidotropic	Pathogenic
<i>Pseudomonas</i>	<i>Achromobacter</i>
<i>Zavarzinia</i>	<i>Alcaligenes</i>
Oligocarbophilic	<i>Agrobacterium</i>
<i>Aquaspirillum</i>	<i>Campylobacter</i>
<i>Hyphomicrobium</i>	<i>Eikenella</i>
Fermentative	<i>Flavobacterium</i>
<i>Empedobacter</i>	<i>Kingella</i>
<i>Azospirillum</i>	<i>Moraxella</i>
Facultatively anaerobic	<i>Morococcus</i>
<i>Alteromonas</i>	<i>Neisseria</i>
<i>Pseudomonas</i>	<i>Ochrobactrum</i>
Aerobic	<i>Oligella</i>
<i>Paracoccus</i>	<i>Pseudomonas</i>
<i>Alcaligenes</i>	<i>Sphingobacterium</i>
	<i>Tsukamurella</i>

COUPLED TO GLUCOSE UTILIZATION

Aerobic respiration:



$$\Delta G^{\circ\prime} = -2870 \text{ kJ} \quad (-686 \text{ kcal})$$

1. Complete denitrification:



$$\Delta G^{\circ\prime} = -2669 \text{ kJ} \quad (-638 \text{ kcal})$$

2. Nitrate respiration (to nitrite only)

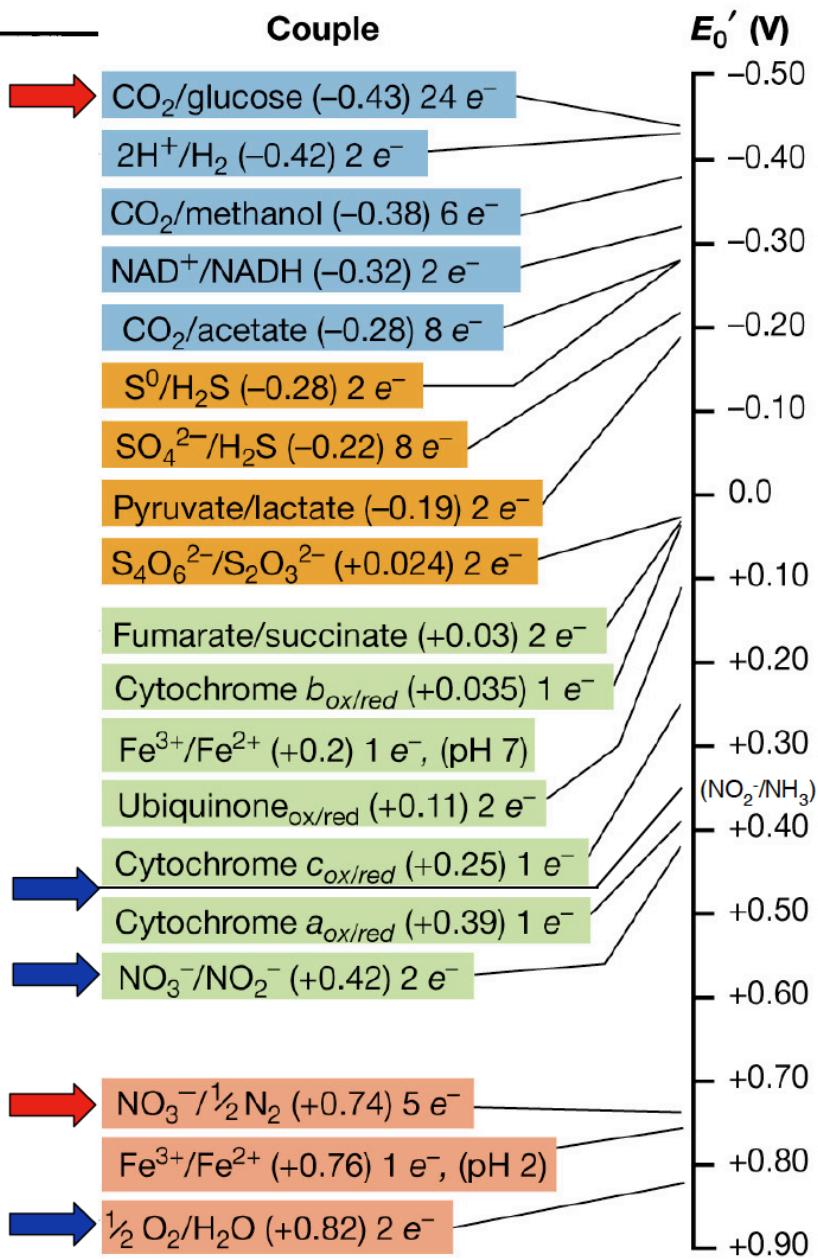


$$\Delta G^{\circ\prime} = -1766 \text{ kJ} \quad (-422 \text{ kcal})$$

3. Nitrate reduction to ammonia



$$\Delta G^{\circ\prime} = -1796 \text{ kJ} \quad (-429 \text{ kcal})$$

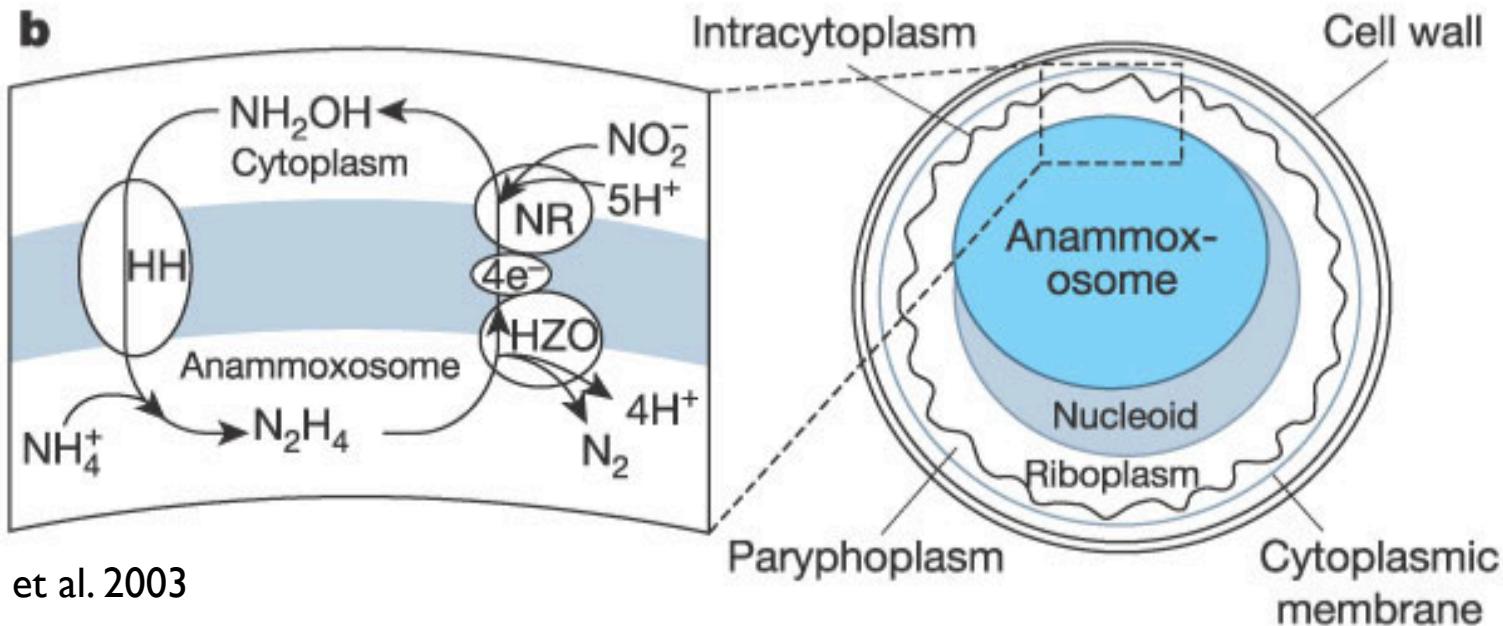
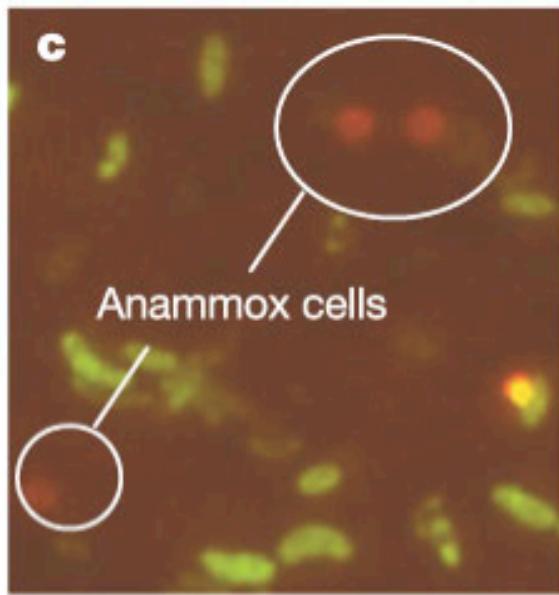
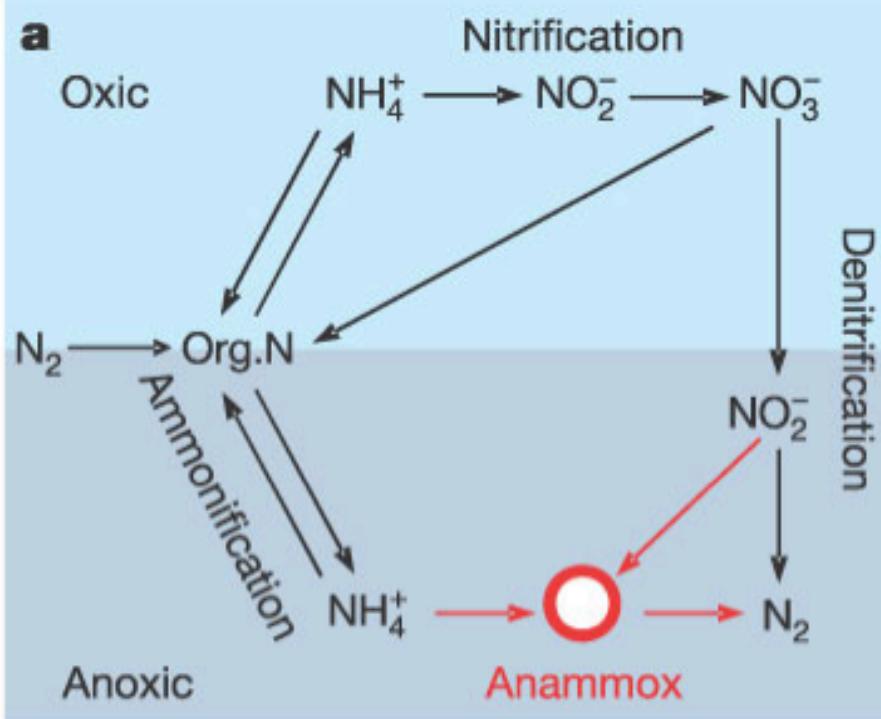


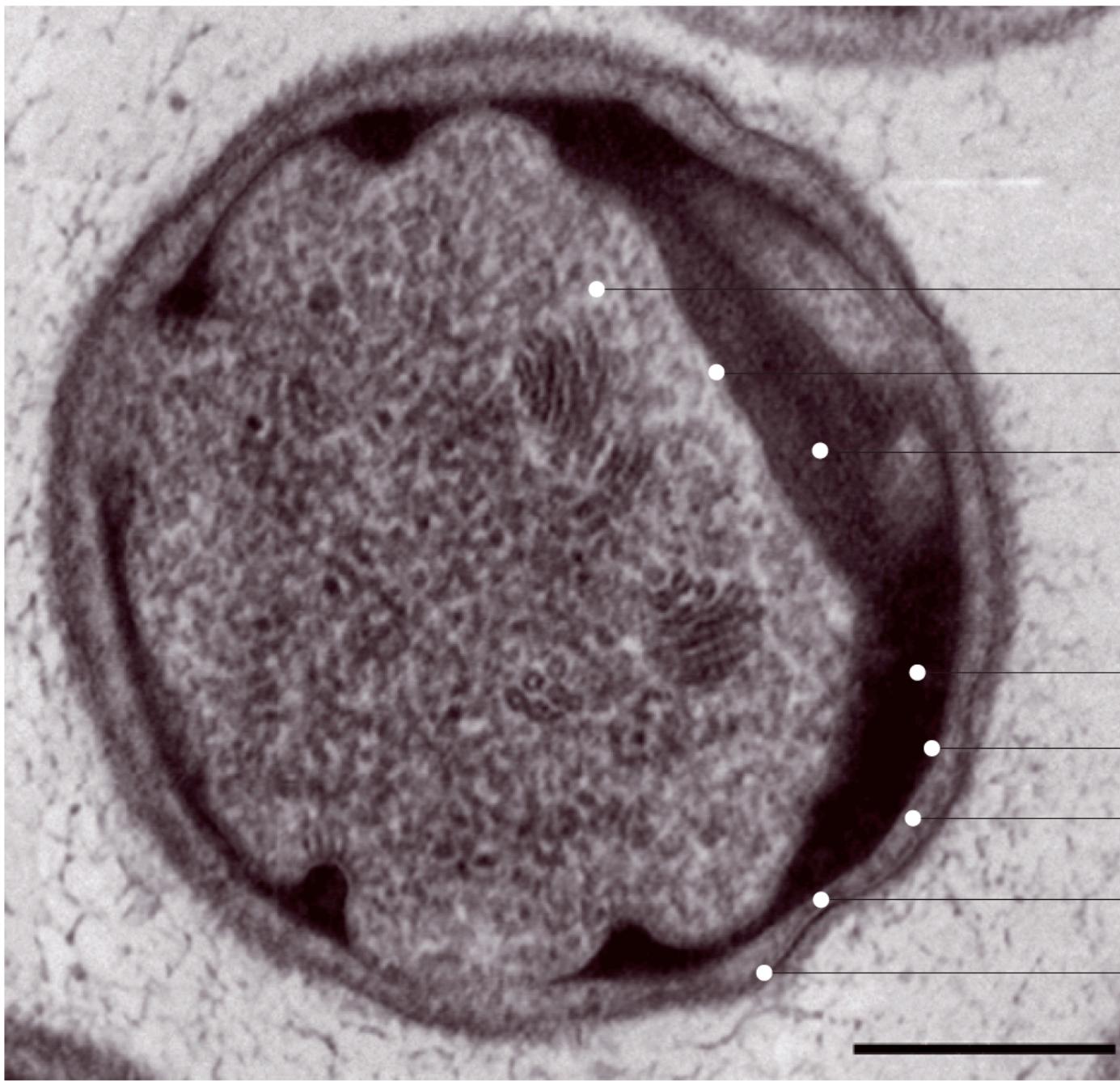
Anaerobic ammonia oxidation (anammox) solves a marine conundrum (?)



- Based on stoichiometric breakdown of Redfield organic matter, there should be a lot more NH_4^+ in anoxic environments than there actually is.
- ‘Discovered’ in 1985, but the responsible bacteria were not identified until 1999
- Chemoautotrophic organisms with an enormous iron requirement.

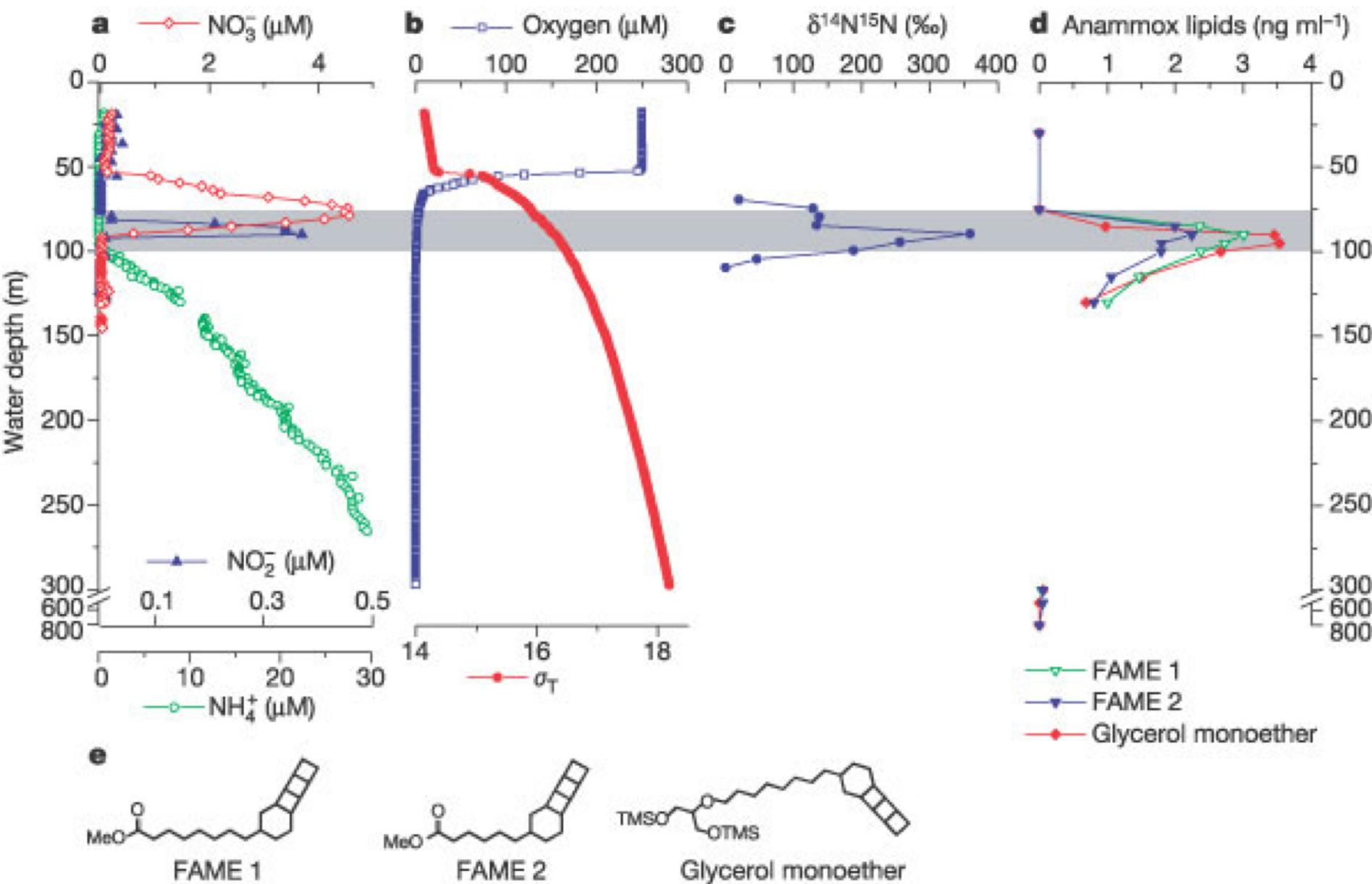




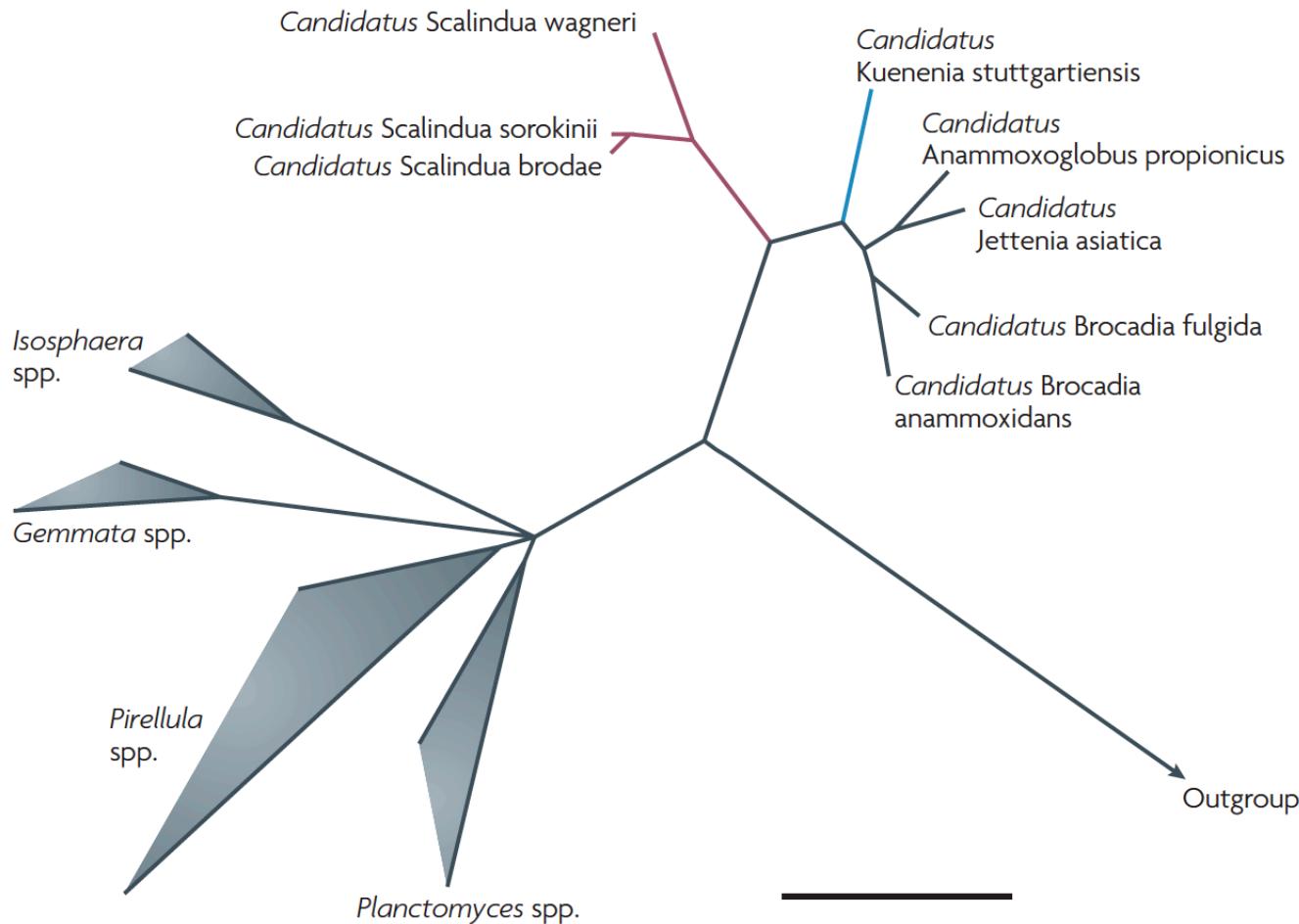


- Anammoxosome
- Anammoxosome membrane
- Nucleoid
- Riboplasm
- Intracytoplasmic membrane
- Paryphoplasm
- Cytoplasmic membrane
- Cell wall

Marine anammox ‘discovered’ in the Black Sea



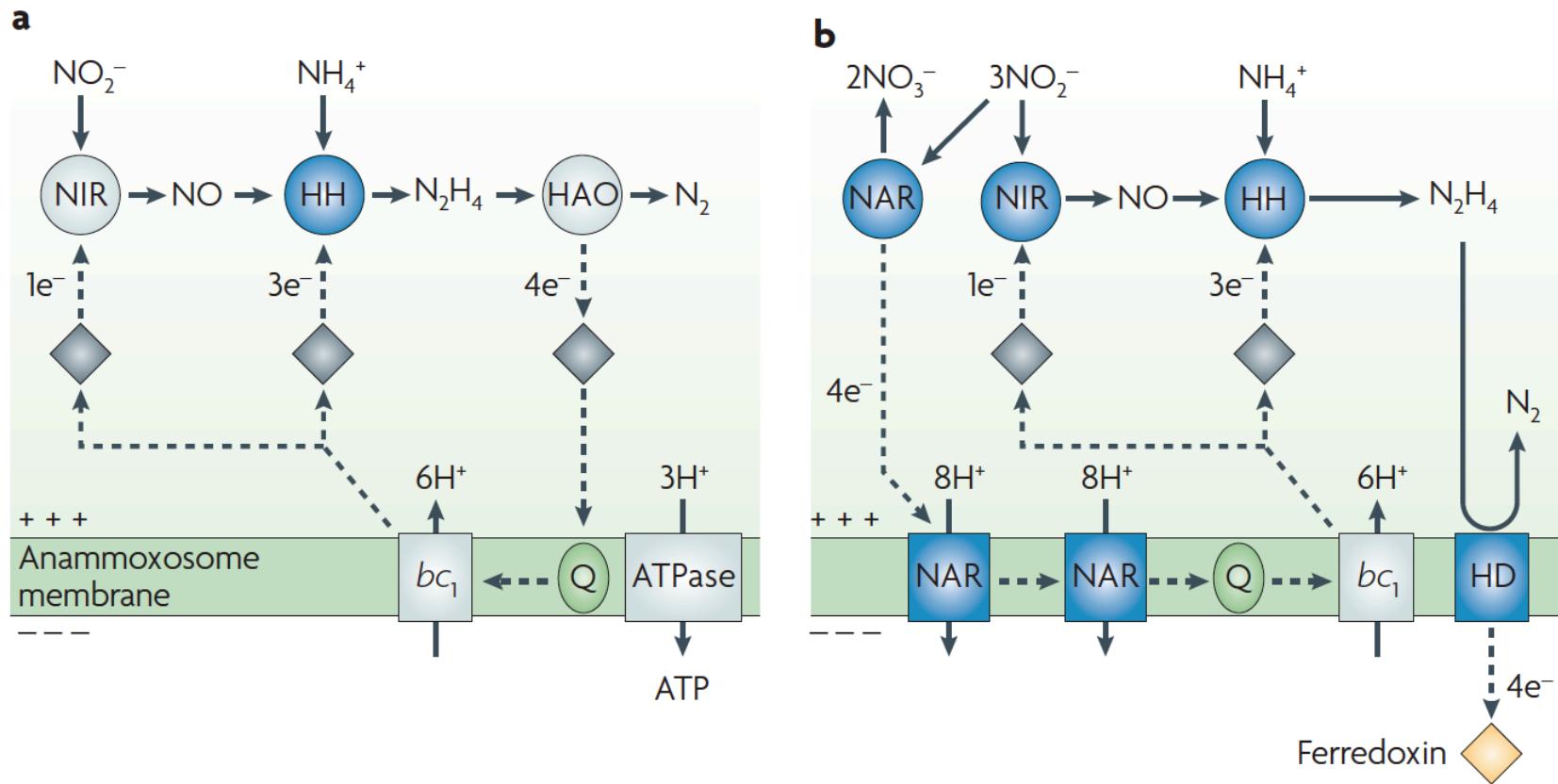
Anammox bacteria are highly divergent from their nearest neighbors



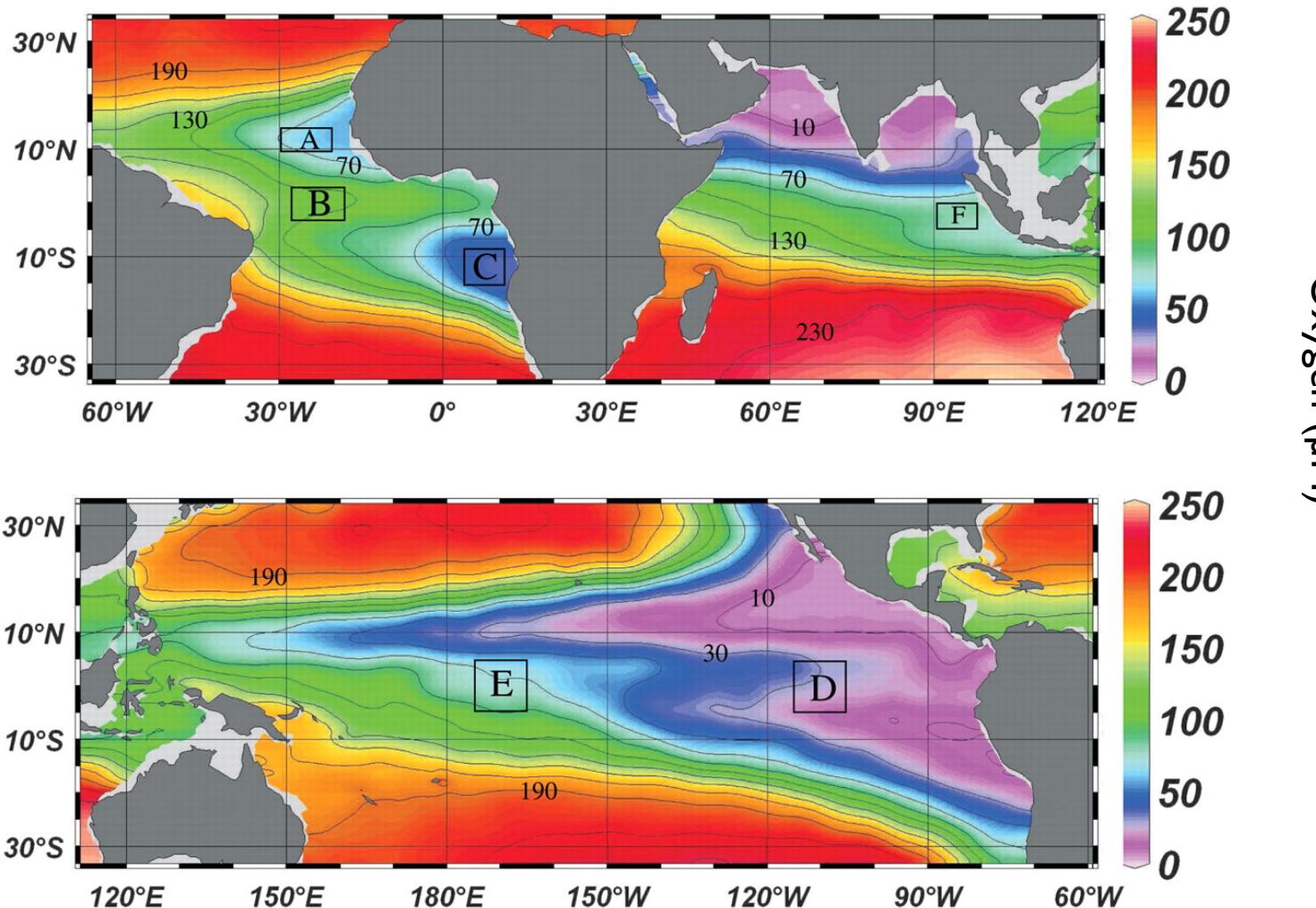
Kuenen 2008

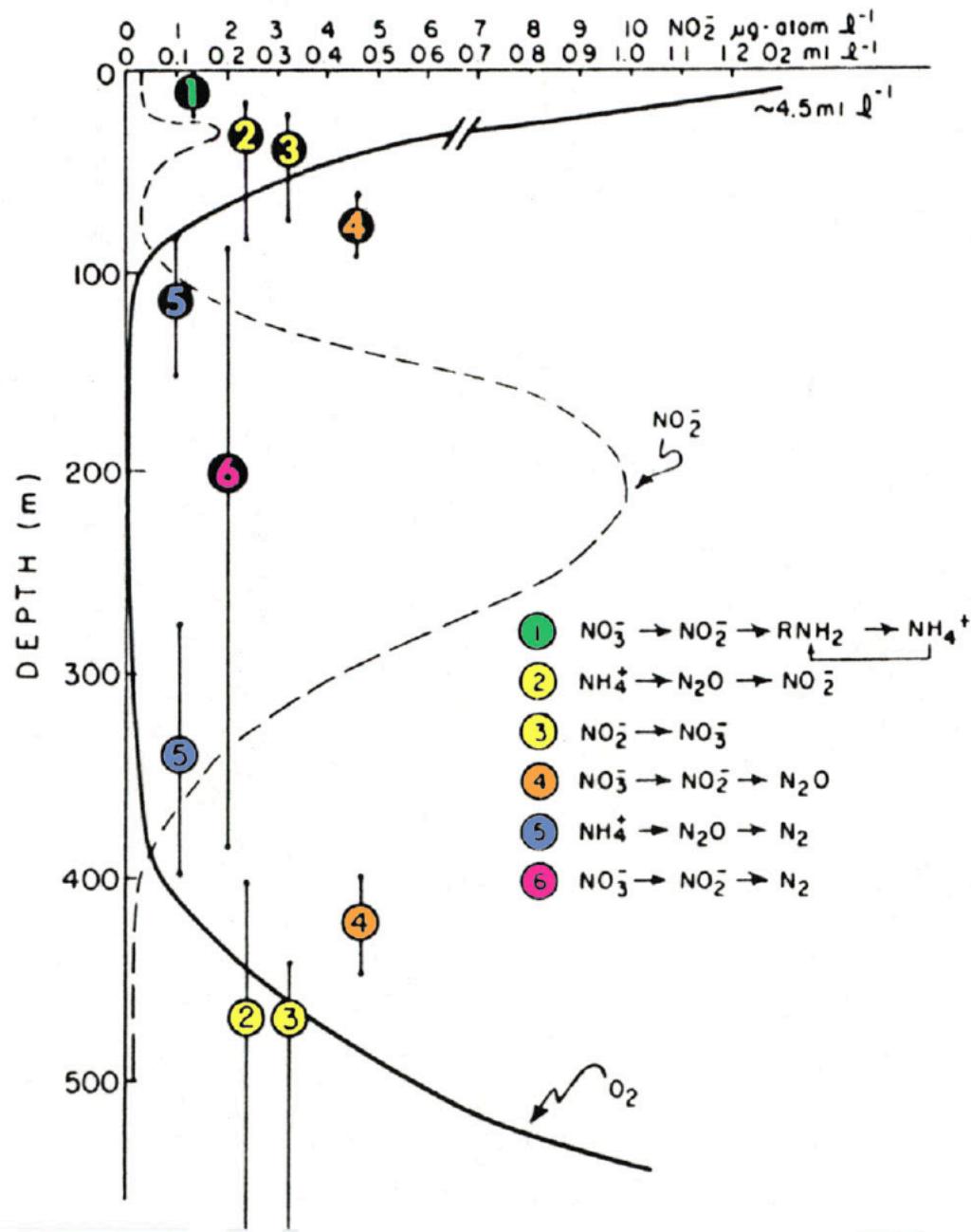
Several functional gene targets for anammox bacteria

Candidates: *hao* (hydrazine oxidoreductase); *nirS*



Putting it all together: Synthetic studies of the marine nitrogen cycle





Oxygen minimum zones were ripe for integrated molecular and biogeochemical studies.

(Codispoti and Christiansen, 1985)

Revising the nitrogen cycle in the Peruvian oxygen minimum zone

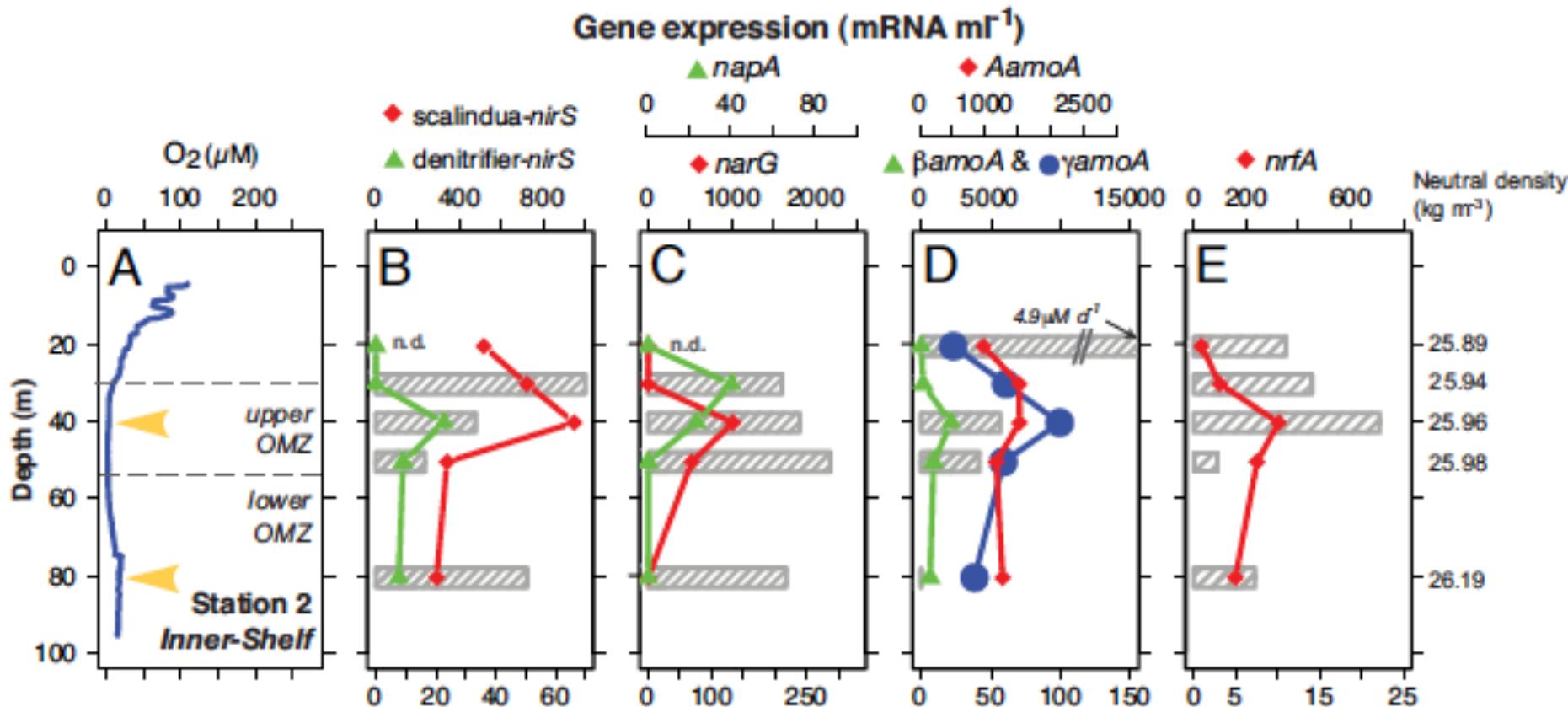
Phyllis Lam^{a,1}, Gaute Lavik^a, Marlene M. Jensen^{a,2}, Jack van de Vossenberg^b, Markus Schmid^{b,3}, Dagmar Woebken^{a,4}, Dimitri Gutiérrez^c, Rudolf Amann^a, Mike S. M. Jetten^b, and Marcel M. M. Kuypers^a

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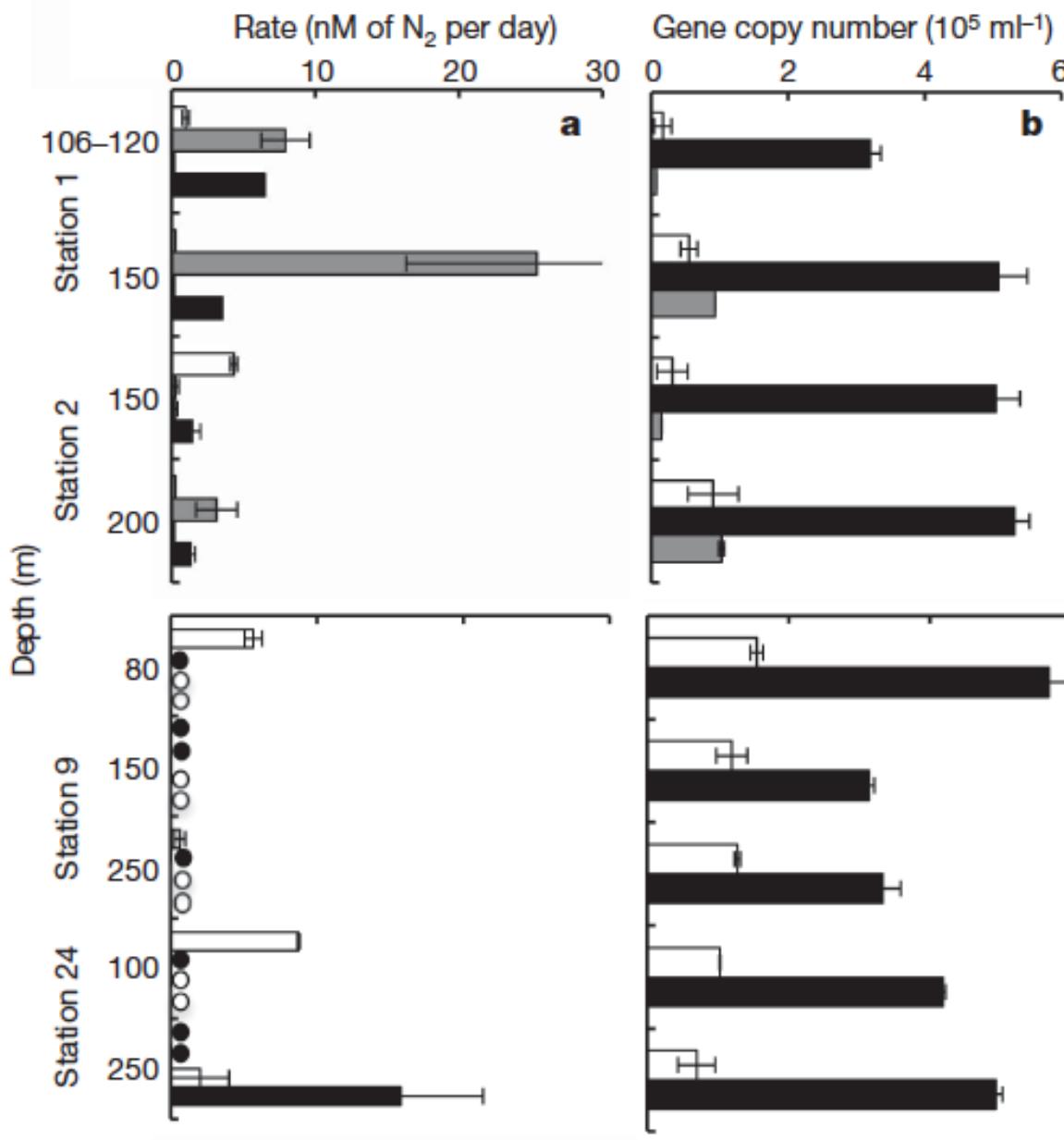
LETTERS

Denitrification as the dominant nitrogen loss process in the Arabian Sea

B. B. Ward¹, A. H. Devol², J. J. Rich³, B. X. Chang², S. E. Bulow¹, Hema Naik⁴, Anil Pratihary⁴ & A. Jayakumar¹



But beware the ‘invisible present’ . . .



Ward et al. 2009

Organic matter C:N sets nitrogen loss pathway

