

# **Extreme diversity and abundance of planktonic diplonemids in the world oceans**

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*Station Biologique de Roscoff, Roscoff, France*

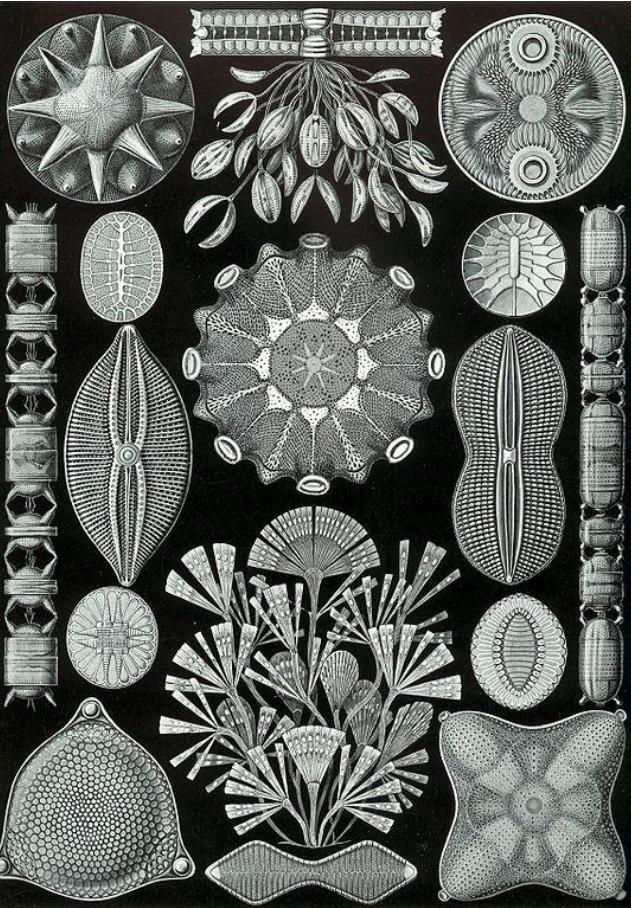
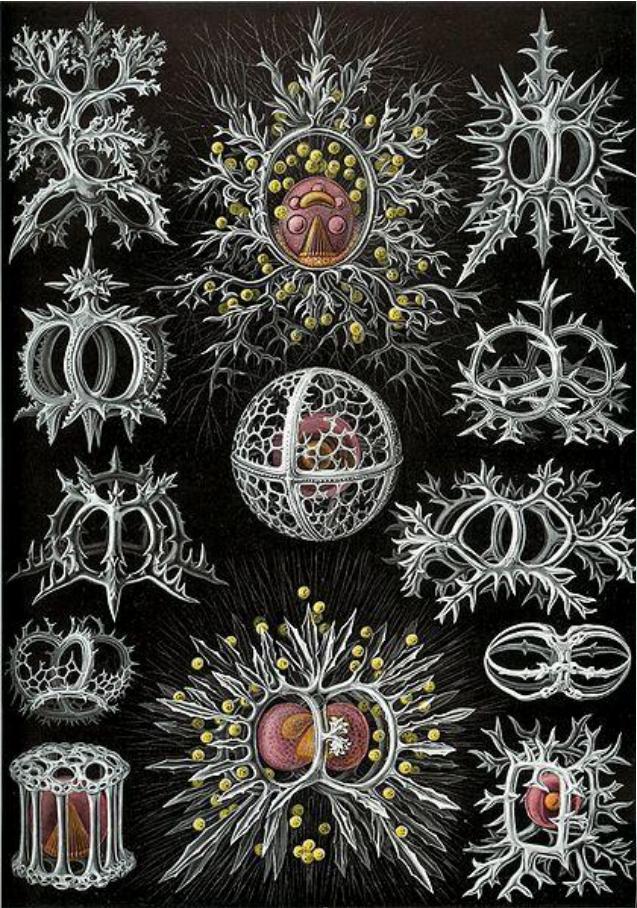
**Funding: EU 7<sup>th</sup> FP (FP7/2007-2013) , Czech Academy, Tara Oceans, Oceanomics**



# BIODIVERSITY

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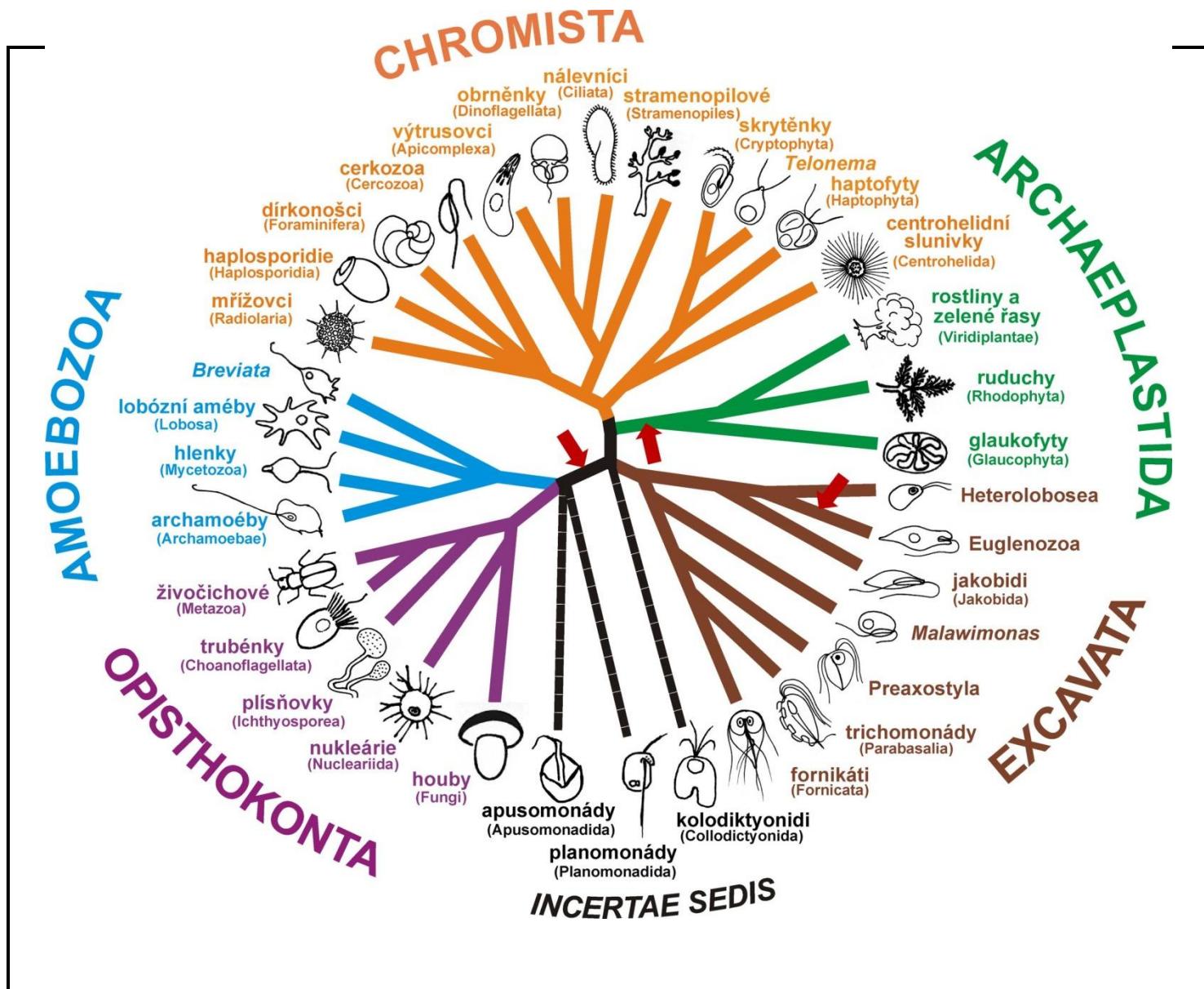


# Biodiversity of world ocean?



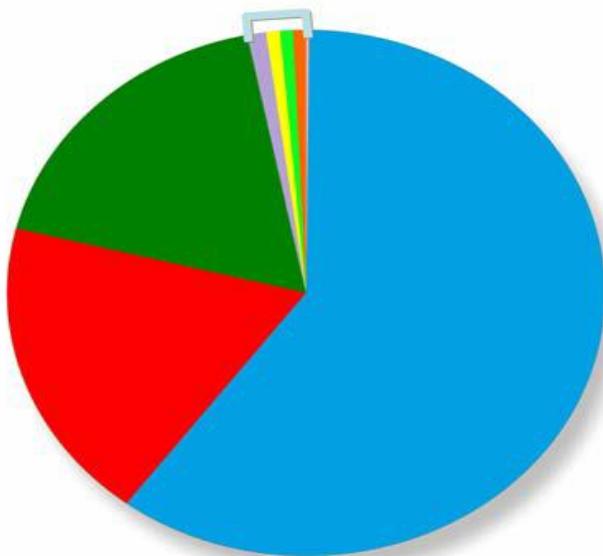


# Current view of eukaryotic diversity

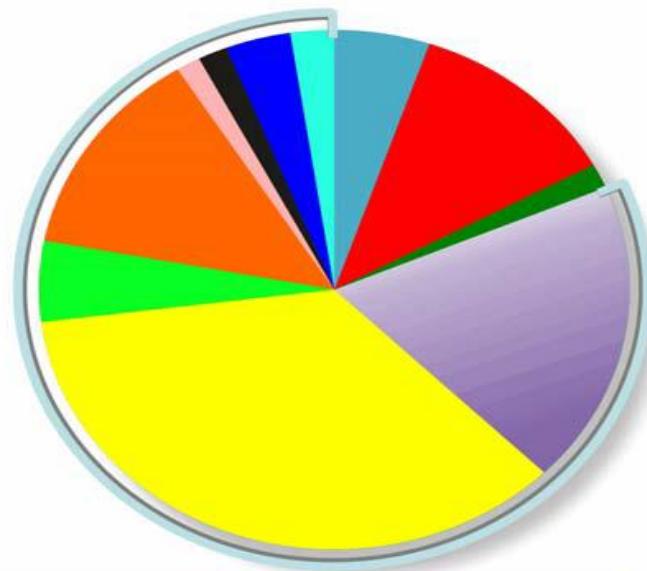


**Enormous diversity in number of species**

A. Catalogued species  
(N<sub>tot</sub> ≈ 2 million)

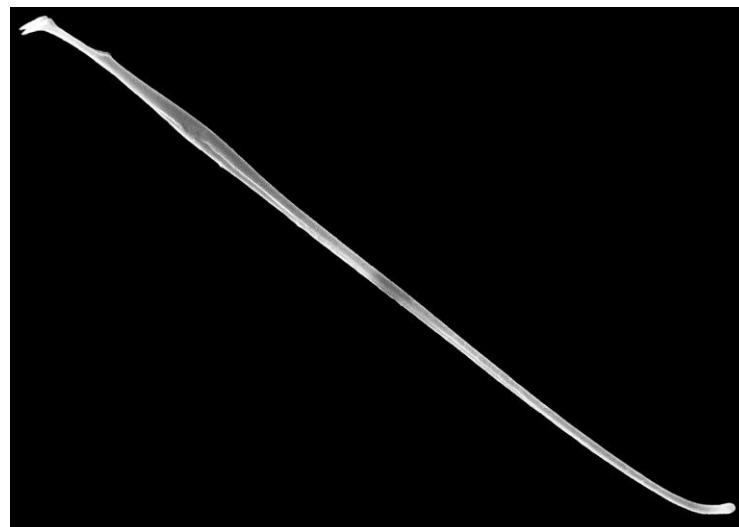
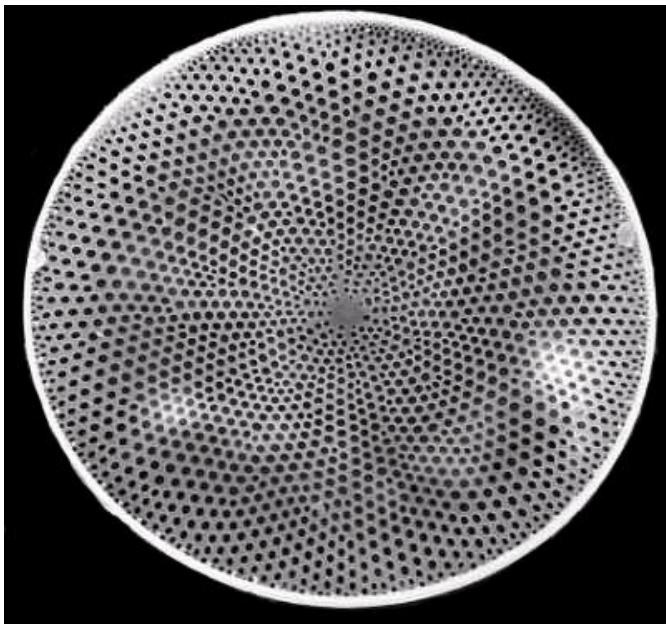
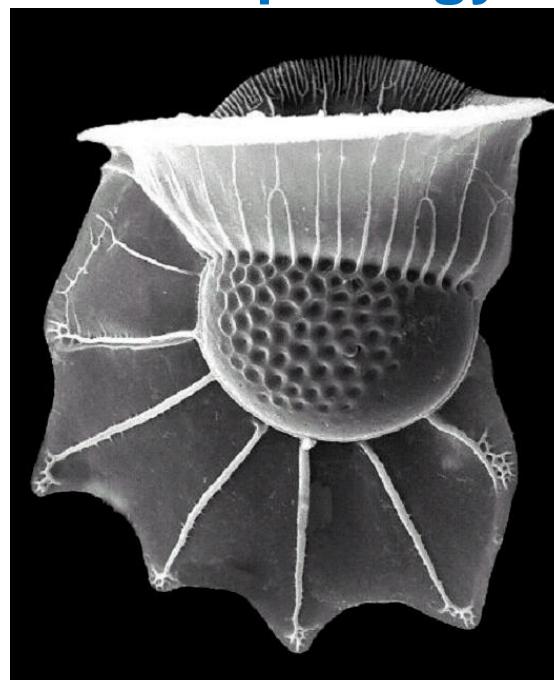
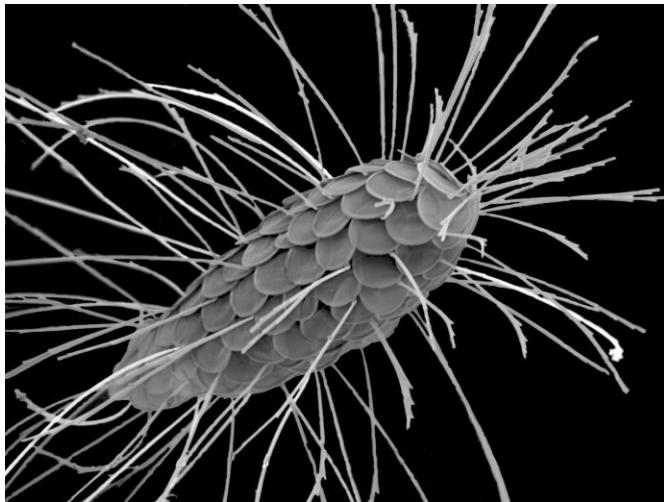


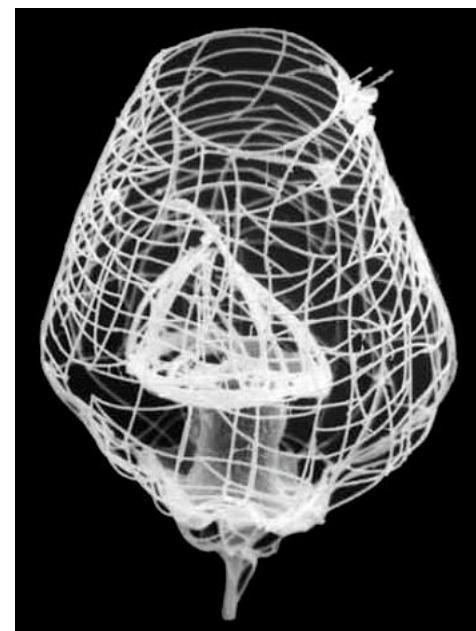
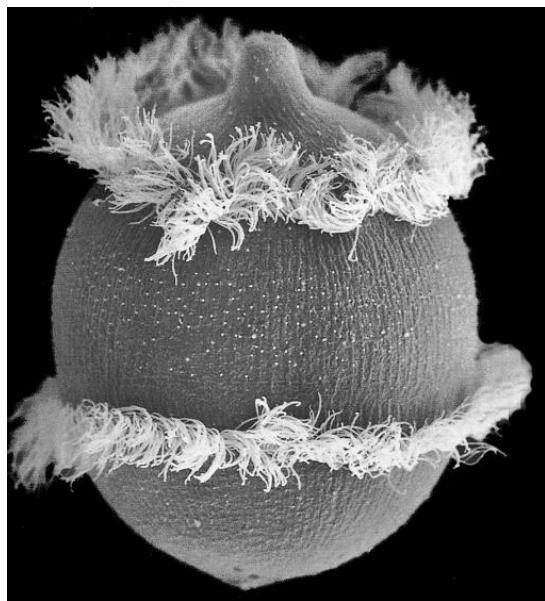
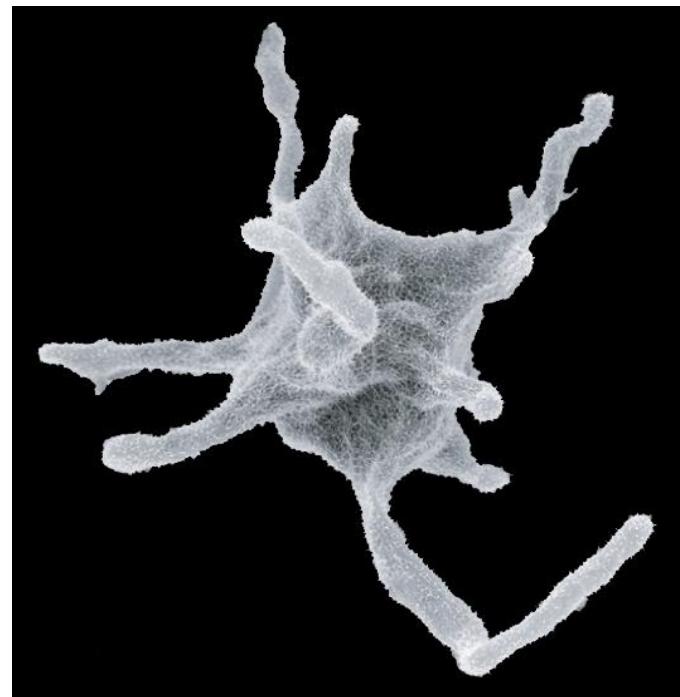
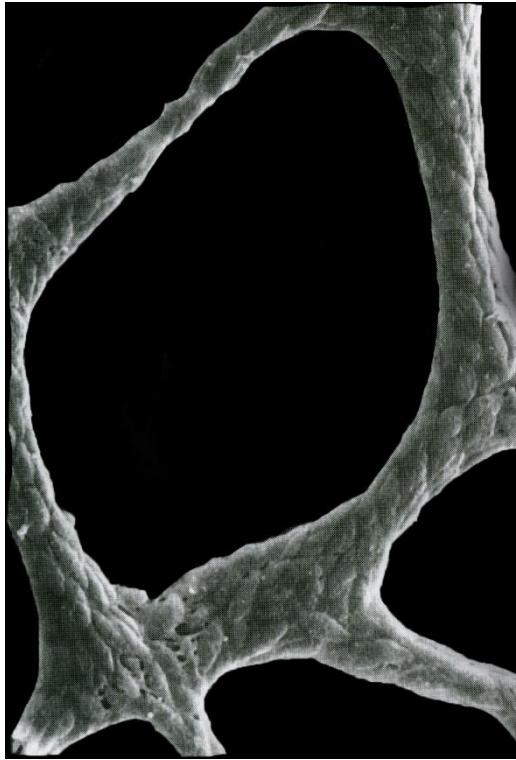
B. Environmental OTUs  
(1430 18S V4 rDNA 97%)

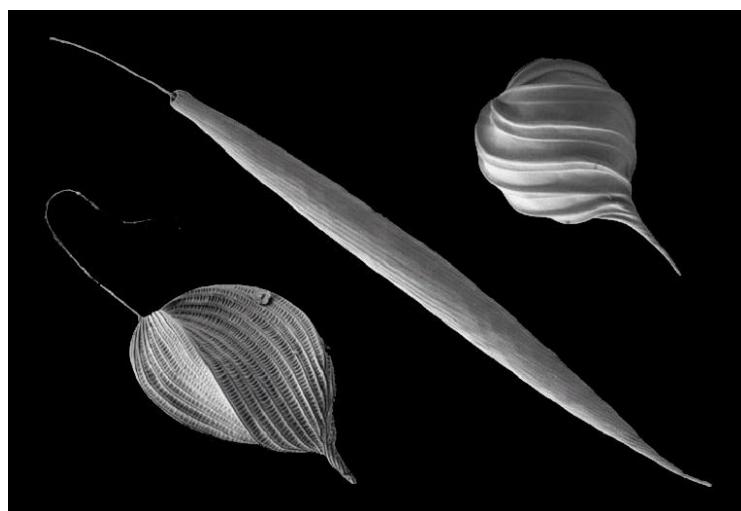
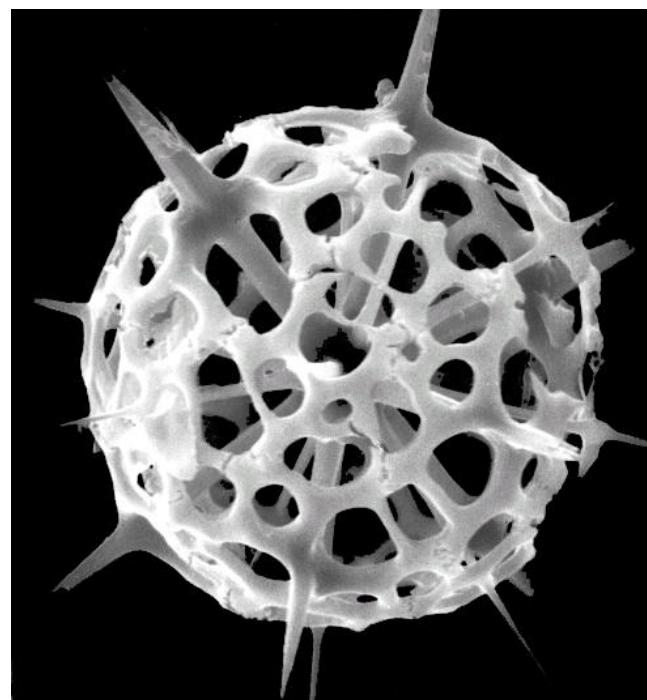
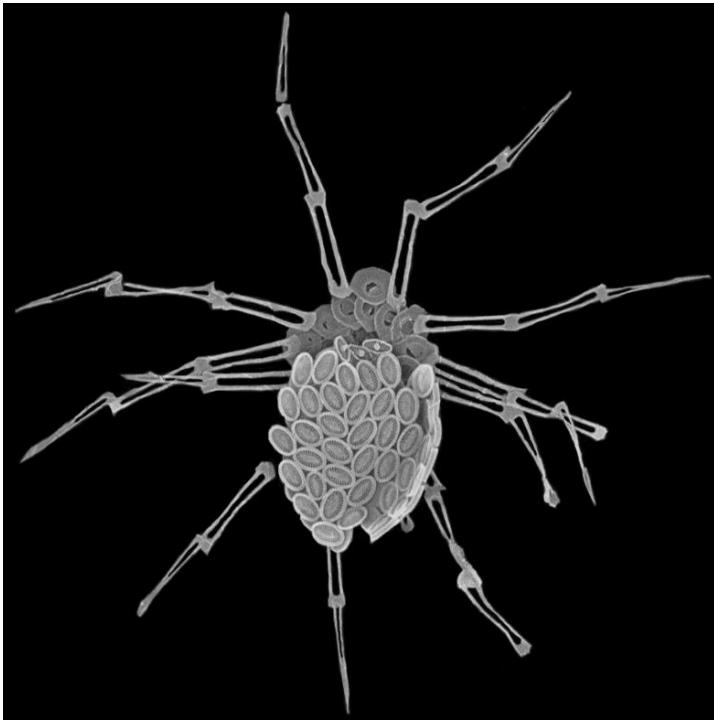


Morphological versus genetic views of total eukaryotic diversity. (A) Relative numbers of described species per eukaryotic supergroup. (B) Relative number of V4 18S rDNA Operational Taxonomic Units (97%) per eukaryotic supergroup, based on 59 rDNA clone library surveys of marine, fresh-water, and terrestrial total eukaryotic biodiversity.

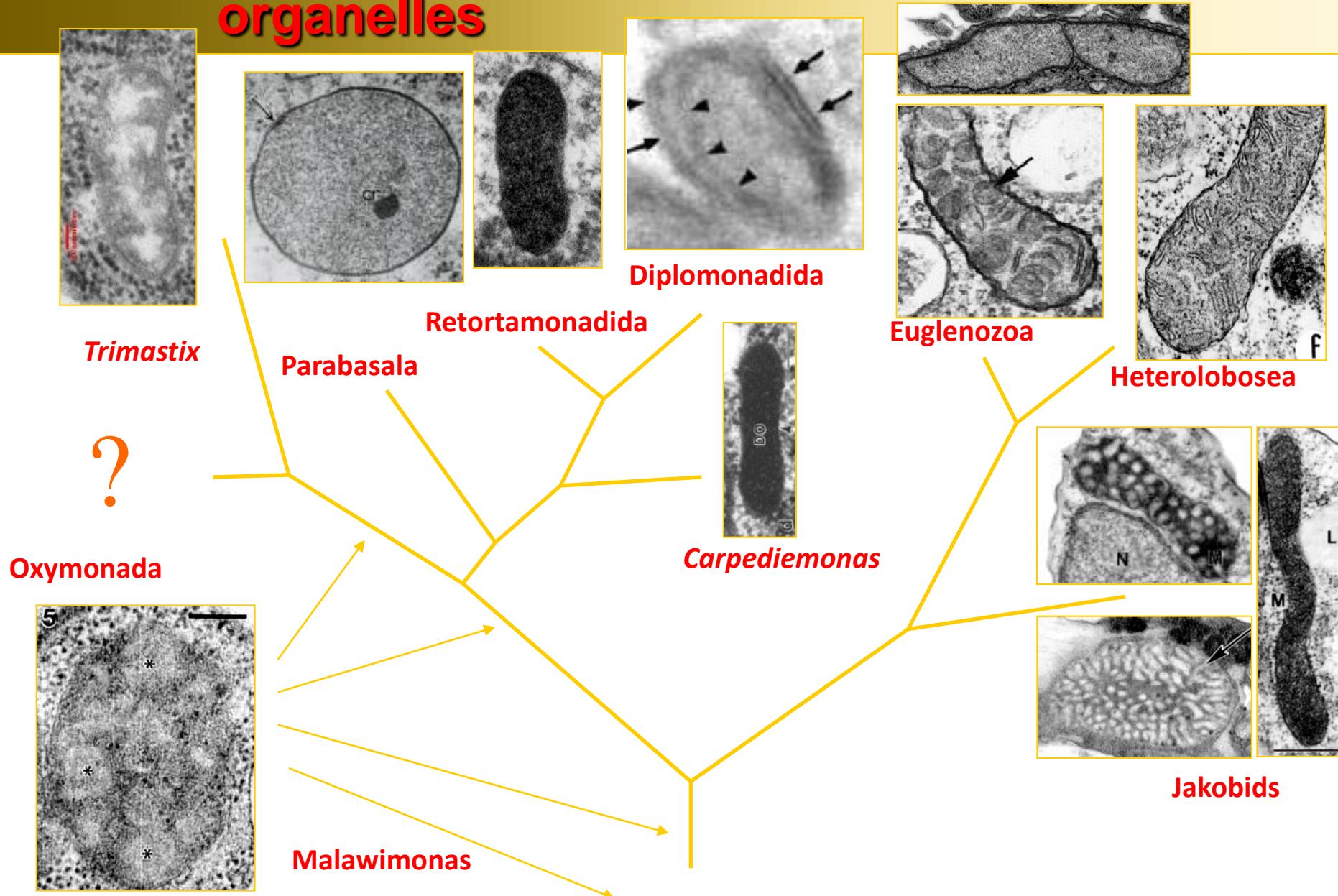
# Enormous diversity of protists in morphology



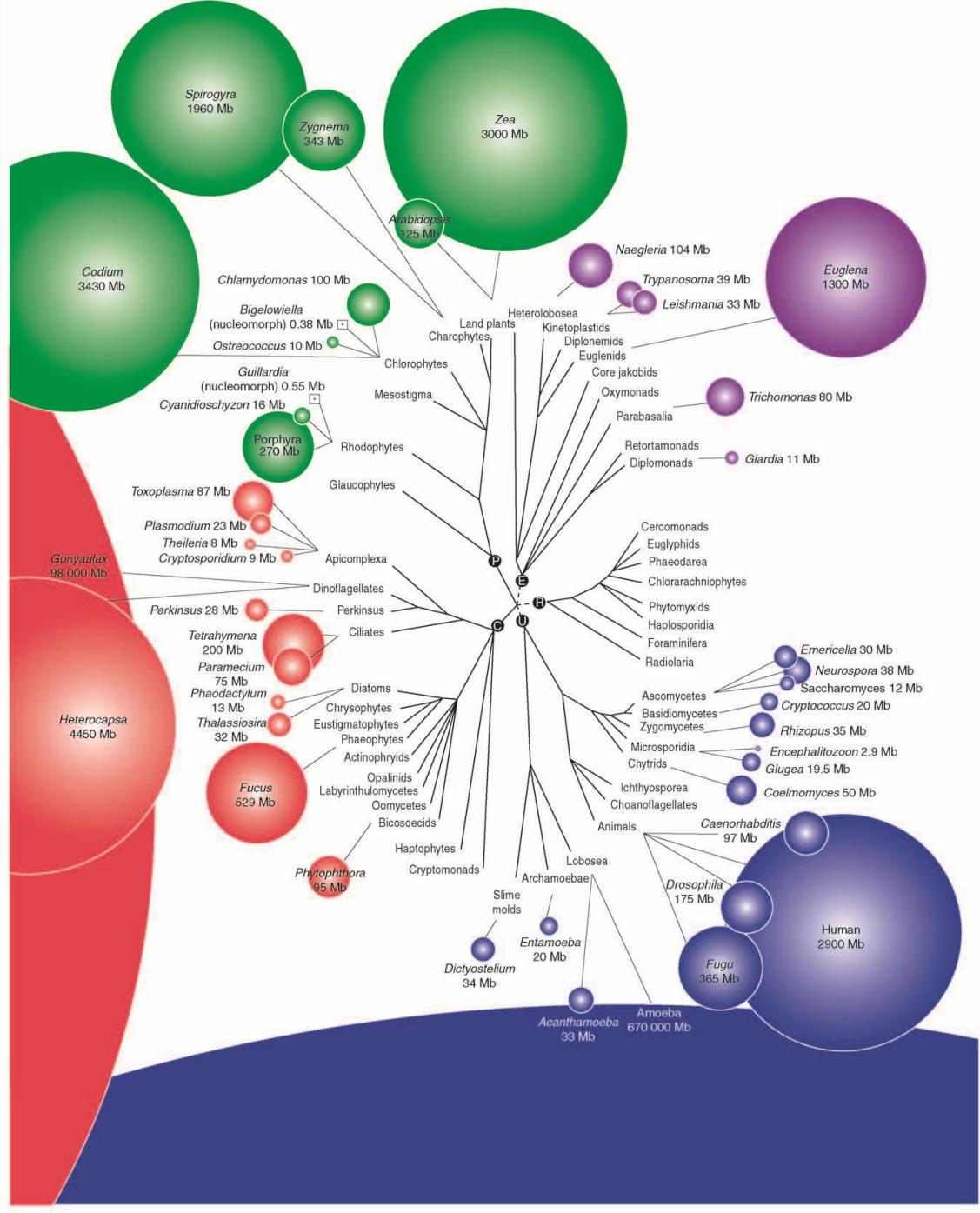




# Unexpected diversity of mitochondrion-derived organelles



# Enormous diversity of genome size in protists





# Eric Karsenti & Etienne Bourgeois



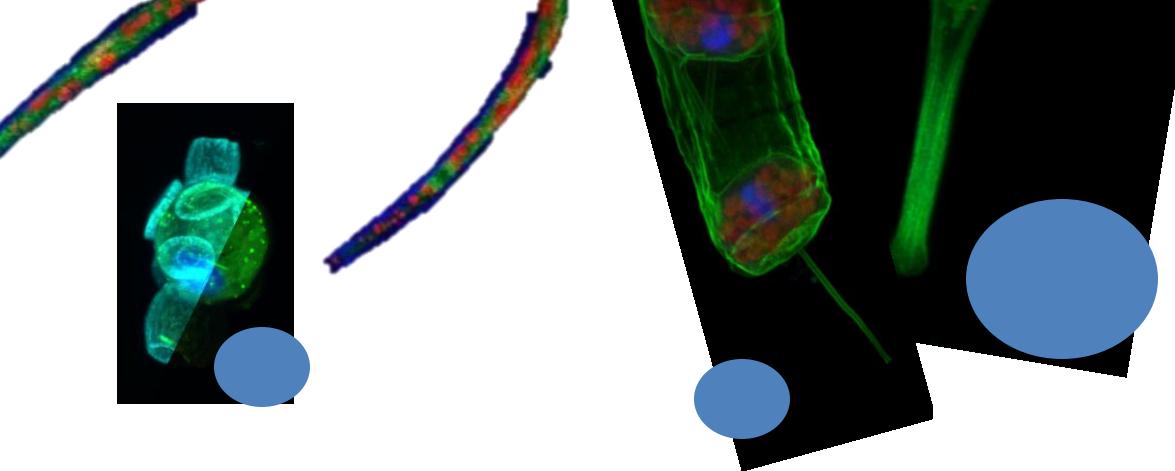
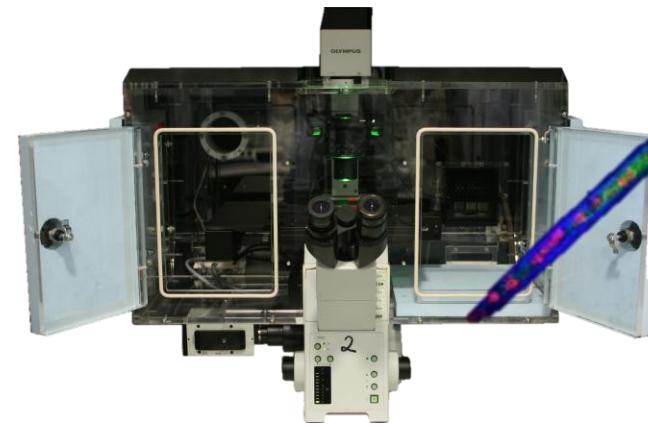
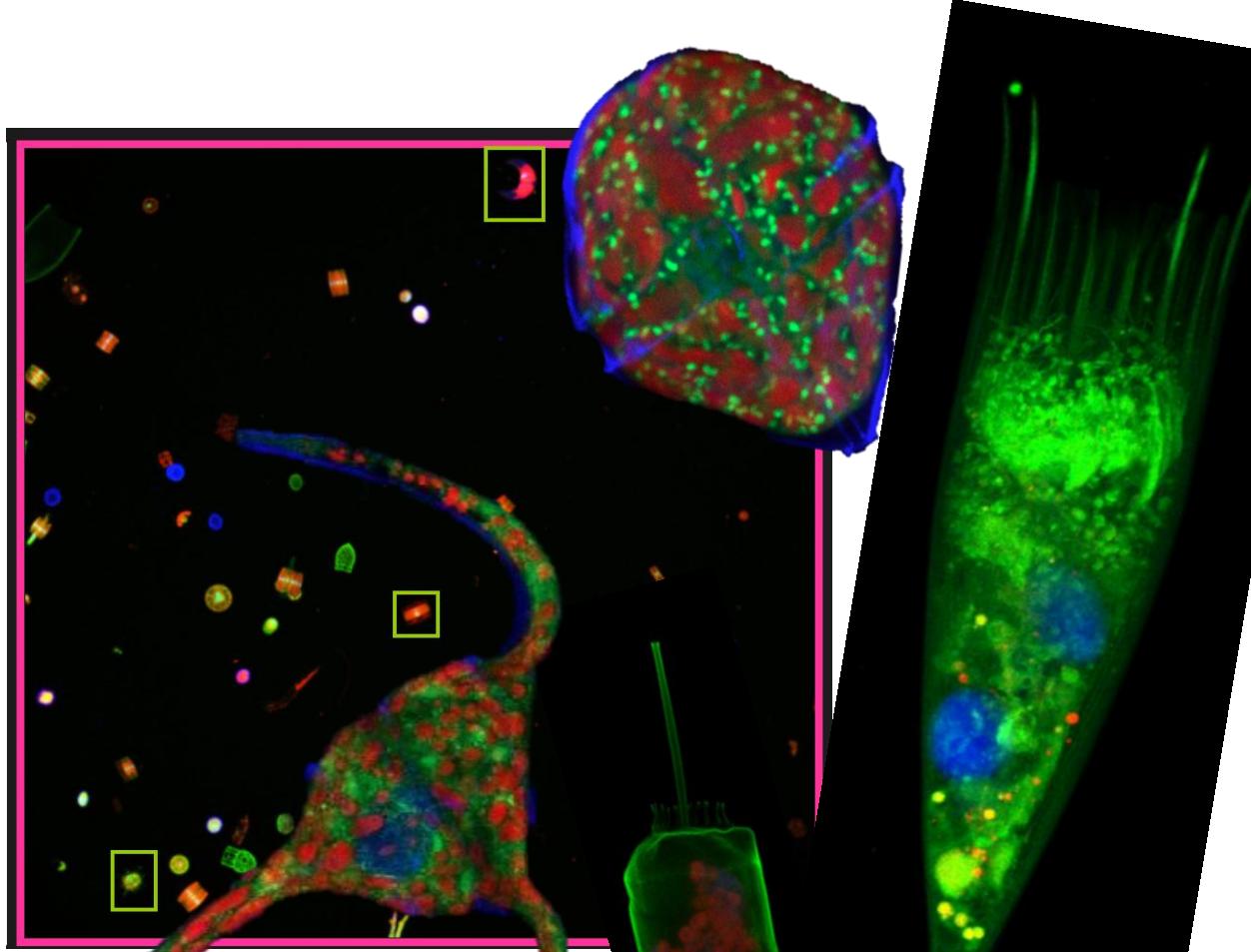
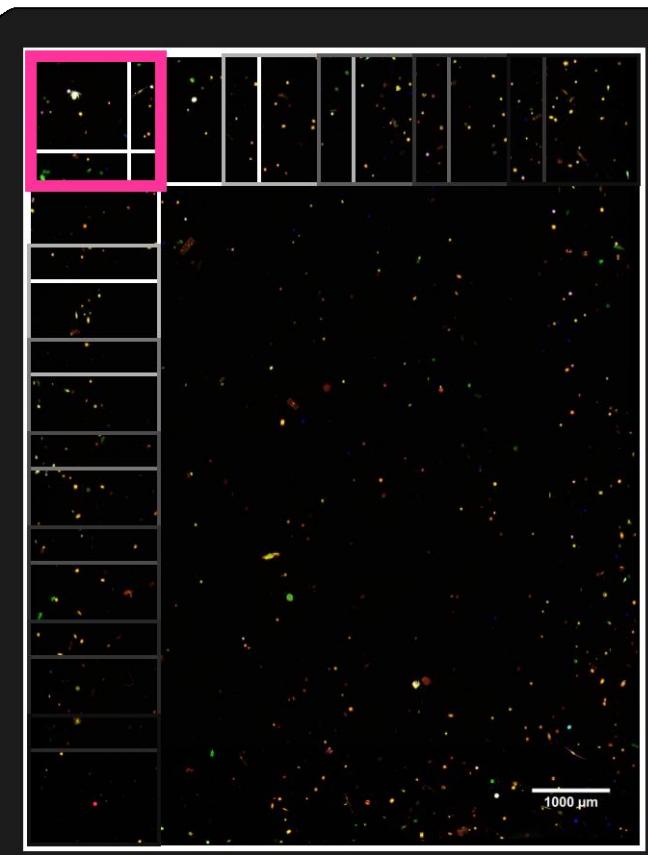
# TARA OCEANS

2009 - 2012



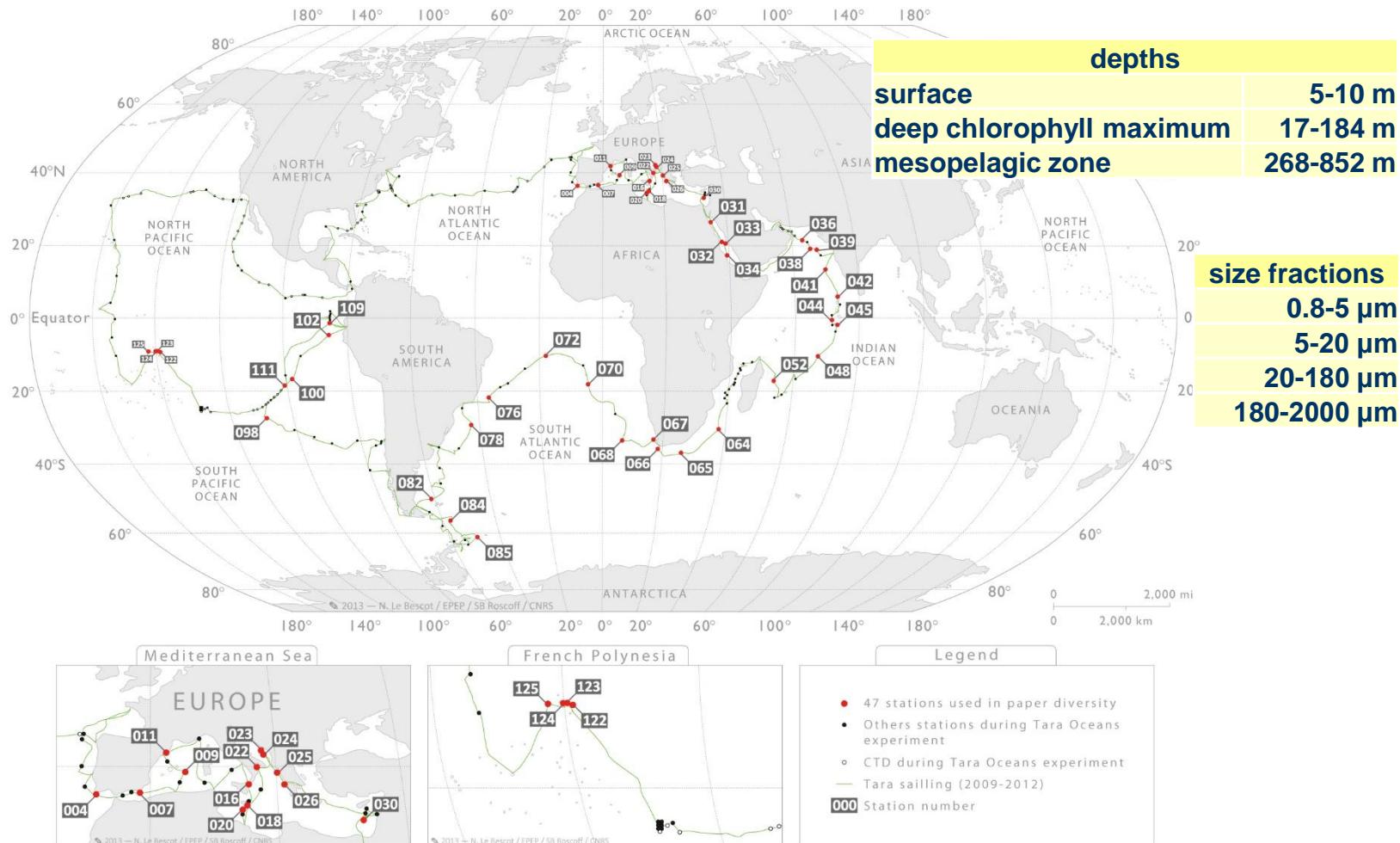




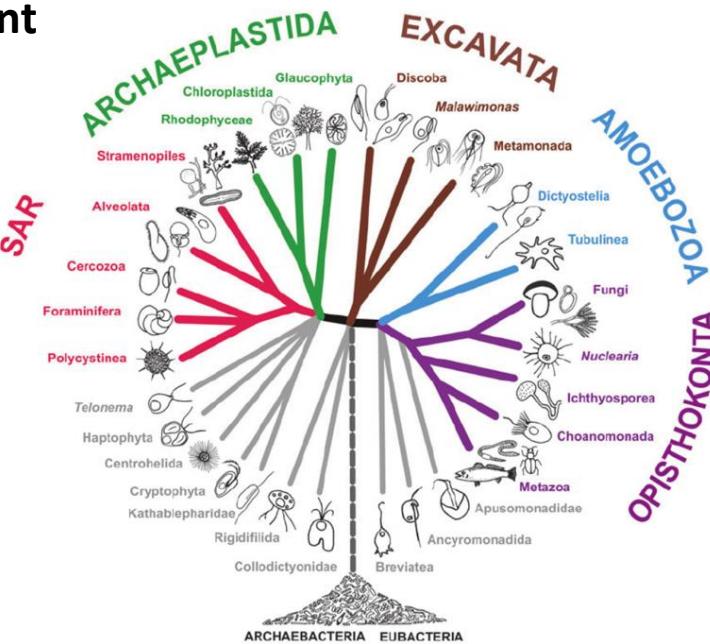
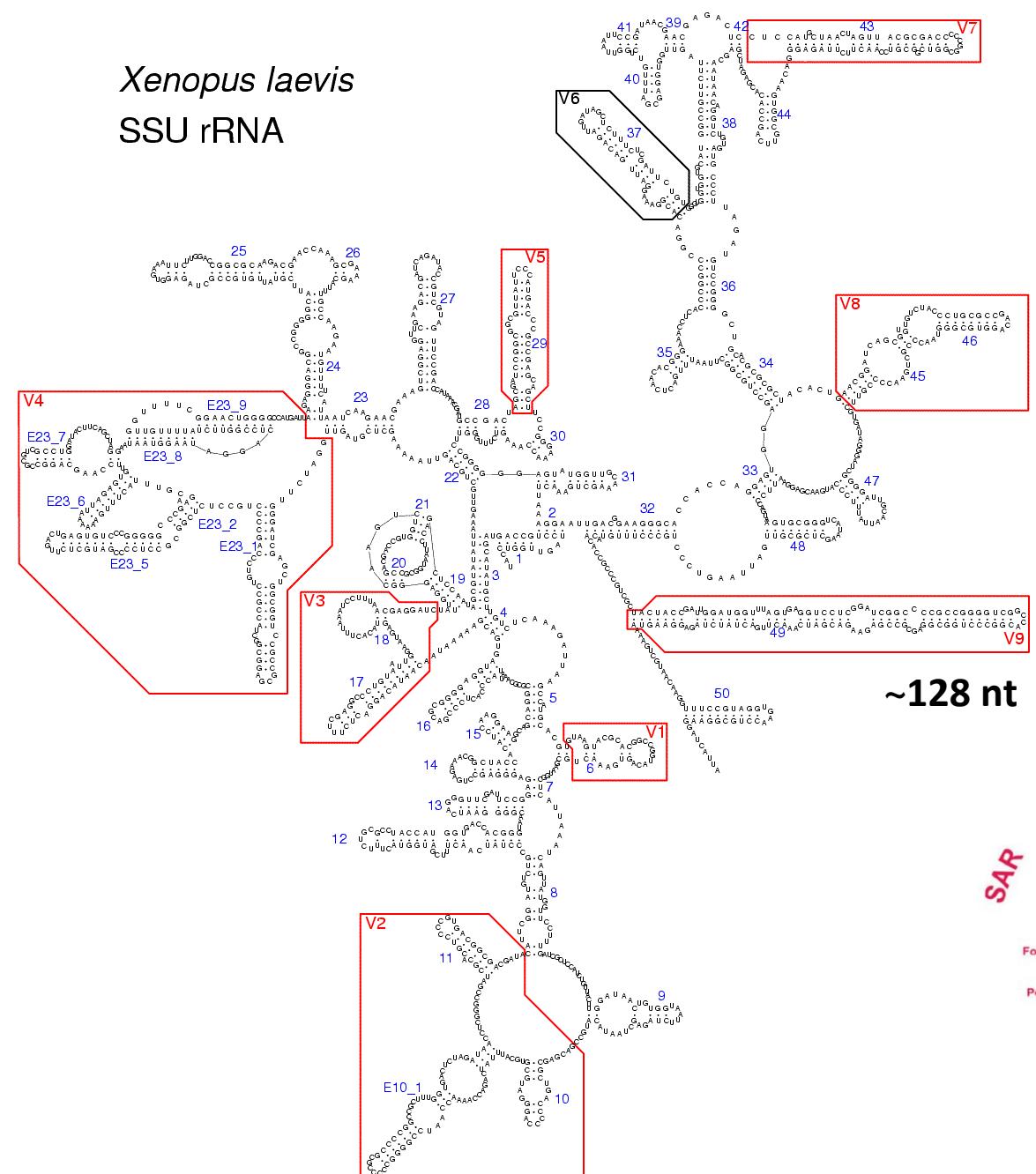


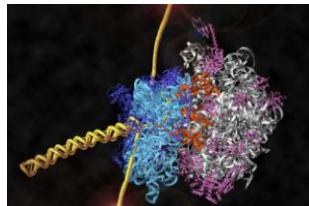
# Tara Oceans Expedition, 2009-2012

381 samples from 49 locations

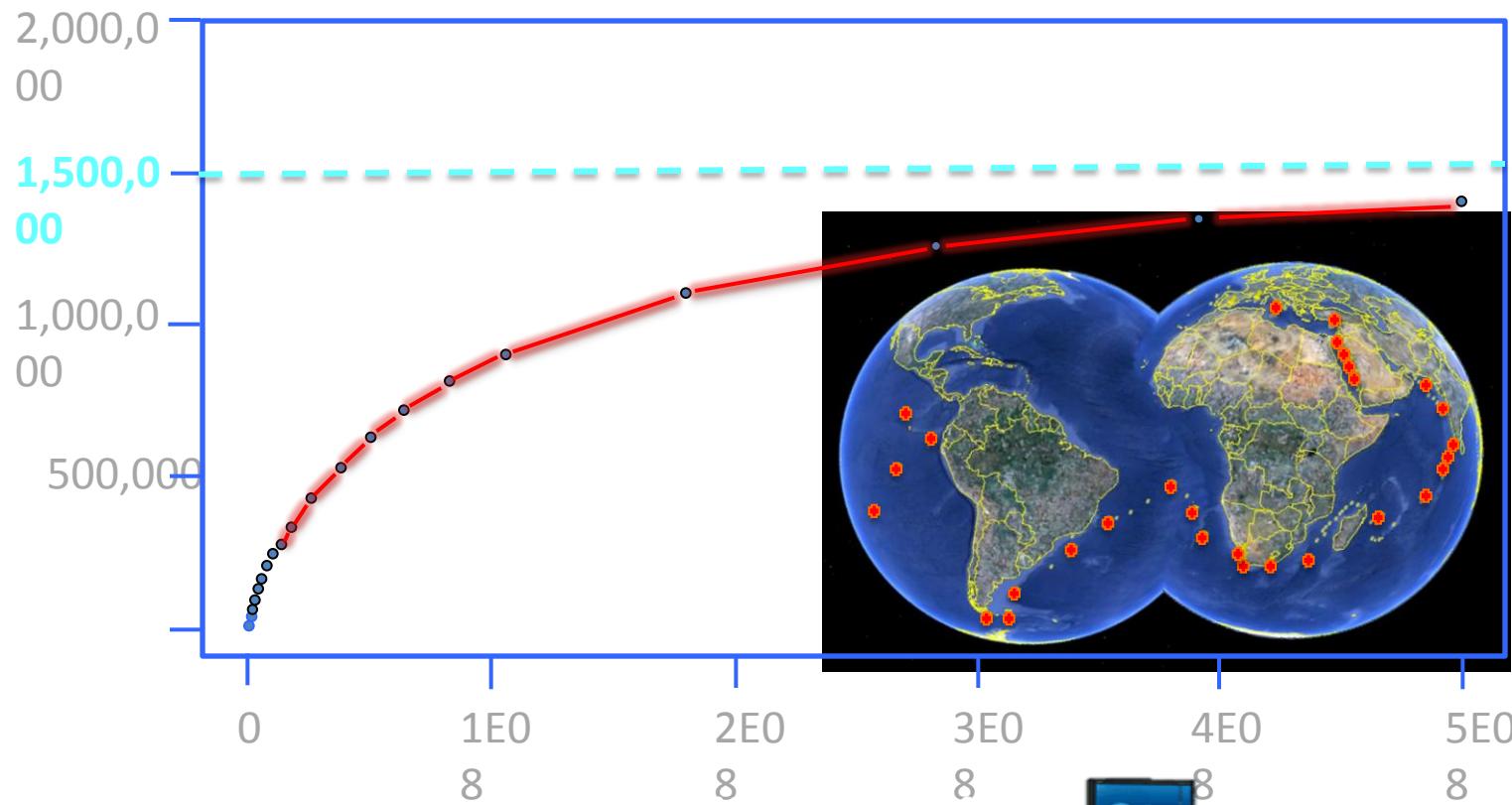


*Xenopus laevis*  
SSU rRNA

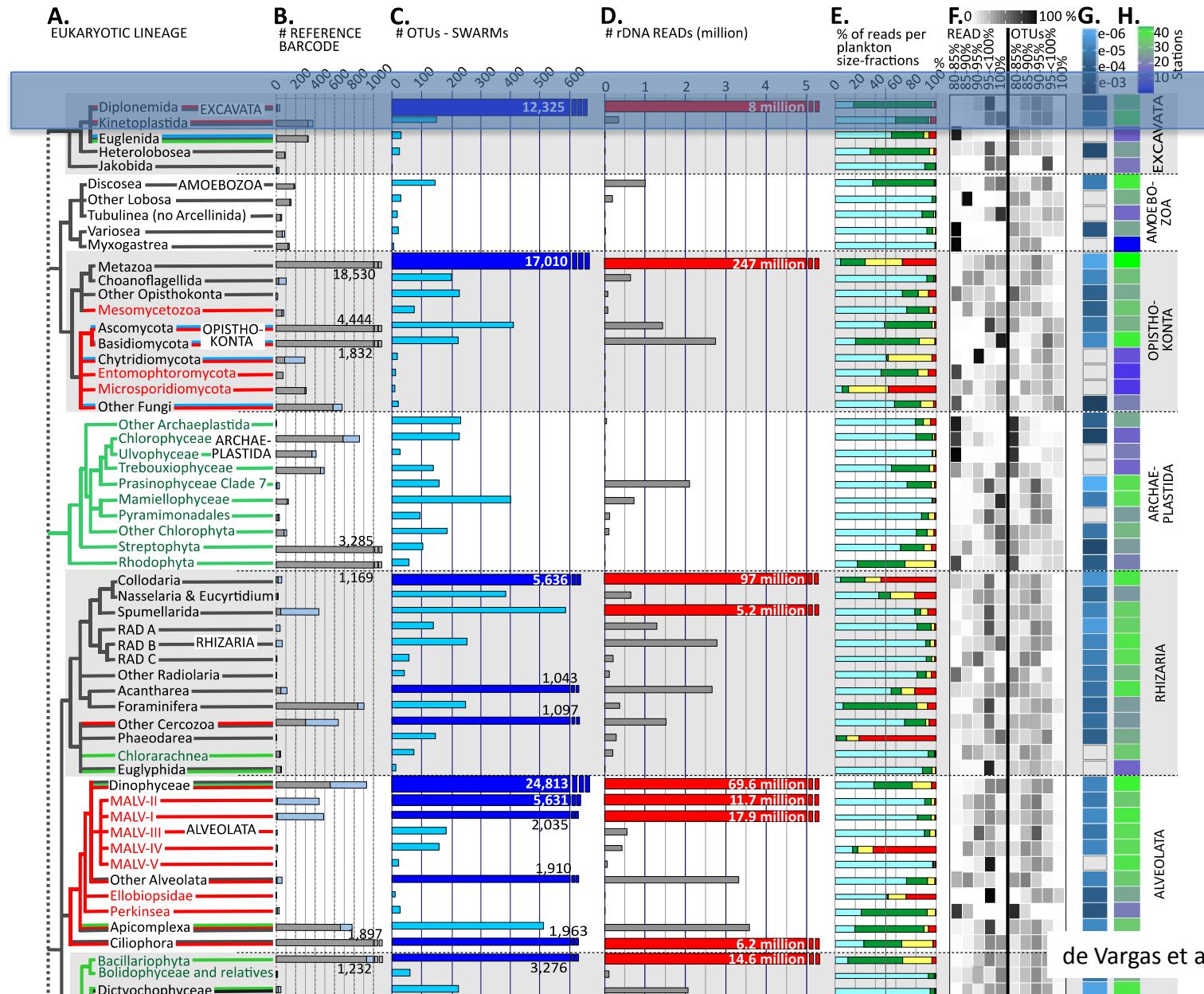




~1.5 million protist + animal taxa in  
photic oceanic plankton



# THE GLOBAL ANALYSIS OF THE V9 DIVERSITY OF THE PHOTIC LAYER



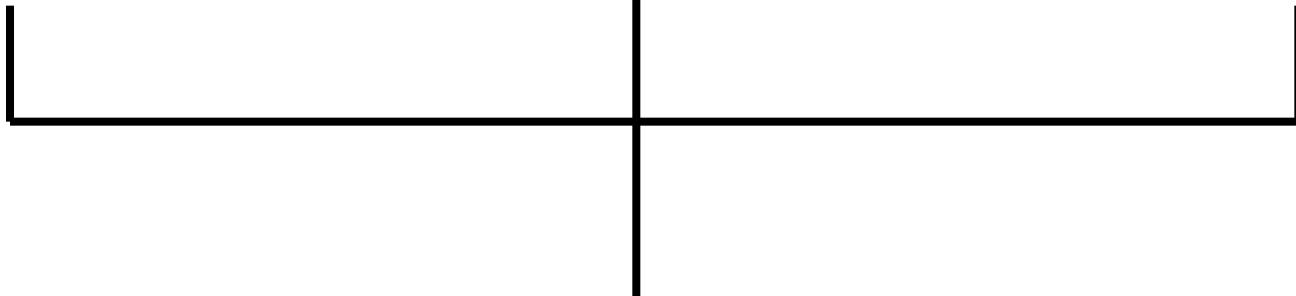
**DIPLODEMIDS?!**

The **Euglenozoa** clade is a primitive eukaryotic group of flagellates.

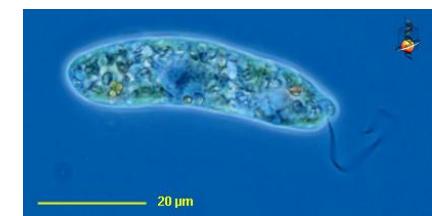


*Trypanoplasma borreli*

Kinetoplastids

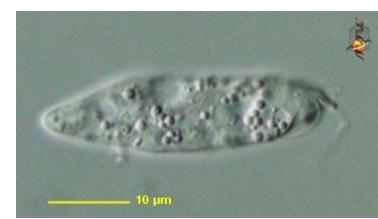


Diplonemids



*Euglena acus*

Euglenids



*Diplonema papillatum*

# Summary of our knowledge of diplonemid morphology

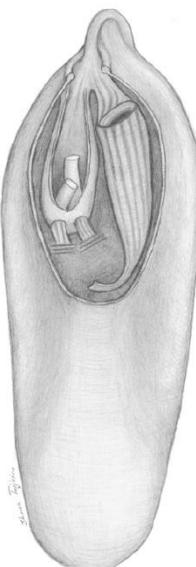
**A**



**B**



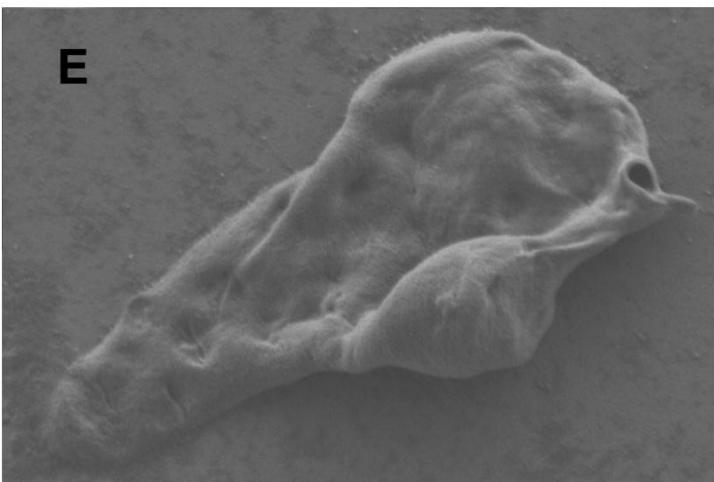
**C**



**D**



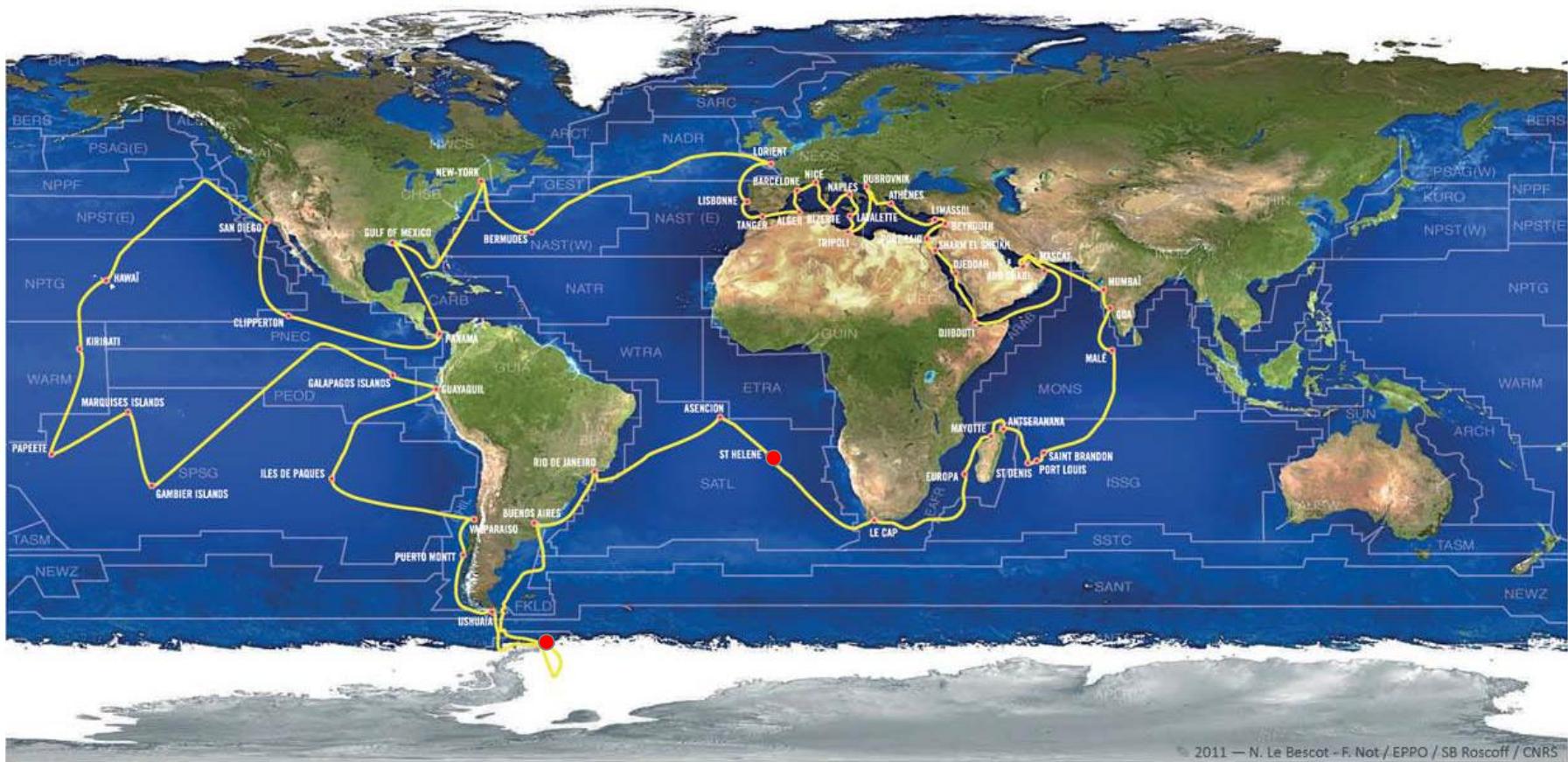
**E**



**A** *Diplonema breviciliata*  
(Simpson 1997);

**B** *Rhynchopus* sp.  
(Simpson 1997);

**C-E** *Rhynchopus euleeides*  
(Roy et al. 2007)



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total abundance: 1,315,922 barcodes

clade: novel diplonemid

sample: 71% of total barcodes in the sample

size fraction: 3 µm - infinity

depth: 784 m

temperature: 0.5 C

total abundance: 845,473 barcodes

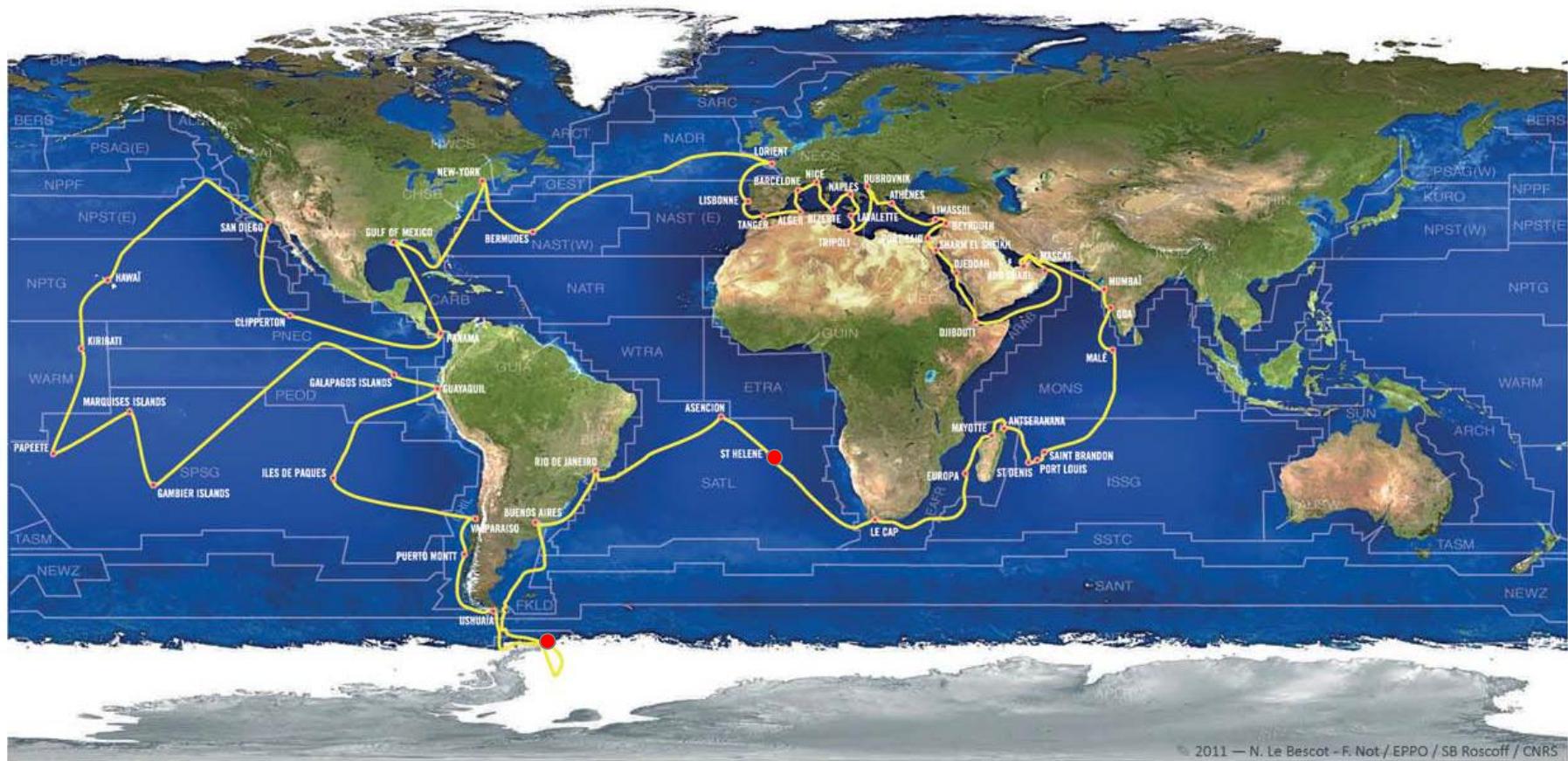
clade: planktonic diplonemid

sample: 51% of total barcodes in the sample

size fraction: 0.8-3 µm

depth: 791 m

temperature: 4.2 C



© 2011 — N. Le Bescot - F. Not / EPPO / SB Roscoff / CNRS

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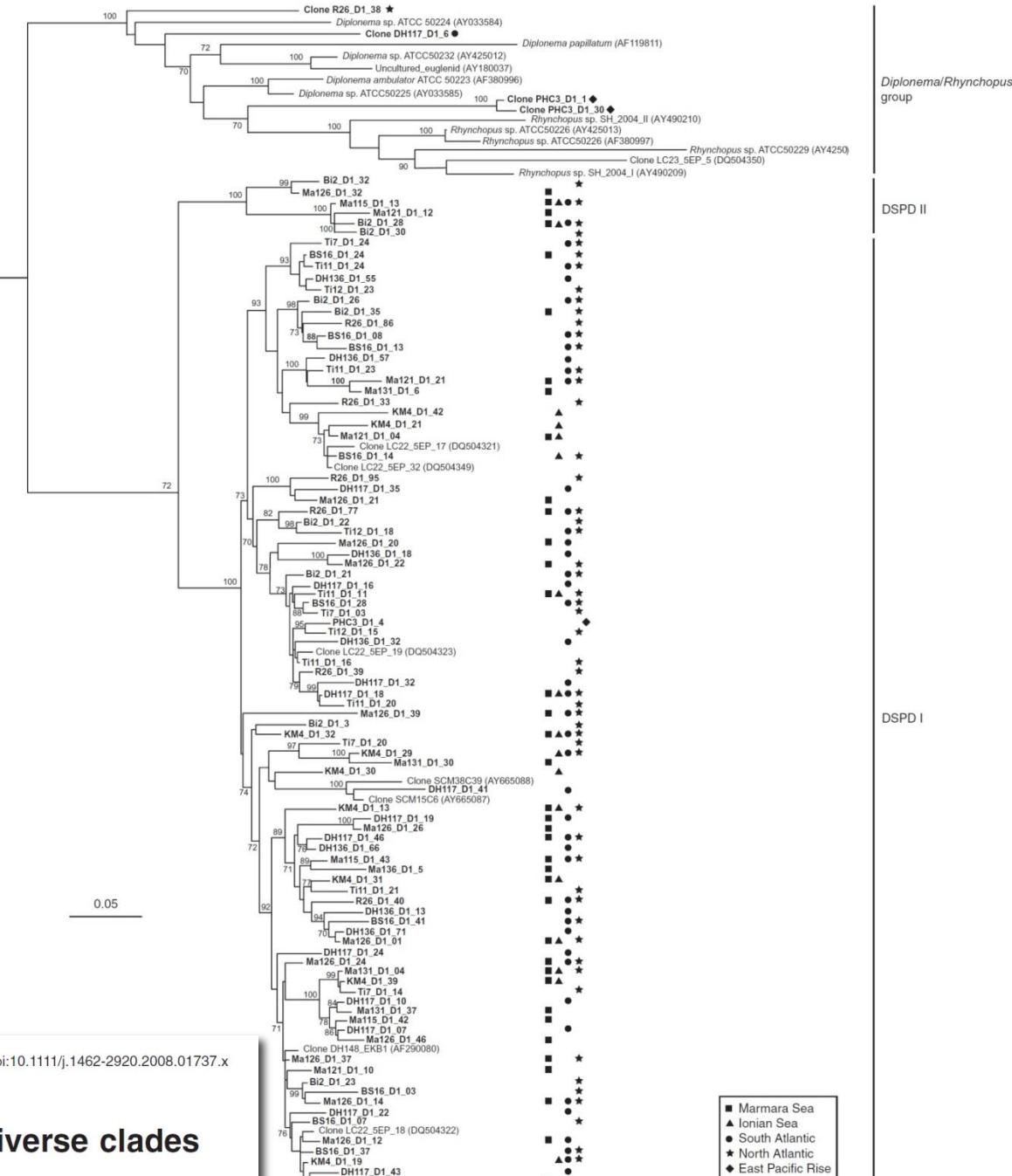
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Enrique Lara,<sup>1</sup> David Moreira,<sup>1</sup>  
Alexander Vereshchaka<sup>2</sup> and  
Purificación López-García<sup>1\*</sup>

doi:10.1111/j.1462-2920.2008.01737.x

Environmental Microbiology (2009) 11(1): 47–55

## Pan-oceanic distribution of new highly diverse clades of deep-sea diplonemids

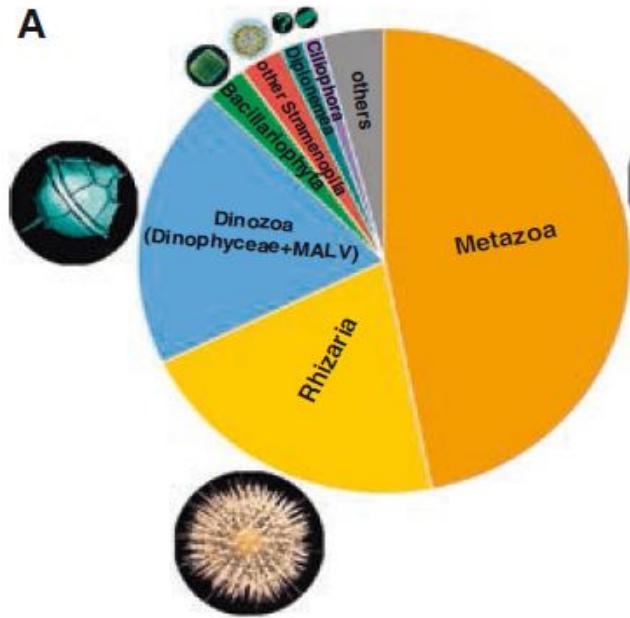
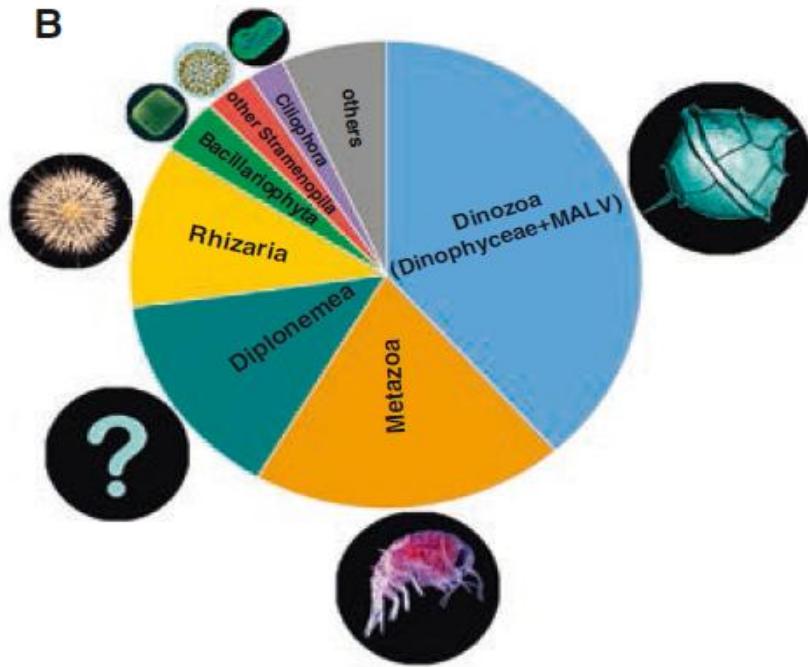
year	authors	location	depth, m
2001	Lopez-Garcia et al.	Drake passage	3000
2007	López-García et al.	North Atlantic Ocean	750
2007	Countway et al.	North Atlantic Ocean	125, 2500
2009	Lara et al.	Marmara Sea	15 - 1250
2009	Lara et al.	Ionian Sea	3000
2009	Lara et al.	South Atlantic Ocean	5 - 1000
2009	Lara et al.	North Atlantic Ocean	2280 - 3500
2009	Lara et al.	East Pacific Rise	1695
2010	Sauvadet et al.	Pacific Ocean	500, 900
2010	Scheckenbach et al.	South Atlantic Ocean	5000
2011	Eloe et al.	Puerto Rico Trench	6000

**156 diplomonmid SSU sequences in GenBank:**

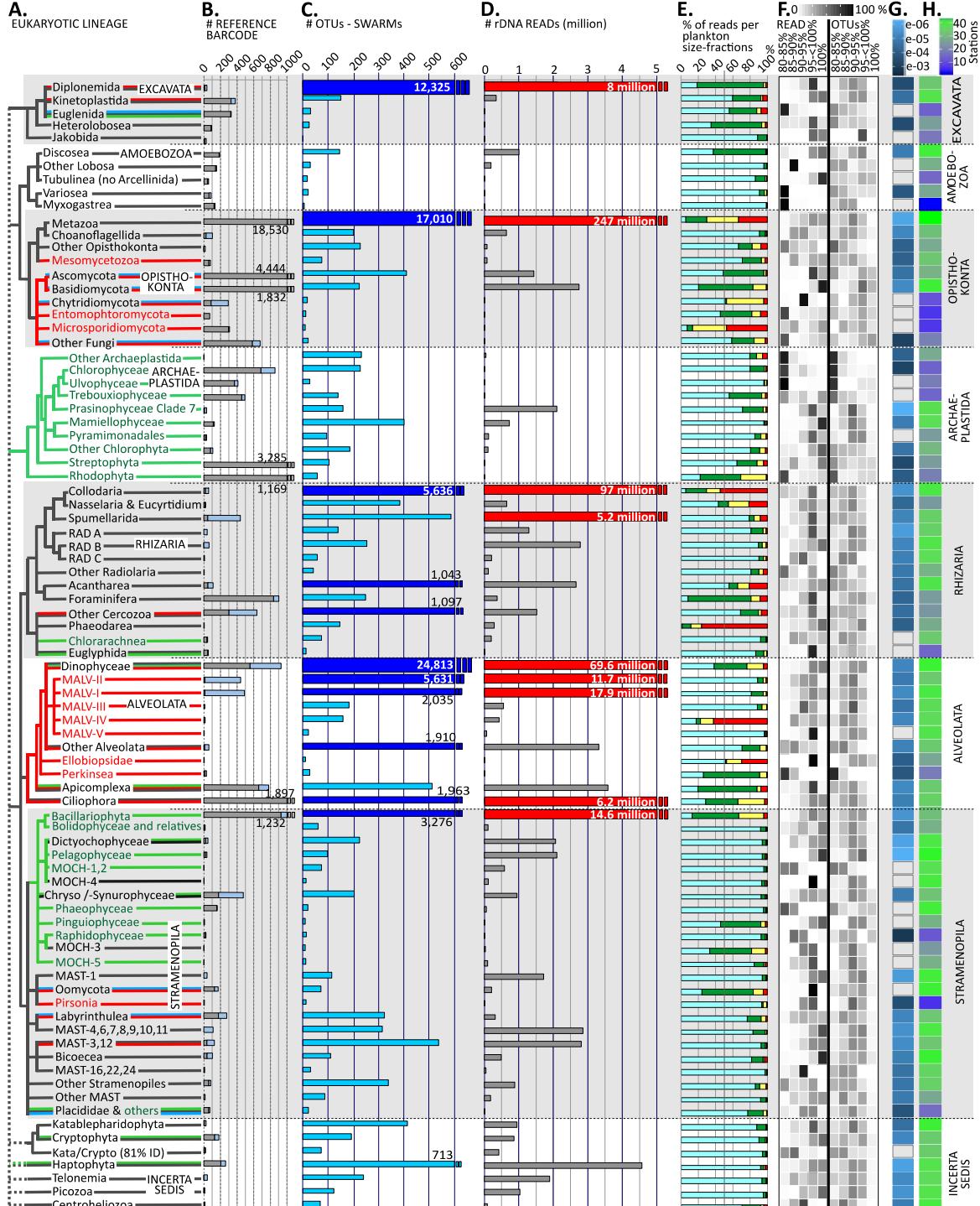
8 *Diplonema* spp.

6 *Rhynchopus* spp.

142 environmental sequences

**A****B**

But diplonemids are the 3<sup>rd</sup> most diverse and 6<sup>th</sup> most abundant eukaryote in the world oceans!



... a few surprises there, e.g. low counts of streptophytes and haptophytes, only diatoms and dinos significant among primary producers.

Obvious winners – metazoans and dino. Amazingly diverse & abundant Rhizarians but first and foremost, **Diplonemids**.

All this in photic zone, but they are deep ,too

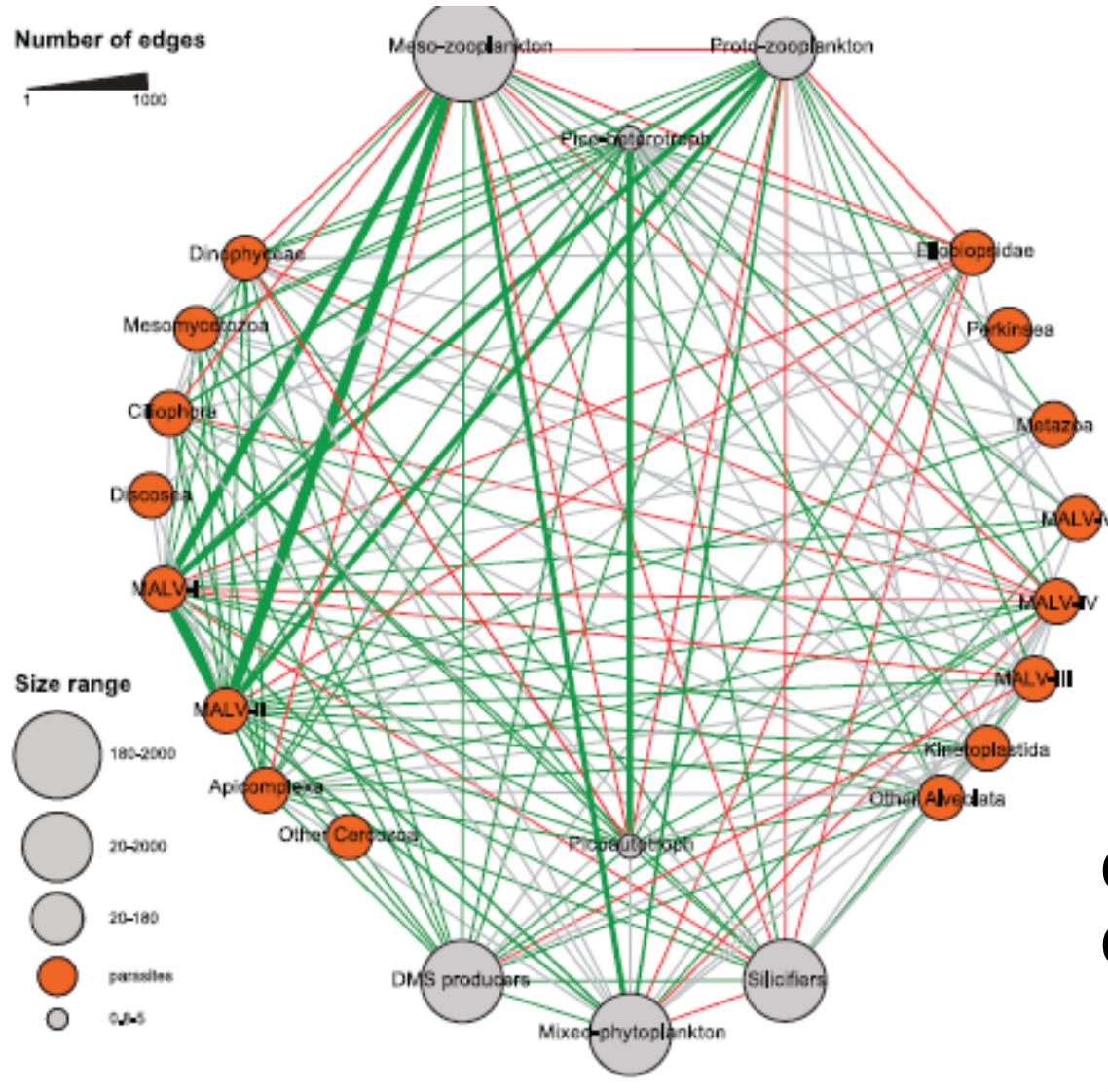
de Vargas et al.  
Science, 2015

# WHAT ARE THEY DOING THERE?

**heterotrophs**

~~phototrophs?~~

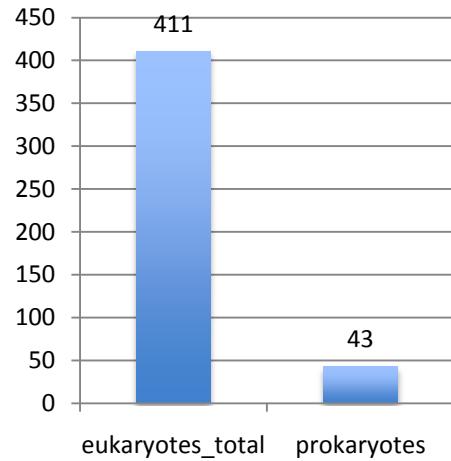
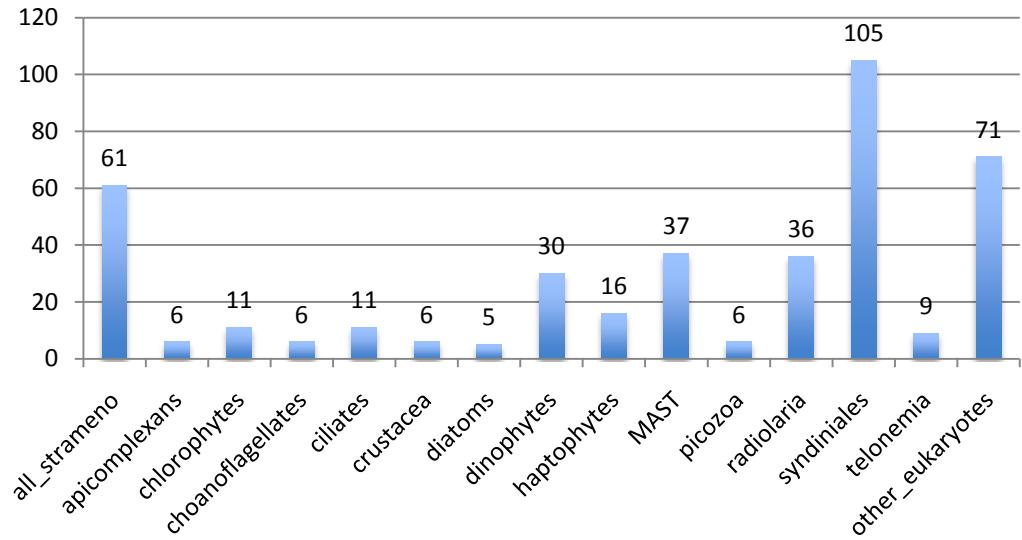
parasites  
symbionts  
predators  
detritovores  
bacteriovores  
...



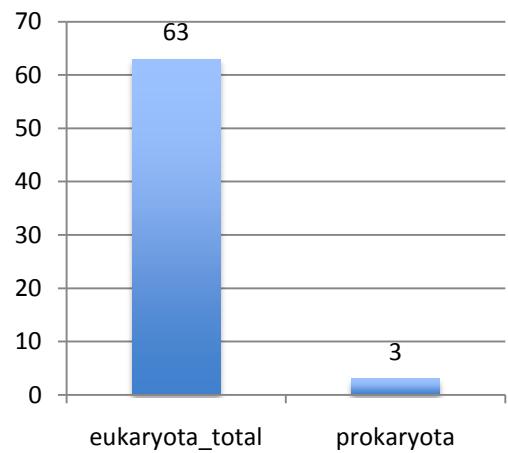
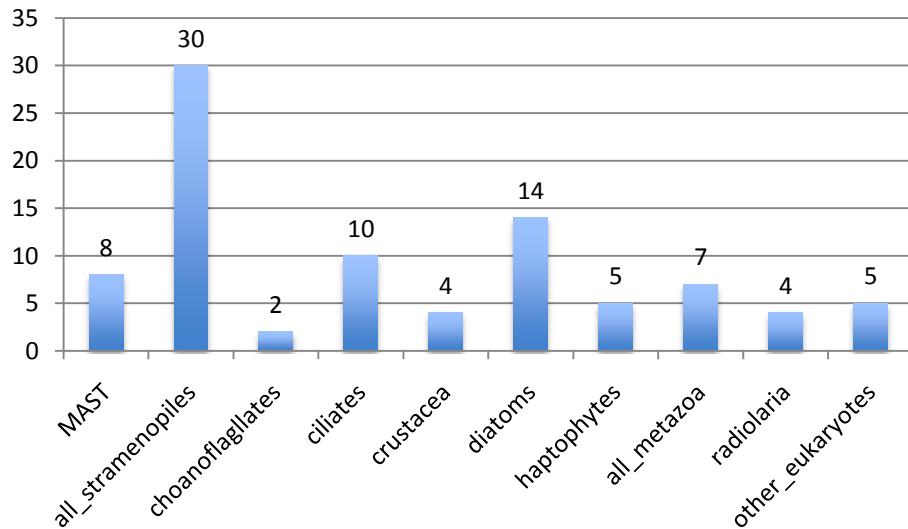
Co-occurrence plot says very little if anything in case of diplonemids

# In-silico estimates of interactions, general numbers

## copresence



## mutual exclusion



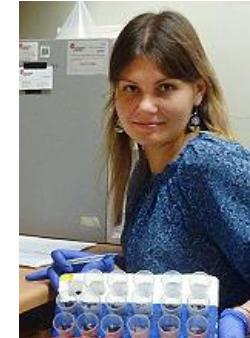
# Conclusions

- A clade sister to classic diplomonemids is the most abundant group of Excavata in the World Ocean plankton.
- As compared to other eukaryotes, it emerges as a hyper-diverse clade, the third by the number of OTUs after dinoflagellates and metazoans.
- Marine planktonic diplomonemids reach very high relative abundance in some stations of the deep non-photic zone, in small size-fractions up to 20 µm.
- This huge group represents an overlooked major player in marine ecosystems, however, nothing is known about their morphology and lifestyle.

# Acknowledgements



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