

The INRAe logo is located in the bottom left corner. It consists of three overlapping circles in different shades of green (light green, medium green, and dark teal). To the right of the circles, the acronym "INRAe" is written in a bold, teal, sans-serif font.

INRAe



Diatom metabarcoding for biomonitoring : 1st part

F. Rimet

Reminders:
Biology
Taxonomy
Biomonitoring



Reminders

- 1- Diatom biology
- 2- Taxonomy
- 3- Classical diatom biomonitoring



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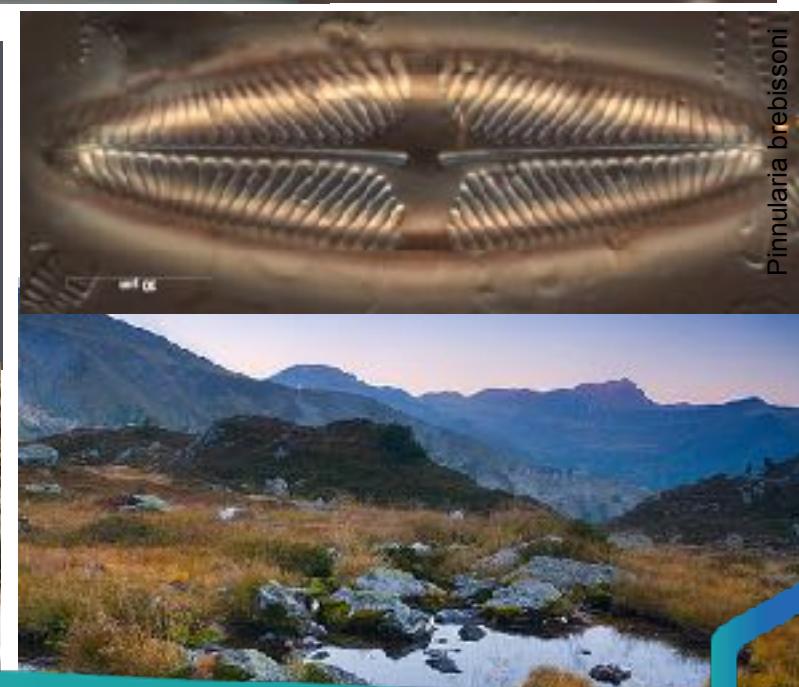
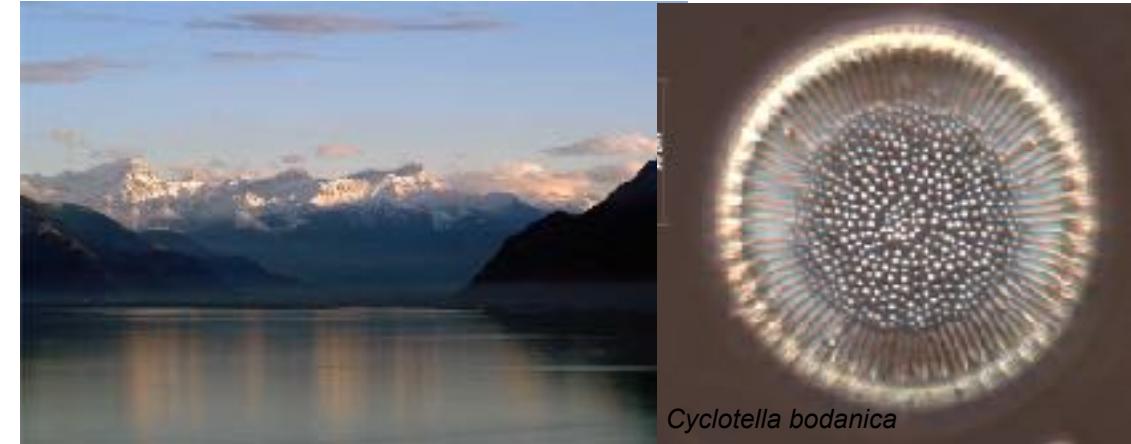
www.biolaweb.com

Diversity and importance in the biosphere

- 100 000 species (Mann & Vanormelingen 2013)
- 25% of the total biomass on earth (Werner 1977)
- Colonize all habitats:
lakes, rivers, oceans, soils,
wet walls, caves ...



Luticola ventricosa

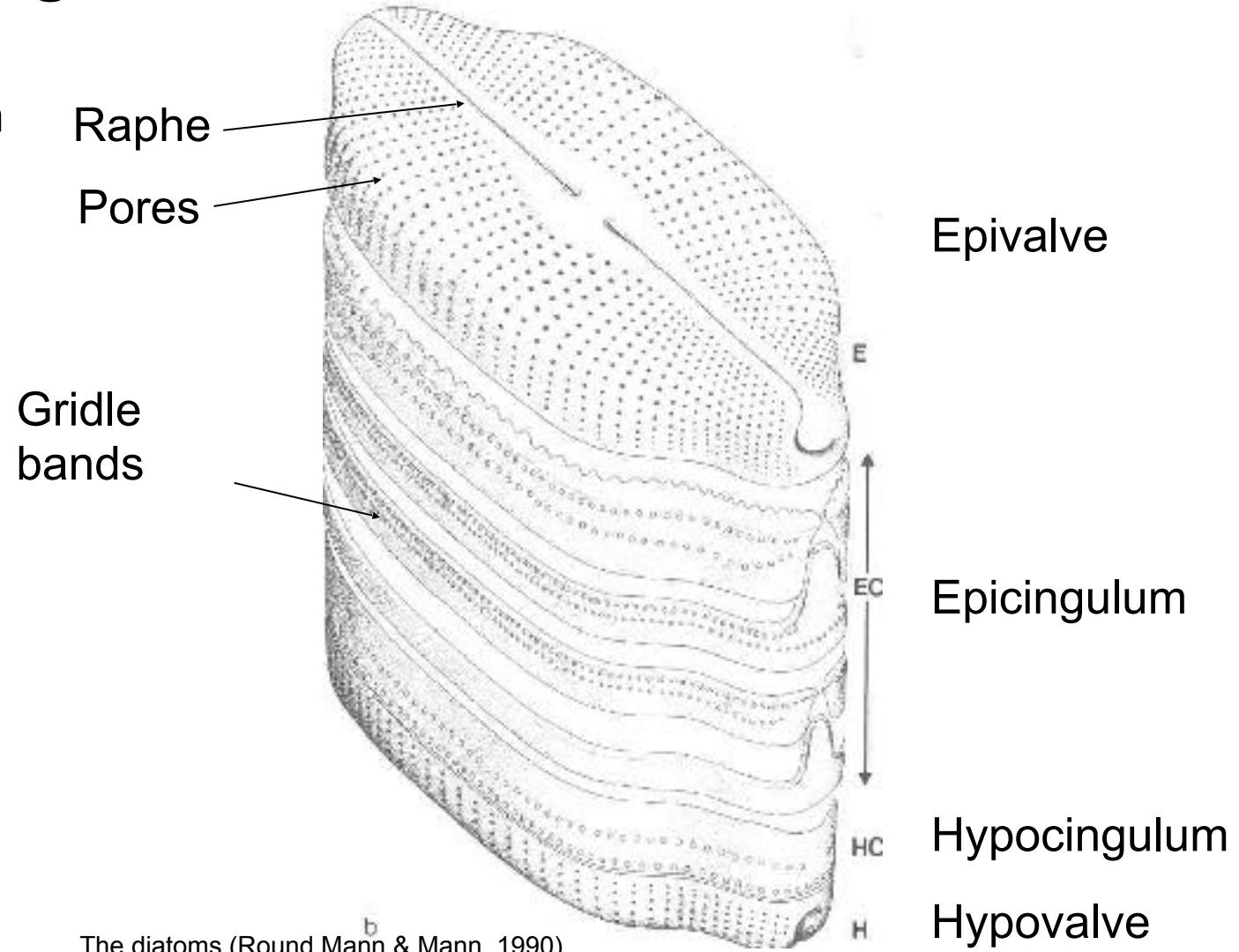
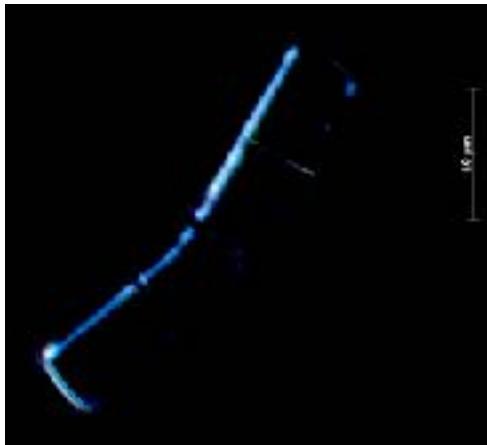


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Cell organisation

Extracellular siliceous skeleton



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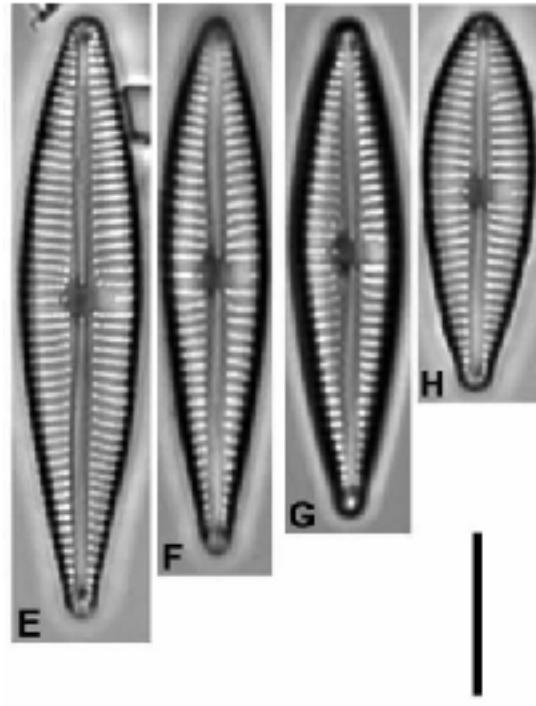
www.biolaweb.com

Life cycle

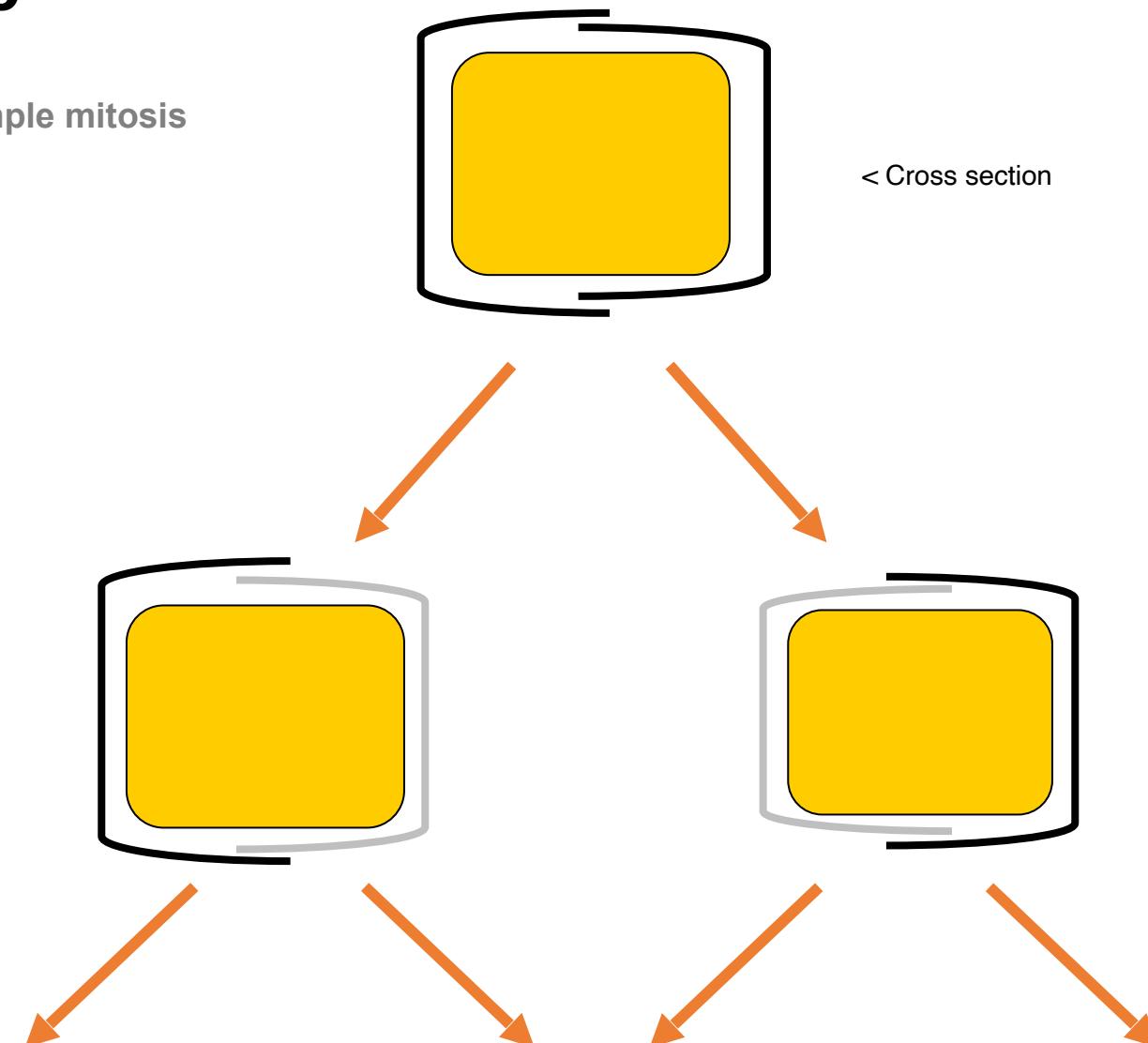
- Reproduction :

Vegetative reproduction: simple mitosis

Cell reduction



Gomphonema parvulum
(culture strain) Rose & Cox 2014



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Life cycle

- Reproduction:

Sexual reproduction (Round, Crawford, Mann 1990)

Sexual
reproduction
(Gametes mating)

Meiosis

$2n$ chromosomes > n chromosome
Production of gametes

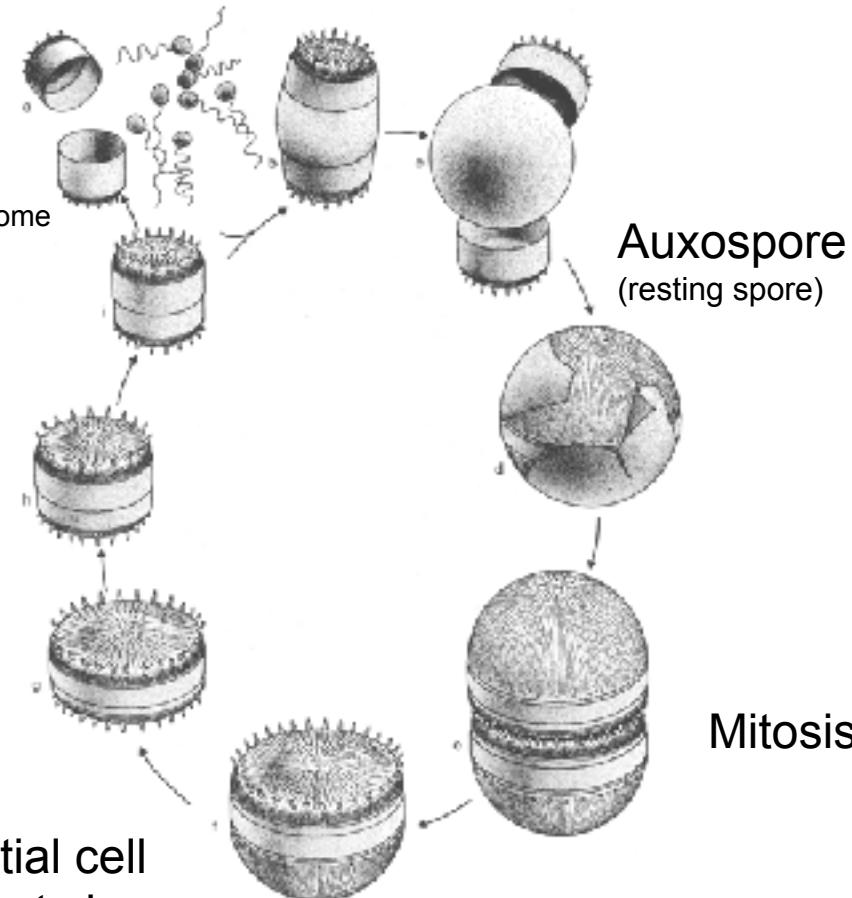
Mitosis

Vegetative cell

Initial cell
largest size

Auxospore
(resting spore)

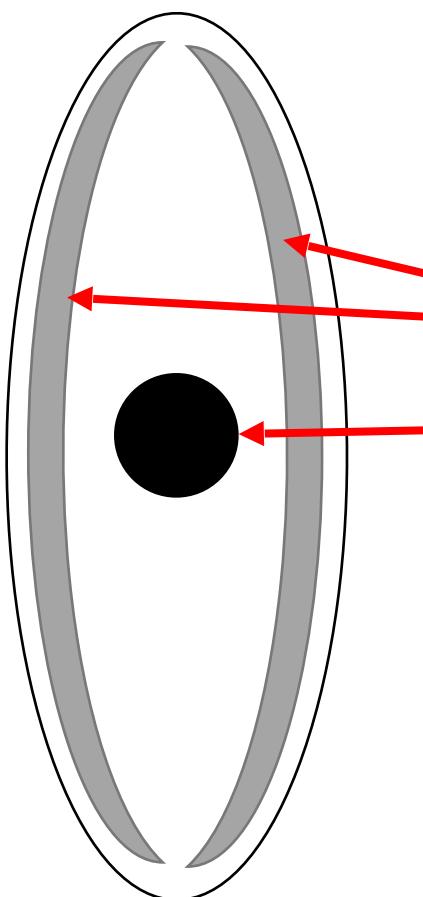
Mitosis



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Cell organisation

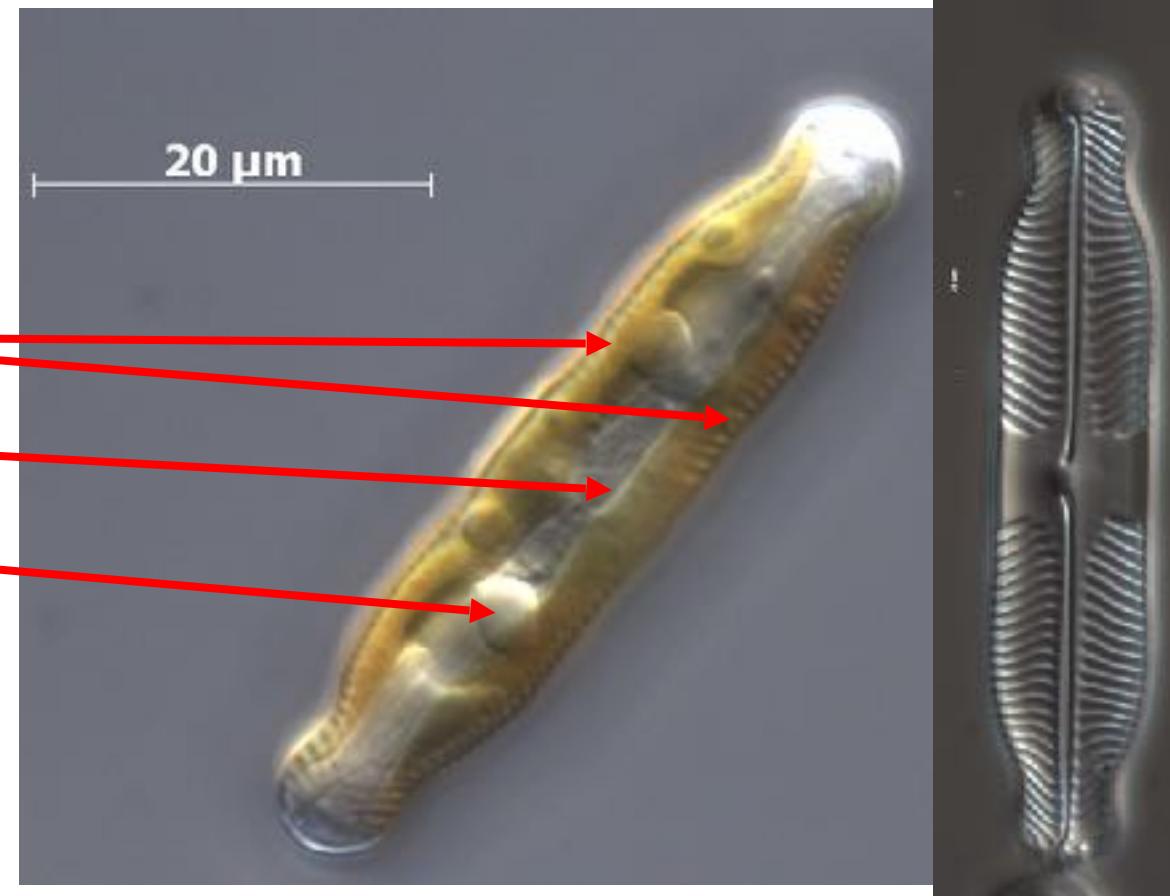


Intracellular organelles

Chloroplasts

Nucleus

Lipidic drops



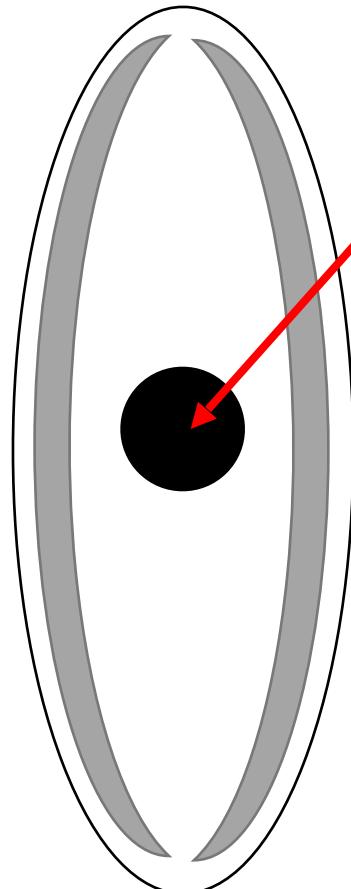
TCC879-*Pinnularia lundii* var. *linearis*



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Cell organisation



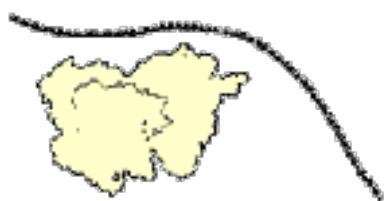
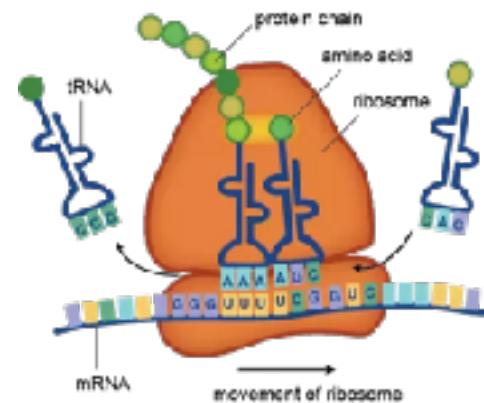
Nucleus :

Several chromosomes (nucleus) have coding genes for ribosomal RNA.
rRNA > ribosome, composed by:

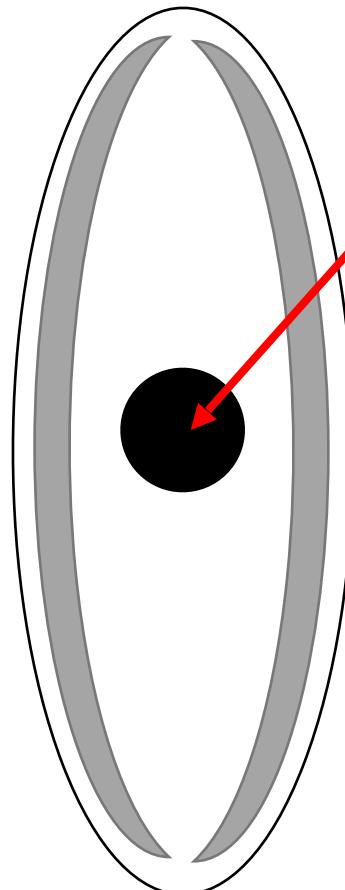
- Large sub-unit (60S) : 28S rRNA, 5,8S rRNA, 5S rRNA
 - Small sub-unit (40S) : 18S rRNA

18S marker is widely used in biodiversity studies because:

- Variable sequences flanked by highly conserved regions allowing the use of universal primers.
 - Numerous copies in the genome: present in large quantities



Cell organisation



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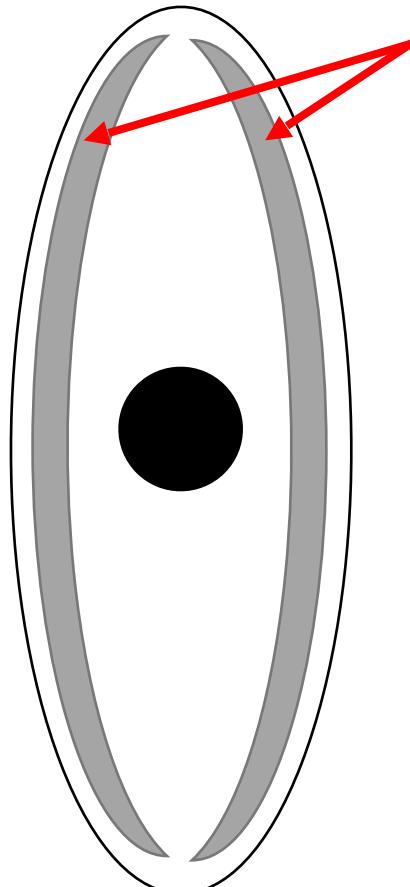
Protist Ribosomal Reference database (PR²)



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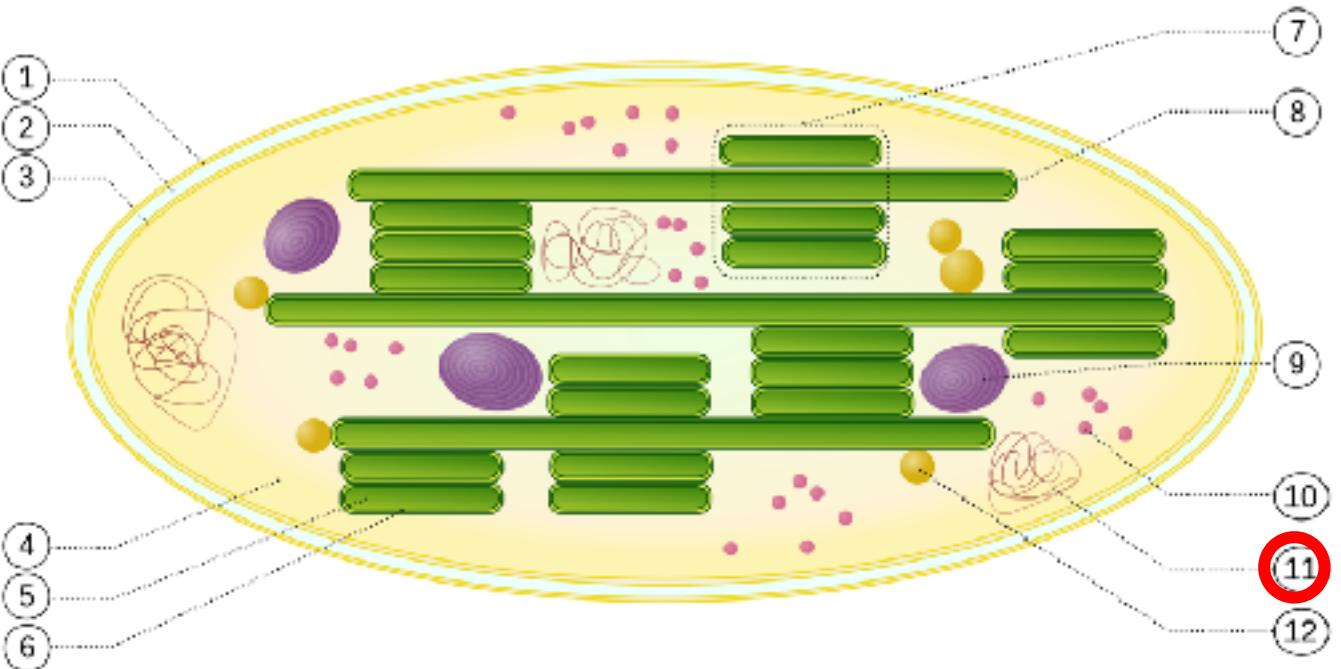
www.biolaweb.com

Cell organisation



Chloroplasts:

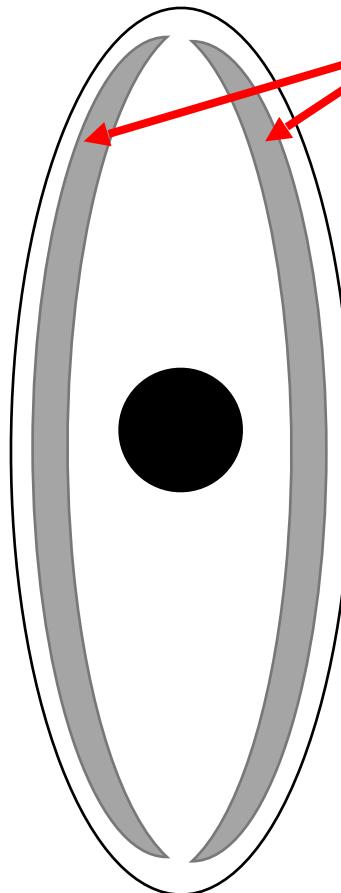
- (1) outer membrane; (2) intermembrane space; (3) Inner membrane
- (4) stroma ; (5) lumen du thylakoïde ; (6) membrane du thylakoïde ;
- (7) granum (thylakoïdes stack(8))
- (9) amidon ; (10) ribosome;
- (11) Chloroplastic DNA ;
- (12) Lipidic droplet



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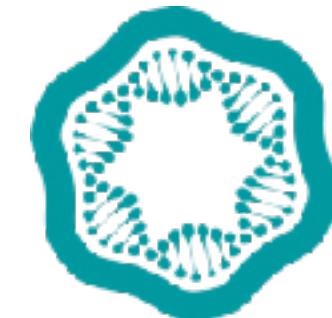
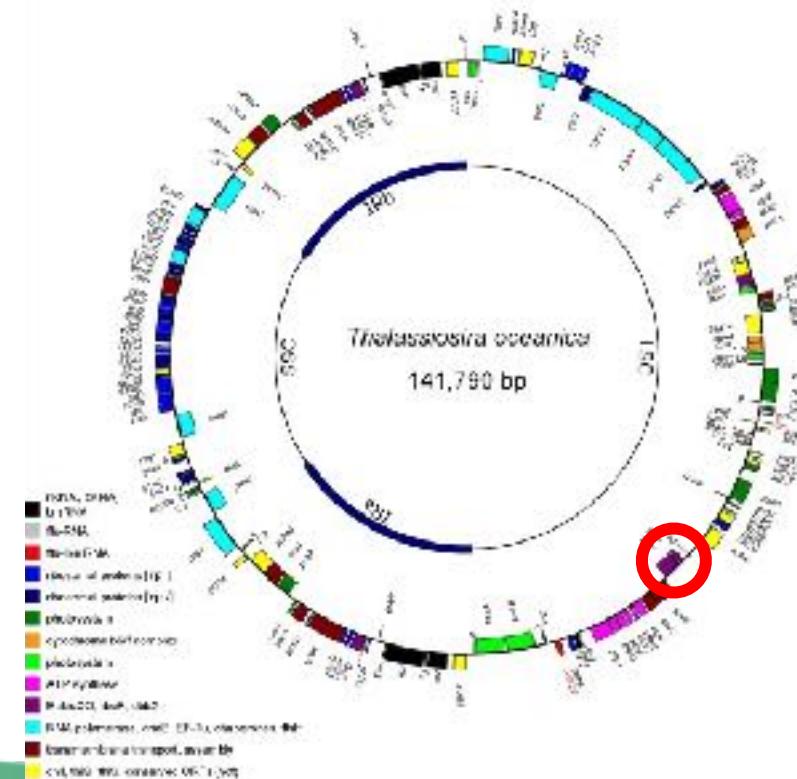
www.biolaweb.com

Cell organisation



Chloroplasts:

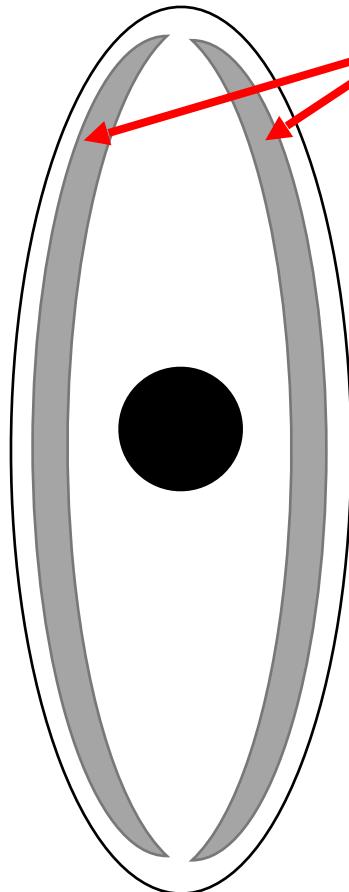
In the chloroplast genome, an important marker: rbcL
Coding the large sub-unit (« L ») of RuBisCo (« rbc »), key enzyme of photosynthesis.



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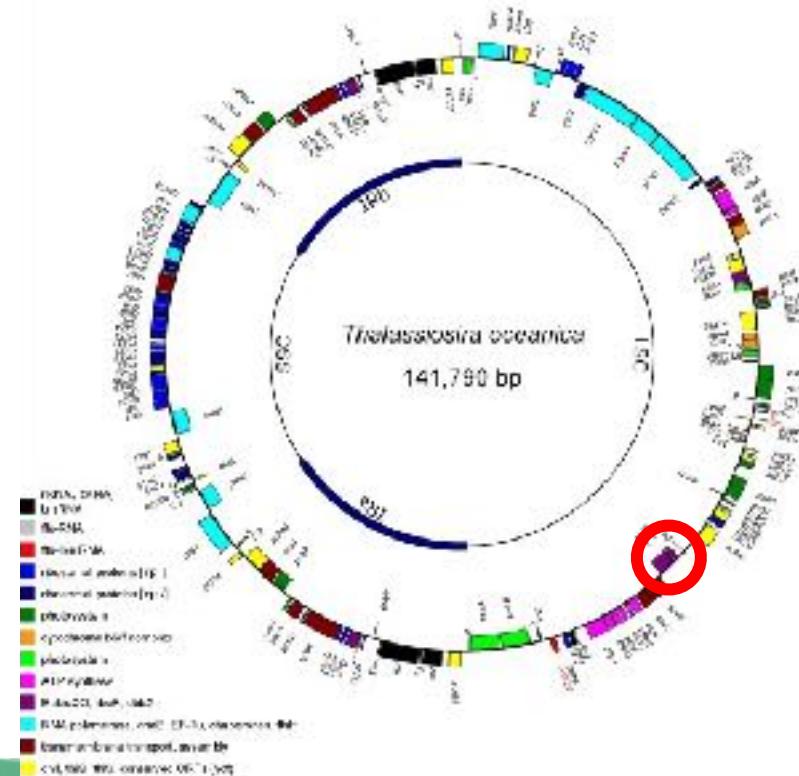
www.biolaweb.com

Cell organisation

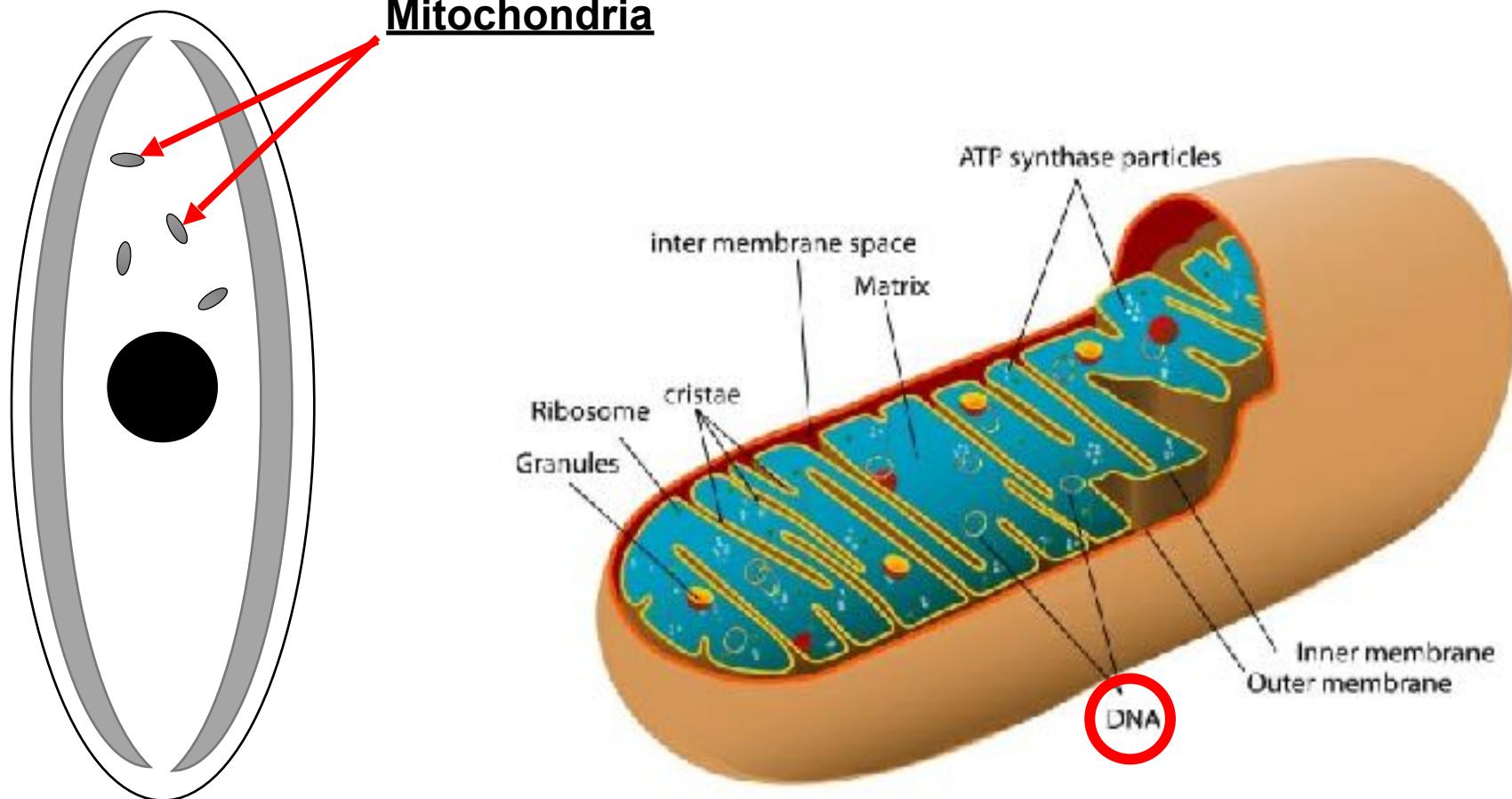


Chloroplasts:

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Coding the large sub-unit (« L ») of RuBisCo (« rbc »), key enzyme of photosynthesis.



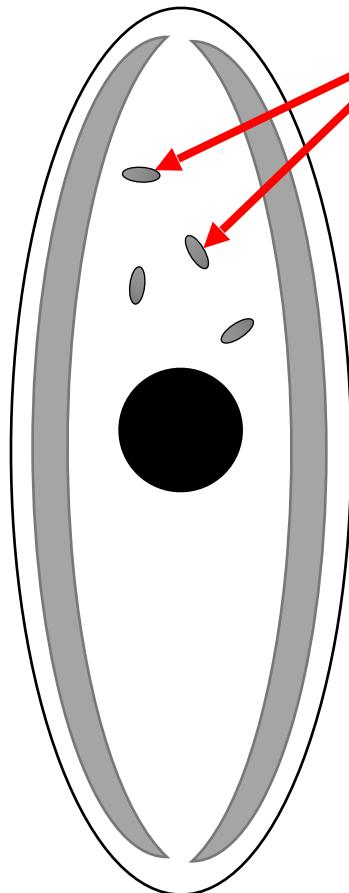
Cell organisation



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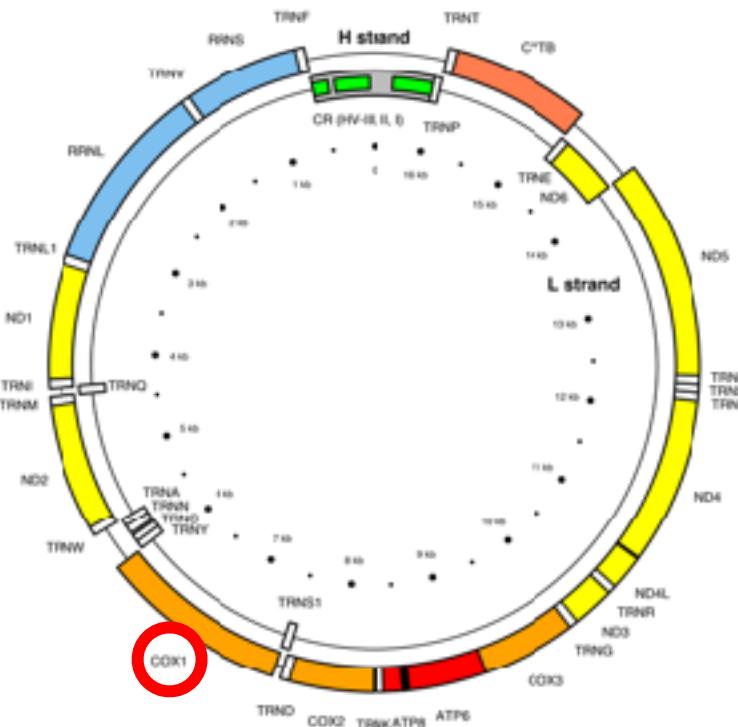
www.biolaweb.com

Cell organisation



Mitochondria:

In the mitochondrial genome, an important marker : Cox1 encodes a subunit of Cytochrome c oxidase, which is involved in the respiration process. It is a key enzyme in aerobic respiration.



BOLD SYSTEMS

DATABASES IDENTIFICATION TAXONOMY WORKBENCH RESOURCES LOGIN □

BARCODE OF LIFE DATA SYSTEM ^{v4}

Advancing biodiversity science through DNA-based species identification.

EXPLORE DATA



DESIGNED TO SUPPORT THE GENERATION & APPLICATION OF DNA BARCODE DATA

ECLD is a cloud-based data storage and analysis platform developed at the Centre for Biodiversity Genomics in Canada. It consists of four main modules: a data portal, an educational portal, a registry of BINs (putative species), and a data collection and analysis workbench.



DATA PORTAL

A data retrieval interface that allows for searching over 9.7M public records in ECLD using multiple search criteria including, but not limited to, geography, taxonomy and depository.



EDUCATION PORTAL

A custom platform for educators and students to explore barcode data and contribute novel barcodes to the ECLD database.



BIN DATABASE

A searchable database of Barcode Index Numbers (BINs), sequence clusters that closely approximate species.



WORKBENCH

A data collection and analysis environment that supports the assembly and validation of DNA barcodes and other sequences.



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Public Data Portal - RecordList | +

boldsystems.org/index.php/Public_BINSearch/searchrecords

BOLDSYSTEMS DATABASES IDENTIFICATION TAXONOMY WORKBENCH RESOURCES LOGIN 🔍

PUBLIC DATA PORTAL

PUBLIC DATA 🔍 SEARCH

The Public Data Portal supports queries based on taxonomic/scientific names only), geography, attribution fields (i.e. collectors and taxonomists), specimen depositories, project numbers/identifiers, specimen and sequence identifiers (imageids, museumids, processed, etc). Free-text searches are allowed; the system will try to extract meaningful terms from the search. Search terms should be separated by a space; search terms containing more than one word ("United States" or "Mus musculus") should be wrapped in quotes.

Search examples (Searches are case insensitive):

- **Diptera**: will return all the mosquitoes in the database
- **Mammalia Aves**: will return all the mammals and birds in the database. Using two terms under the same context (taxonomic in this case), returns all records that match either term.
- **Aves Peru**: will return all the birds collected in Peru. Using two terms under a different context (taxonomic and geographic), returns records that only match both terms.
- **Hawaii <Chordata**: will return all non-chordates, or invertebrates, collected in Hawaii. It is often necessary to exclude records based on a search criteria and is accomplished by adding a minus sign immediately in front of the search term.
- ***Biodiversity Institute of Ontario* Lepidoptera Florida**: will return all moth and butterflies in the Biodiversity Institute of Ontario collection that were collected in Florida. Complex queries are possible by combining multiple search terms.

See also Guidelines:

- No spaces in terms when using a multiterm search (correct: "United States" Mammalia)
- Negative symbol (-) is an exclusion operator which must be preceded by a space
- If the same term appears in multiple contexts (i.e. "Florida" appears in both the geography field and in "University of Florida"), when using a free text search for the state Florida, employ the context clarification "[geo]" (e.g. Florida[geo]). Other context clarification tags are [tax], [ids], [inst], [researcher].

Help 🔍

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25/02/2021

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Chia21.C13 (C1) Record List | Public Data Portal +

bioinfo.systera.org/index.php/Public_SearchTerm

BOLD SYSTEMS DATABASES IDENTIFICATION TAXONOMY WORKBENCH RESOURCES LOGIN Q

Specimens: DYC XML TSV
Sequences: FASTA TRACE
Combined: XML CSV
MB3C Generate from

Barbelliphyla PUBLIC DATA SEARCH

Records 1 to 100 Page 1 2 3 4 5 6 7 8 9 10 Requests per Page 100 ▾

ACHL0049-19 • *Barbelliphyla* (C040P.B88)
Taxonomy: Radiotrophophyta, Barbelliphylidae, Barbelliphyla
Identifiers: ACHL0049 [sampled], BSLF44 [id];
Zoogeography: Université Paris-Diderot, Chiroptologie des milieux continentaux
Collected in France

ACHP0001-16 • *Advanthidium* (186129.215.851.ncbi.nlm.nih.gov)
Taxonomy: Barbelliphophyta, Barbelliphylidae, Advanthidiidae
Identifiers: SPITS13_647mexB1sampled, PvO105 [museum];
Zoogeography: Ghent University Protistology and Acoustic Ecology
Collected in Norway, Svalbard, Svalbard, Svalbard

ACHP0003-19 • *Advanthidium* (186129.215.851.ncbi.nlm.nih.gov)
Taxonomy: Barbelliphophyta, Barbelliphylidae, Advanthidiidae, Advanthidiaceae, Advanthidiidae
Identifiers: SPITS13_M0_10 [sampled], PvO105 [museum];
Zoogeography: Ghent University Protistology and Acoustic Ecology
Collected in Norway, Svalbard, Svalbard

ACHP0003-16 • *Advanthidium* (186129.215.851.ncbi.nlm.nih.gov)
Taxonomy: Radiotrophophyta, Radiotrophophyletes, Advanthidiidae, Advanthidiaceae, Advanthidiidae
Identifiers: SPITS13_M04algal_1 [sampled], PvO105 [museum];
Zoogeography: Ghent University Protistology and Acoustic Ecology
Collected in Norway, Svalbard, Svalbard

ACHP0004-15 • *Advanthidium* (186129.215.851.ncbi.nlm.nih.gov)
Taxonomy: Radiotrophophyta, Radiotrophophyletes, Advanthidiidae, Advanthidiaceae, Advanthidiidae
Identifiers: SPITS13_M02plus8_10 [sampled], PvO105 [museum];
Zoogeography: Ghent University Protistology and Acoustic Ecology
Collected in Norway, Svalbard, Svalbard

ACHP0005-15 • *Advanthidium* (186129.215.851.ncbi.nlm.nih.gov)
Taxonomy: Radiotrophophyta, Radiotrophophyletes, Advanthidiidae, Advanthidiaceae, Advanthidiidae
Identifiers: SPITS13_M03_10 [sampled], PvO105 [museum];
Zoogeography: Ghent University Protistology and Acoustic Ecology
Collected in Norway, Svalbard, Svalbard

ACPD0006-15 • *PSAMMOPHYLUM INHABENS* (186129.215.851.ncbi.nlm.nih.gov)
Taxonomy: Radiotrophophyta, Radiotrophophyletes, Advanthidiidae, Advanthidiaceae, Advanthidiidae

Results Summary

Found 1,319 published records, with 4,810 records with sequences forming 235 BINS (103). with specimens from 49 countries, deposited in 22 institutions.

Of these records, 3,649 have species names, and represent 730 species.

Specimen Distribution



Data Summary



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ASTBR002-13 - *Asterionellopsis tenellifera* [185:374, SP:425,ITS:788,rcL:718]

Taxonomy: Bacillariophyta, Bacillariophyceae, Fragilariales, Fragilariacae, Asterionellopsidales
Identifiers: IC_NPAA: 2[unspecified], M82[feldid]
Depository: Research Collection of I. Kaczmarska
Collected in: Canada, New Brunswick, Bay of Fundy

ASTBR003-13 - *Asterionellopsis tenellifera*

Taxonomy: Bacillariophyta, Bacillariophyceae, Fragilariales, Fragilariacae, Asterionellopsidales
Identifiers: IC_NPAA: 3[unspecified], M83[feldid]
Depository: Research Collection of I. Kaczmarska
Collected in: Canada, New Brunswick, Bay of Fundy

ASTBR004-13 - *Asterionellopsis tenellifera* [173:70]

Taxonomy: Bacillariophyta, Bacillariophyceae, Fragilariales, Fragilariacae, Asterionellopsidales
Identifiers: IC_NPAA: 4[unspecified], M87[feldid]
Depository: Research Collection of I. Kaczmarska
Collected in: Canada, New Brunswick, Bay of Fundy

ASTBR005-13 - *Asterionellopsis tenellifera* [185:374, SP:425,ITS:788,rcL:718]

Taxonomy: Bacillariophyta, Bacillariophyceae, Fragilariales, Fragilariacae, Asterionellopsidales
Identifiers: IC_NPAA: 5[unspecified], M88[feldid]
Depository: Research Collection of I. Kaczmarska
Collected in: Canada, New Brunswick, Bay of Fundy

ASTBR006-13 - *Asterionellopsis tenellifera* [185:374, SP:425,ITS:788,rcL:718]

Taxonomy: Bacillariophyta, Bacillariophyceae, Fragilariales, Fragilariacae, Asterionellopsidales
Identifiers: IC_NPAA: 6[unspecified], M89[feldid]
Depository: Research Collection of I. Kaczmarska
Collected in: Canada, New Brunswick, Bay of Fundy

ASTBR007-13 - *Asterionellopsis tenellifera* [185:374, SP:425,ITS:788,rcL:718]

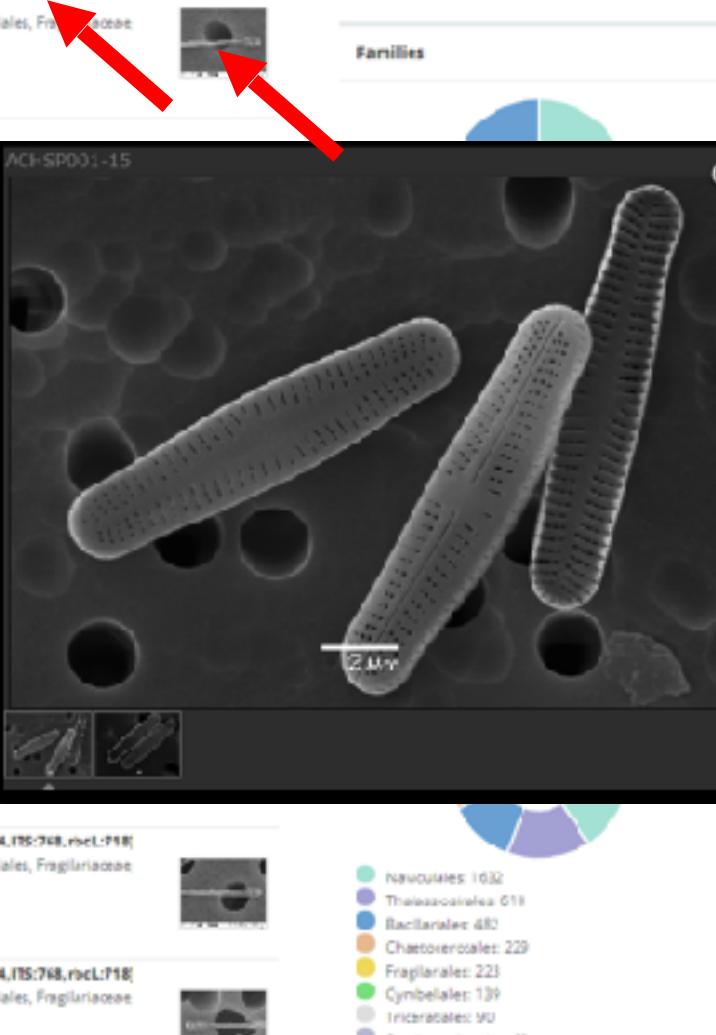
Taxonomy: Bacillariophyta, Bacillariophyceae, Fragilariales, Fragilariacae, Asterionellopsidales
Identifiers: IC_NPAA: 7[unspecified], M90[feldid]
Depository: Research Collection of I. Kaczmarska
Collected in: Canada, New Brunswick, Bay of Fundy

ASTBR008-13 - *Asterionellopsis tenellifera* [185:374,ITS:788,rcL:718]

Taxonomy: Bacillariophyta, Bacillariophyceae, Fragilariales, Fragilariacae, Asterionellopsidales
Identifiers: IC_NPAA: 8[unspecified], M91[feldid]
Depository: Research Collection of I. Kaczmarska
Collected in: Canada, New Brunswick, Bay of Fundy

ASTBR009-13 - *Asterionellopsis tenellifera* [185:374,ITS:788,rcL:718]

Taxonomy: Bacillariophyta, Bacillariophyceae, Fragilariales, Fragilariacae, Asterionellopsidales
Identifiers: IC_NPAA: 9[unspecified], M92[feldid]



Families



naviculales	1632
Thraustostomatidae	611
Bacillariales	482
Chaetoceratales	229
Fragilariales	223
Cymbellales	139
Tritrichales	90
Pseudosolenites	45





Reminders

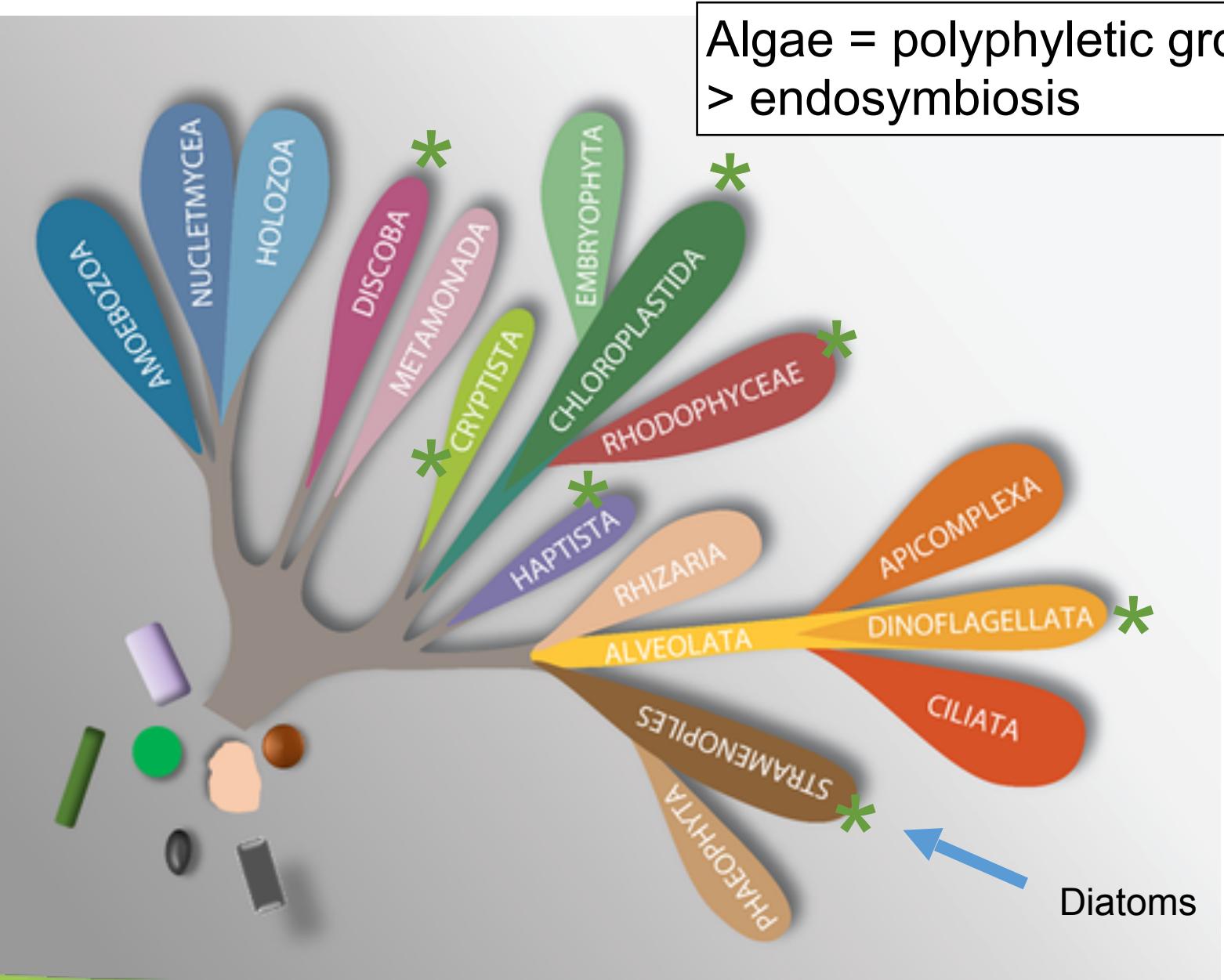
- 1- Diatom biology
- 2- Taxonomy
- 3- Classical diatom biomonitoring



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Adl et al. 2019.
Revisions to the
Classification,
Nomenclature,
and Diversity of
Eukaryotes. JEM



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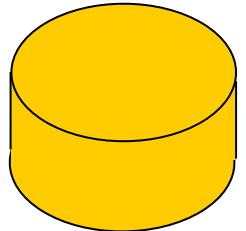
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Taxonomy

- Classification: 4 classes
 - Coscinodiscophyceae
 - Mediophyceae
 - Fragilapiophyceae
 - Bacillariophyceae



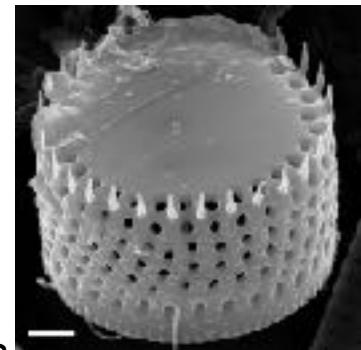
Taxonomy



Coscinodiscophyceae :

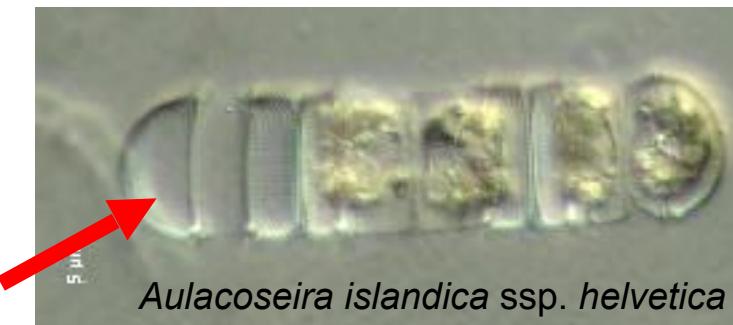
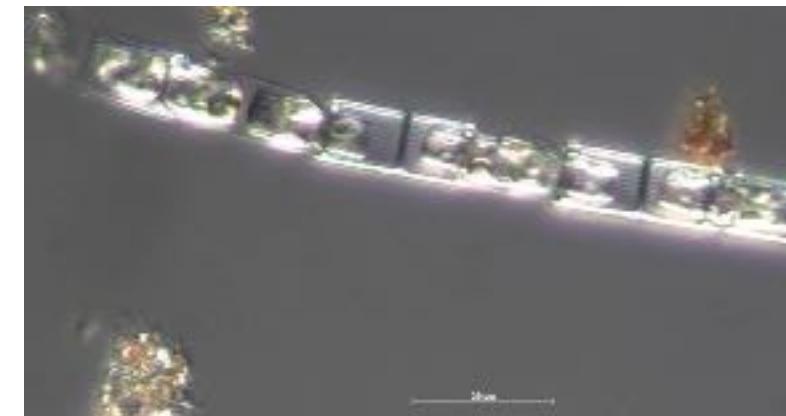
- *Aulacoseira* : filamentous, pores on the mantle, Spines between valve/mantle

- *A. alpigena*



Aulacoseira islandica (Leman fev 2010)

- *Aulacoseira muzzarellensis* (Canal Marne à Maixe 30-07-09)



Aulacoseira islandica ssp. *helvetica*



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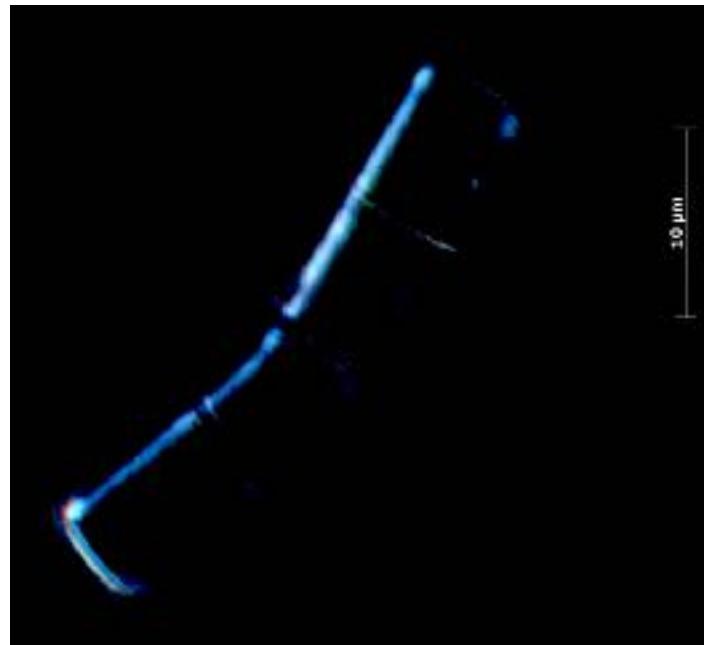
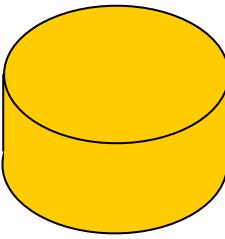
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Taxonomy

- **Coscinodiscophyceae :**

- *Melosira* : filaments, no pores

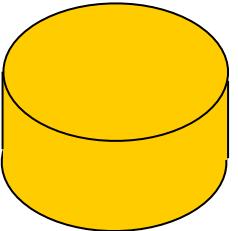
M. varians (TCC538), *lentic rivers, shore of calme lakes*



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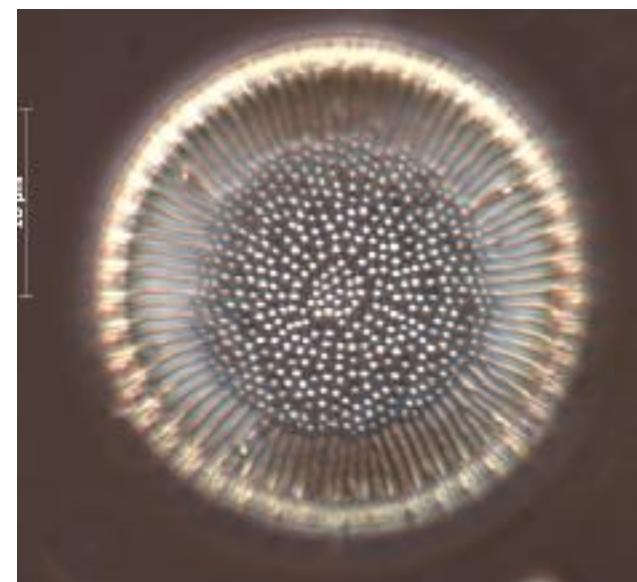
Taxonomy



- Mediophyceae :
- *Cyclotella* : single cell or in chains, planktonic



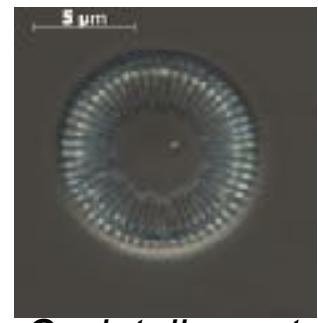
Cyclotella meneghiniana TCC358



Cyclotella lemanensis



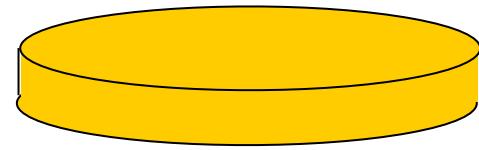
Cyclotella comensis



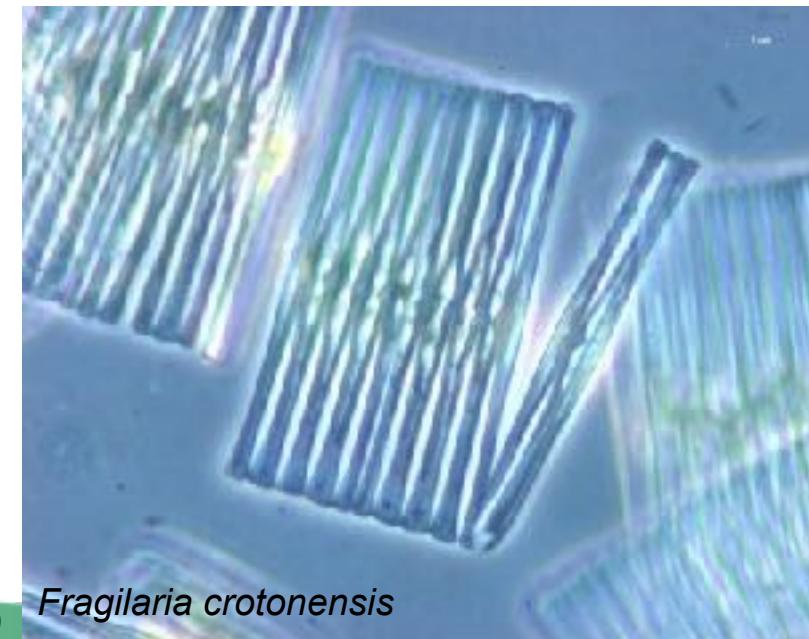
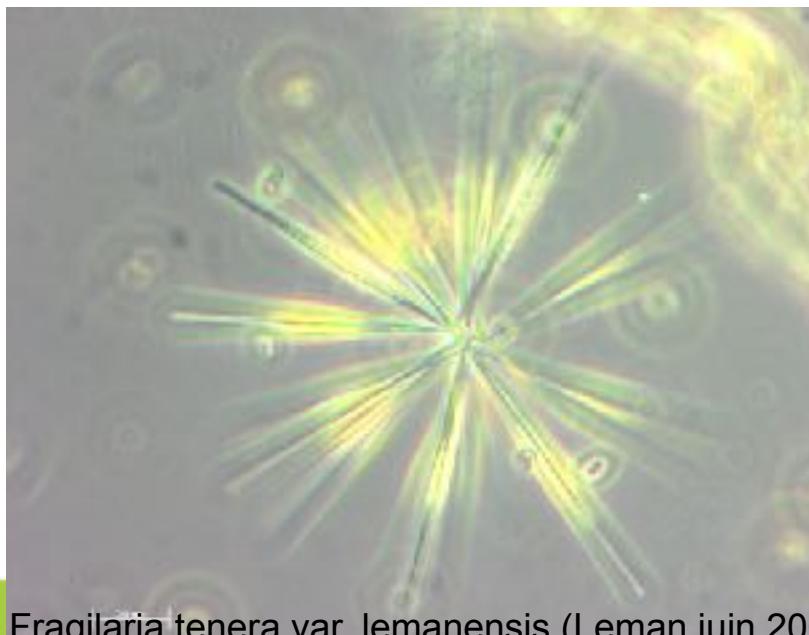
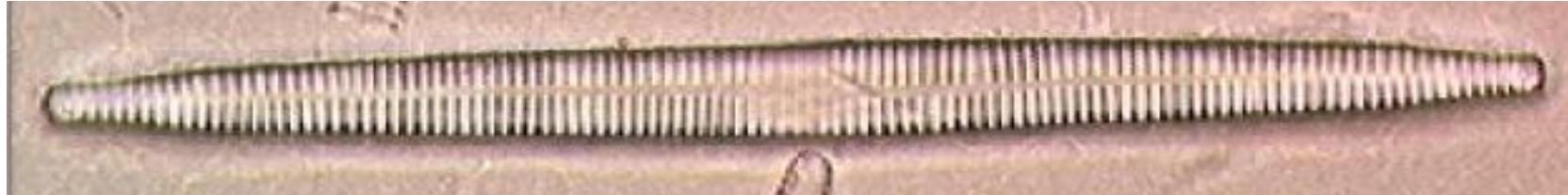
Cyclotella costei



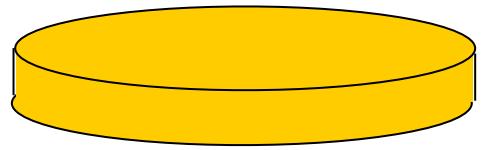
Taxonomy



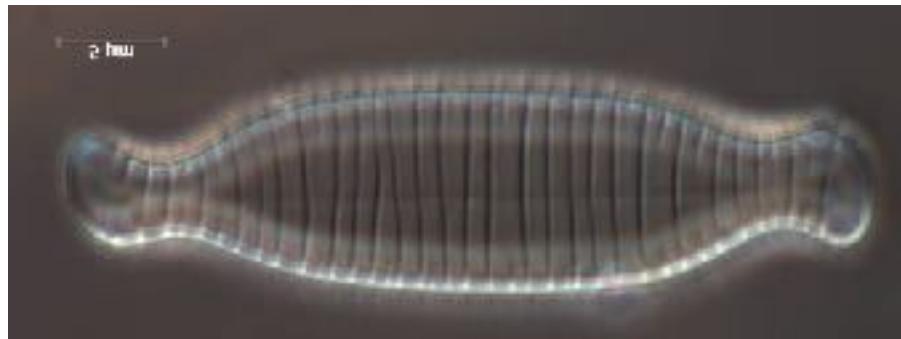
- **Fragilariphyceae** : F. Fragiliaceae
 - No raphe in this family
 - *Fragilaria/Ulnaria* : presence of a pseudo-raphe



Taxonomy



- **Fragilario phyceae** : F. Fragilariaeae
Diatoma: strong transversal costae



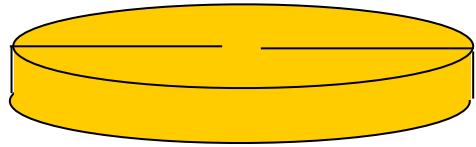
D. ehrenbergii



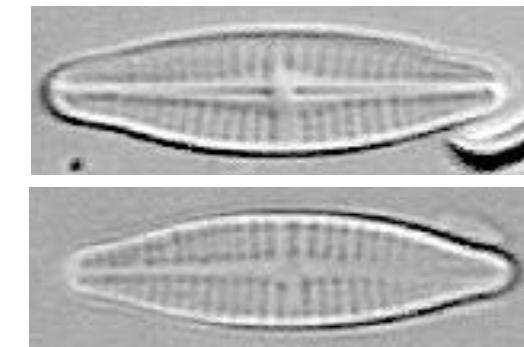
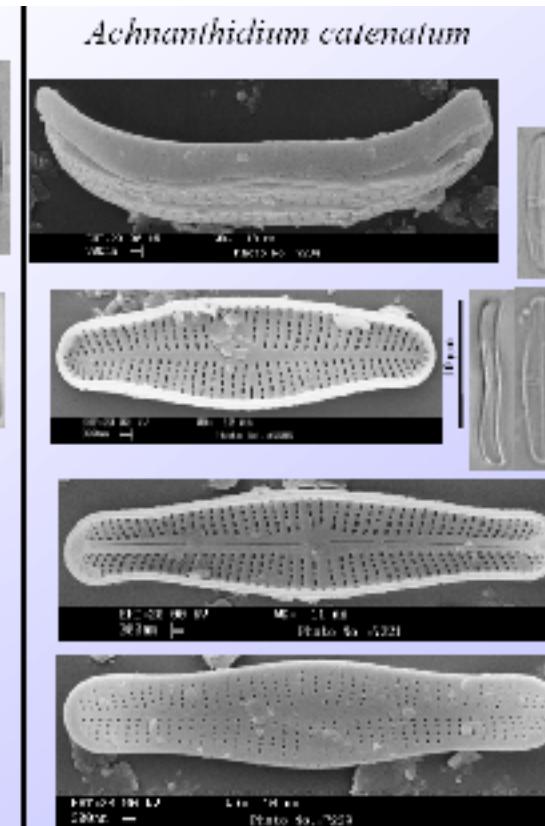
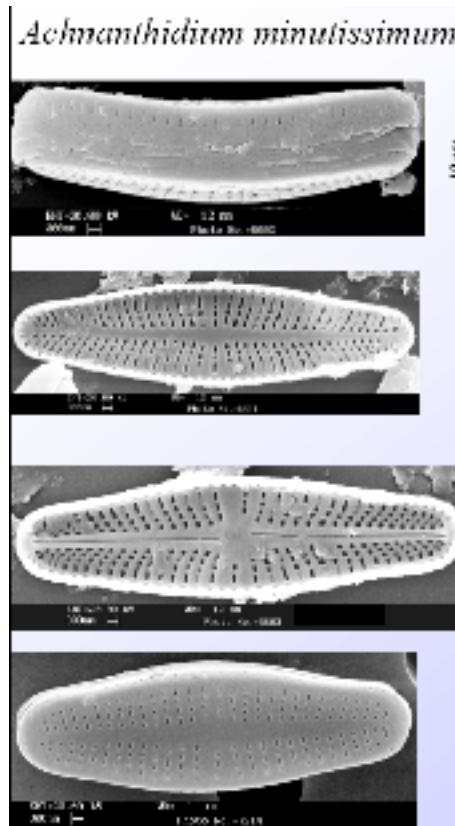
D. elongatum (Bourget 2009)



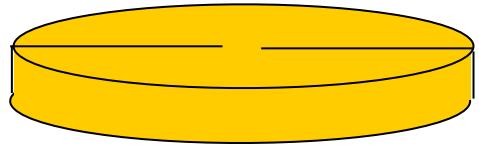
Taxonomy



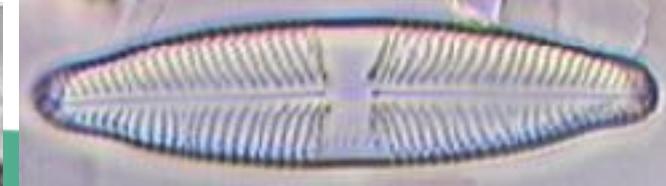
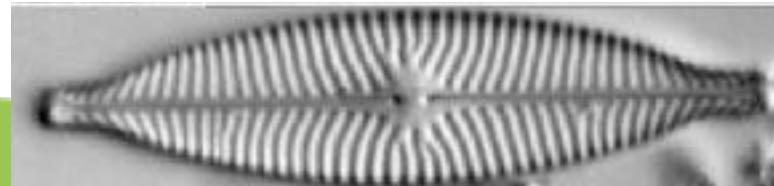
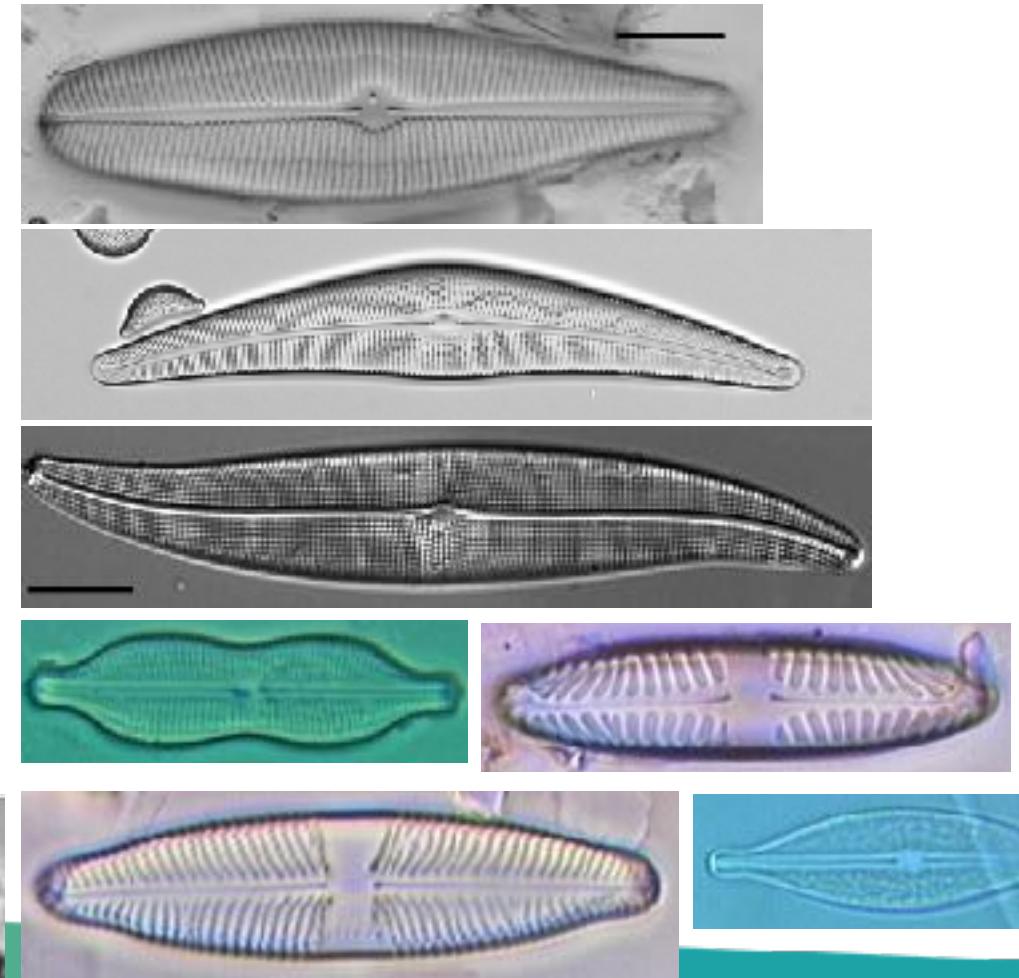
- **Bacillariophyceae:** F. Achnanthaceae, Achnanthidiaceae
Raphe present only on one valve (Monoraphids)



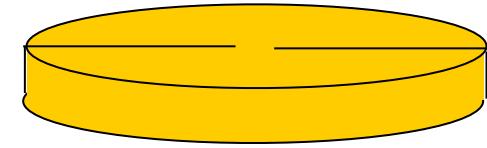
Taxonomy



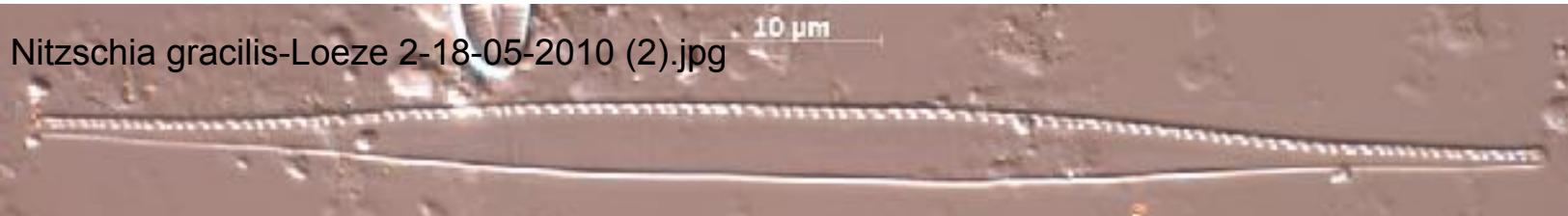
- **Bacillariophyceae** : Biraphids, raphe on the 2 valves
- Heteropolar valves:
- Isopolar valves:
 - Curved raphe
 - Sigmoid raphe
 - Straight raphe
 - Raphé avec 2 crochet au centre : Neidium
 - Cotes lisses : Pinnularia
 - Aire centrale en stauros : Stauroneis
 - Stries irrégulières, très fines : Brachysira
 - Navicula sensu stricto



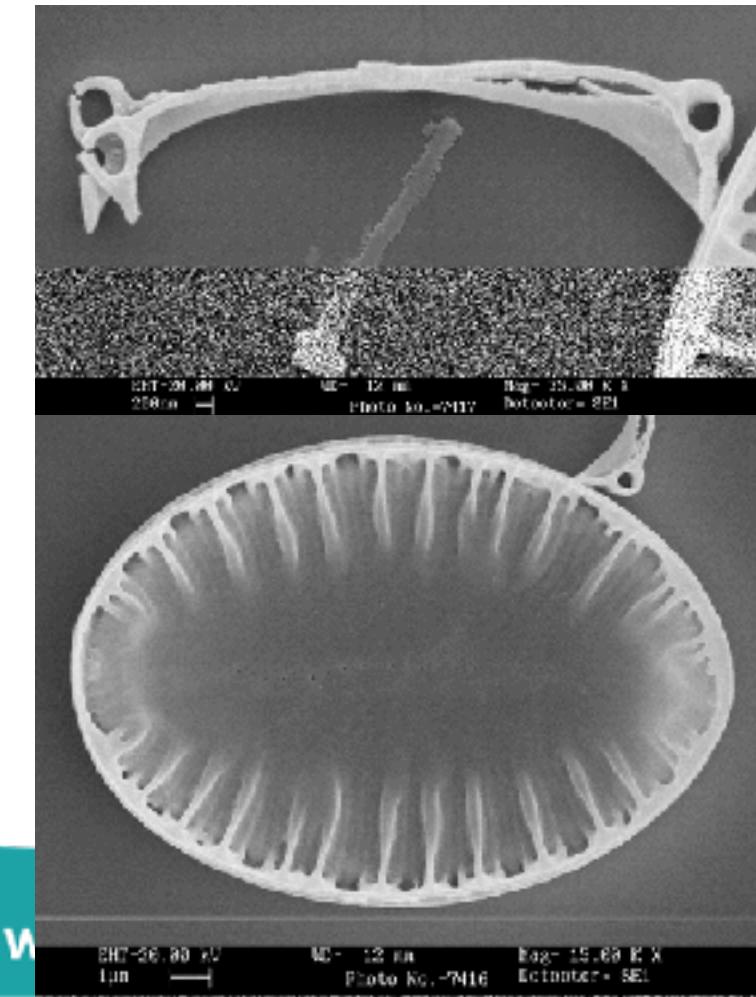
Taxonomy



- **Bacillariophyceae** : Biraphidées, raphé sur les 2 valves
- Heteropolar valves:
- Isopolar valves:
 - Curved raphe
 - Sigmoid raphe
 - Straight raphe
 - Raphé avec 2 crochets au centre : Neidium
 - Cotes lisses : Pinnularia
 - Aire centrale en stauros : Stauroneis
 - Stries irrégulières, très fines : Brachysira
 - Navicula sensu stricto
 - Canal raphe



Nitzschia gracilis-Loeze 2-18-05-2010 (2).jpg





Reminders

- 1- Diatom biology
- 2- Taxonomy
- 3- Classical diatom biomonitoring

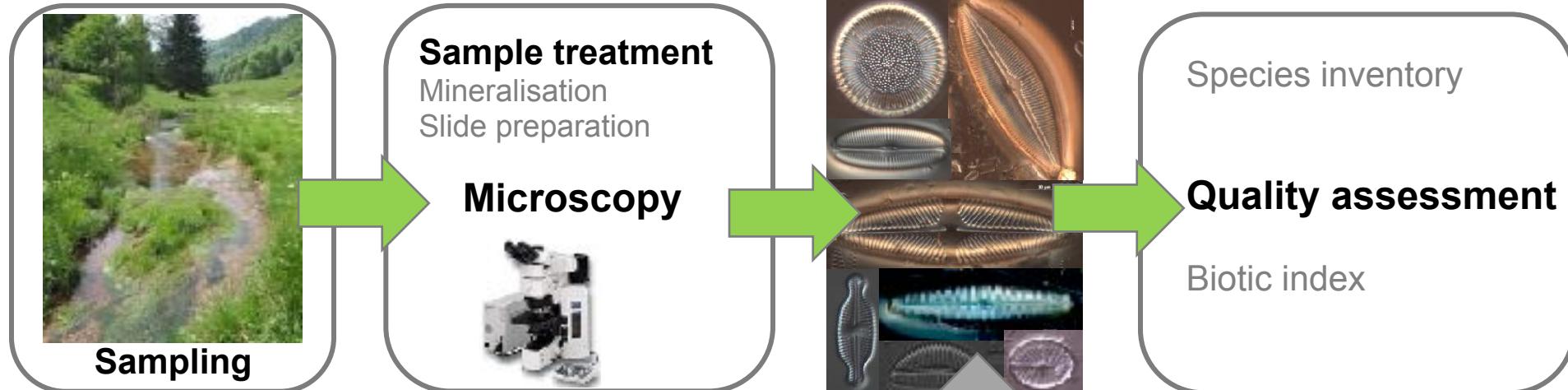


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Usual methodology for diatom biomonitoring

❖ Standardised methodology: microscopy



- Determination of species
- Counting 400 frustules

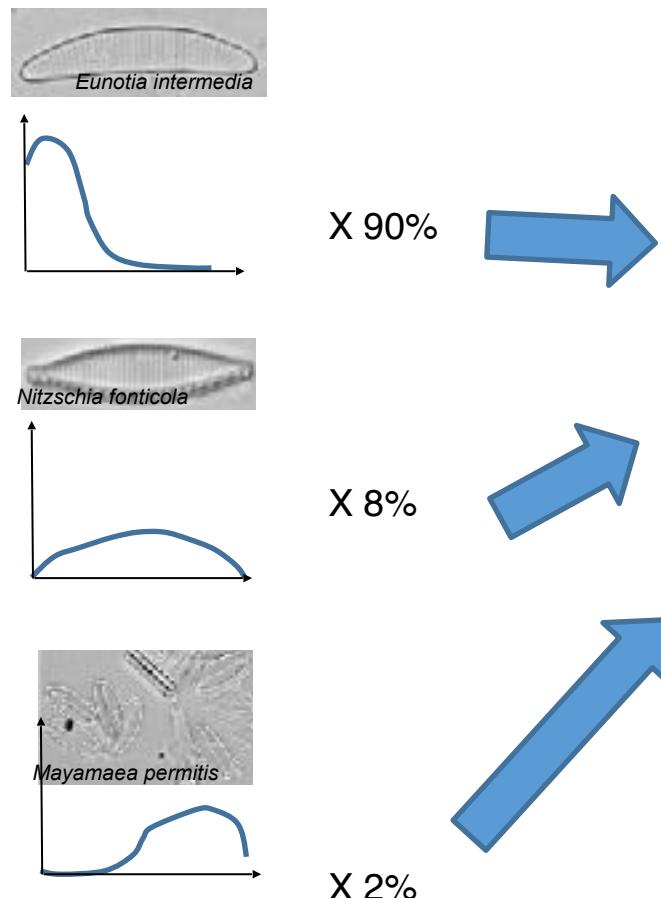
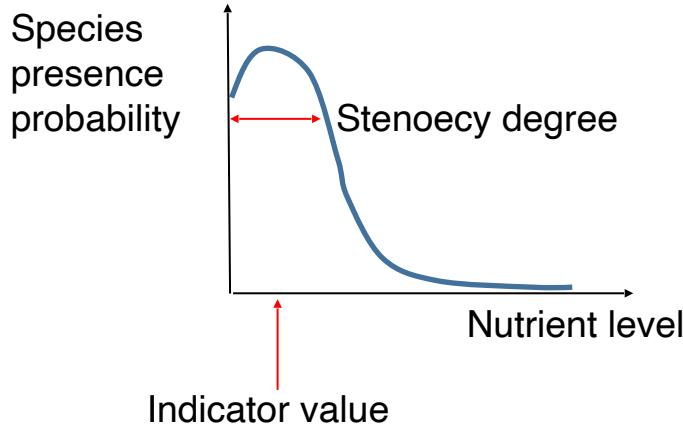


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Principle of a biotic index

Based on the formula of Zelinka & Marvan (1961): ecological profiles are weighted by the abundance of the species in the sample



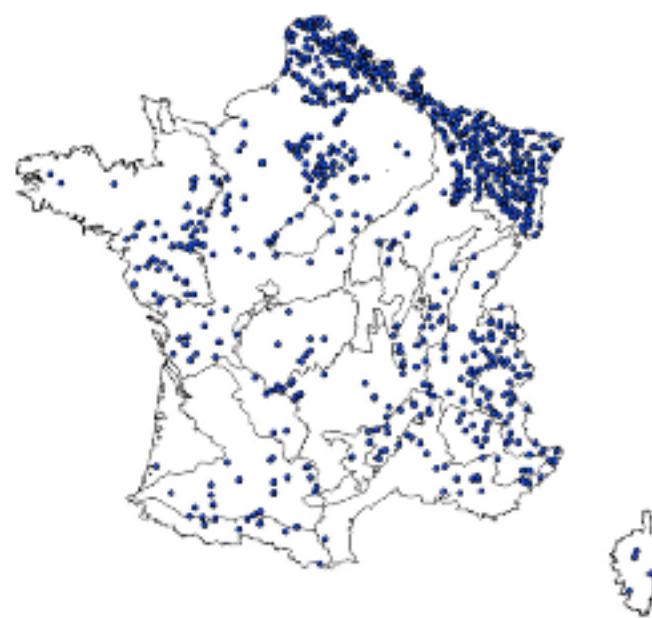
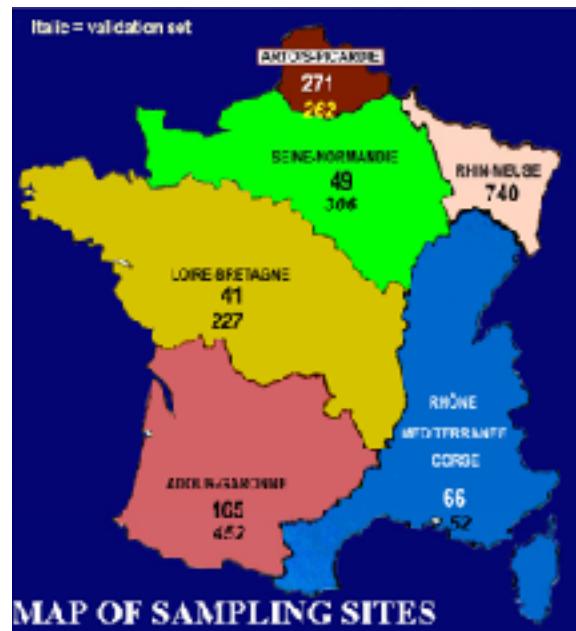
**Weighted average
Index = $\frac{\sum A_j \cdot v_i \cdot s_j}{\sum A_j \cdot v_j}$**

Aj : Species j abundance in the sample,
 sj : stenoecy degree (weight)
 vj : indicator value

Which biotic index is used in France?

IBD: Indice Biologique Diatomée

Index developed in the framework of the WFD, first publication in 2000, and also standardised in 2000 (Afnor 2000).
New versions in 2009 and 2019.



How IBD is used to assess river ecological status?

River status is based on
3 quality parameters

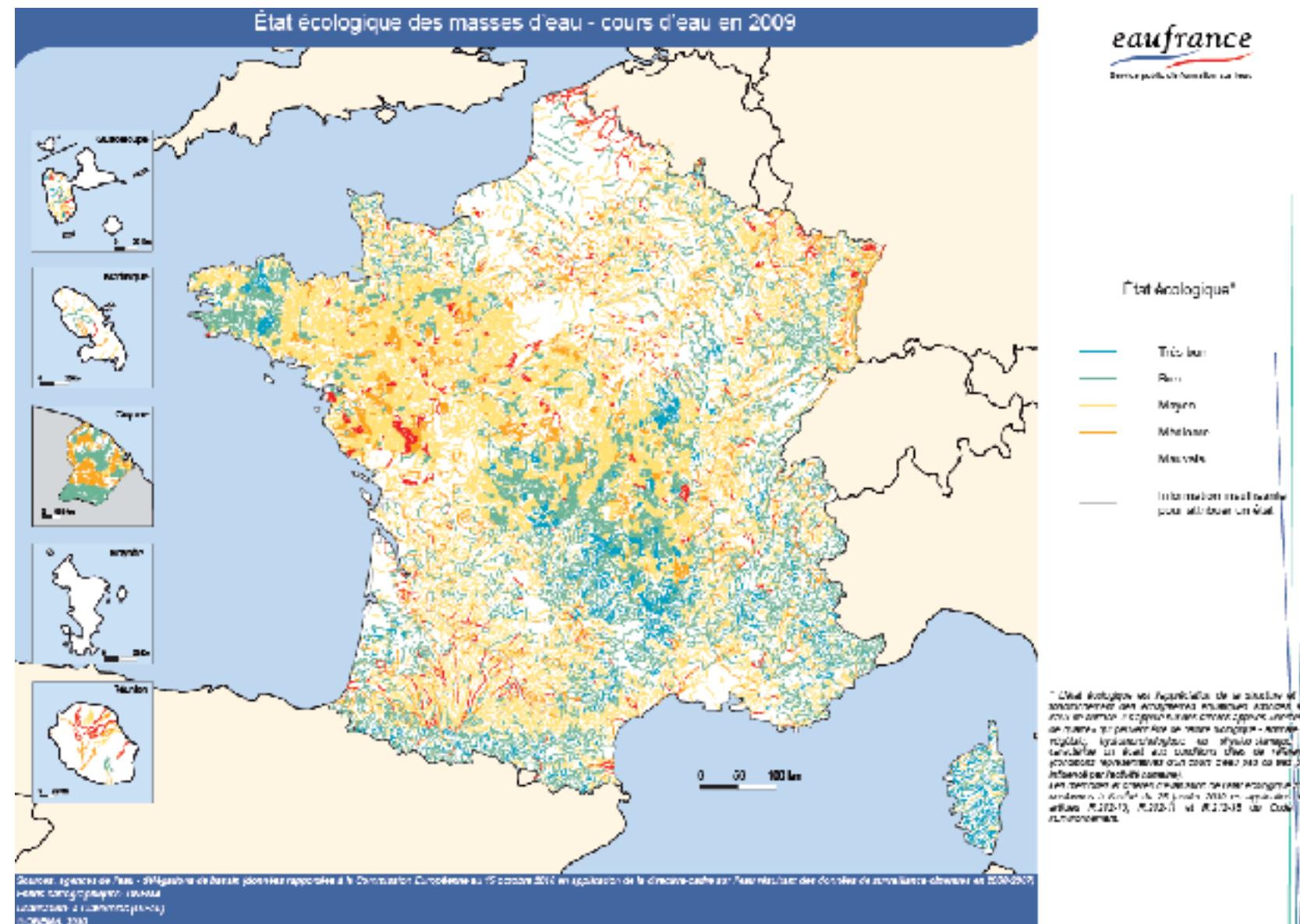
Hydro-morphological quality

Physico-chemical quality

Biological quality

Diatoms
Macroinvertebrates
Fish
Macrophytes

Ecological status of rivers in 2009



3 quality parameters

Hydro-morphological quality

Physico-chemical quality

Biological quality

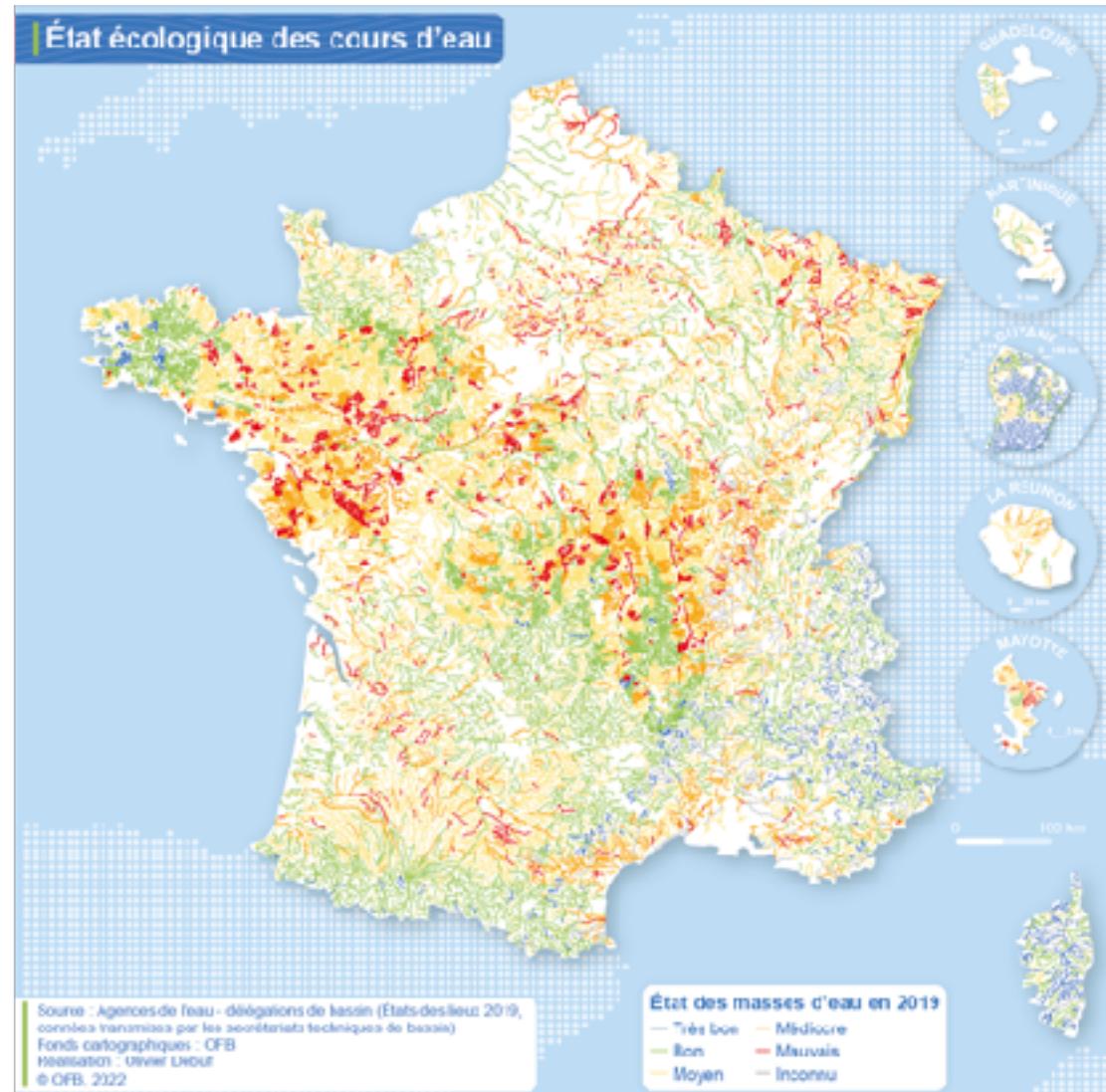
Diatoms

Macroinvertebrates

Fish

Macrophytes

Ecological status of rivers in 2019



3 quality parameters

Hydro-morphological quality

Physico-chemical quality

Biological quality

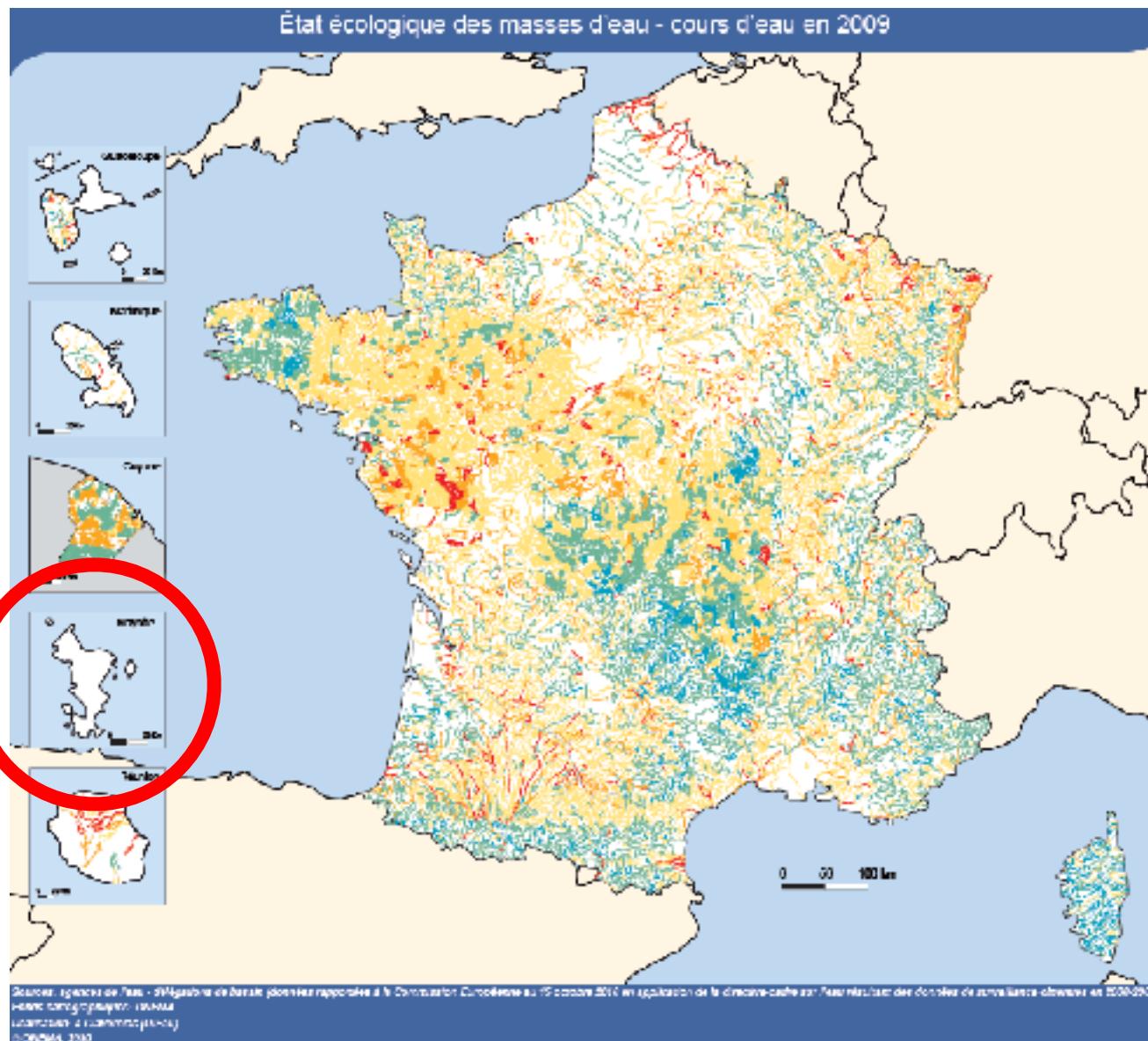
Diatoms

Macroinvertebrates

Fish

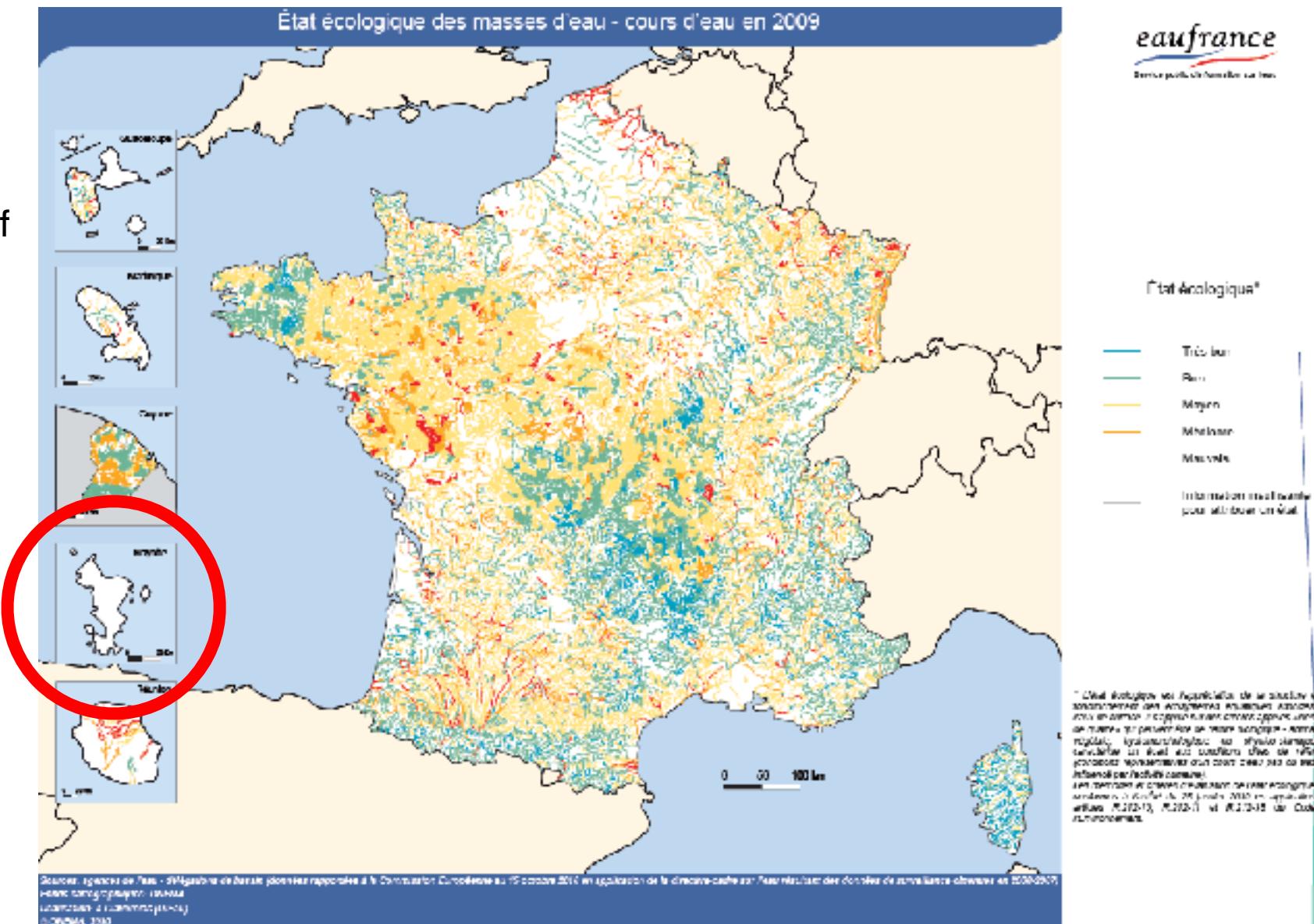
Macrophytes

Ecological status of rivers in 2009

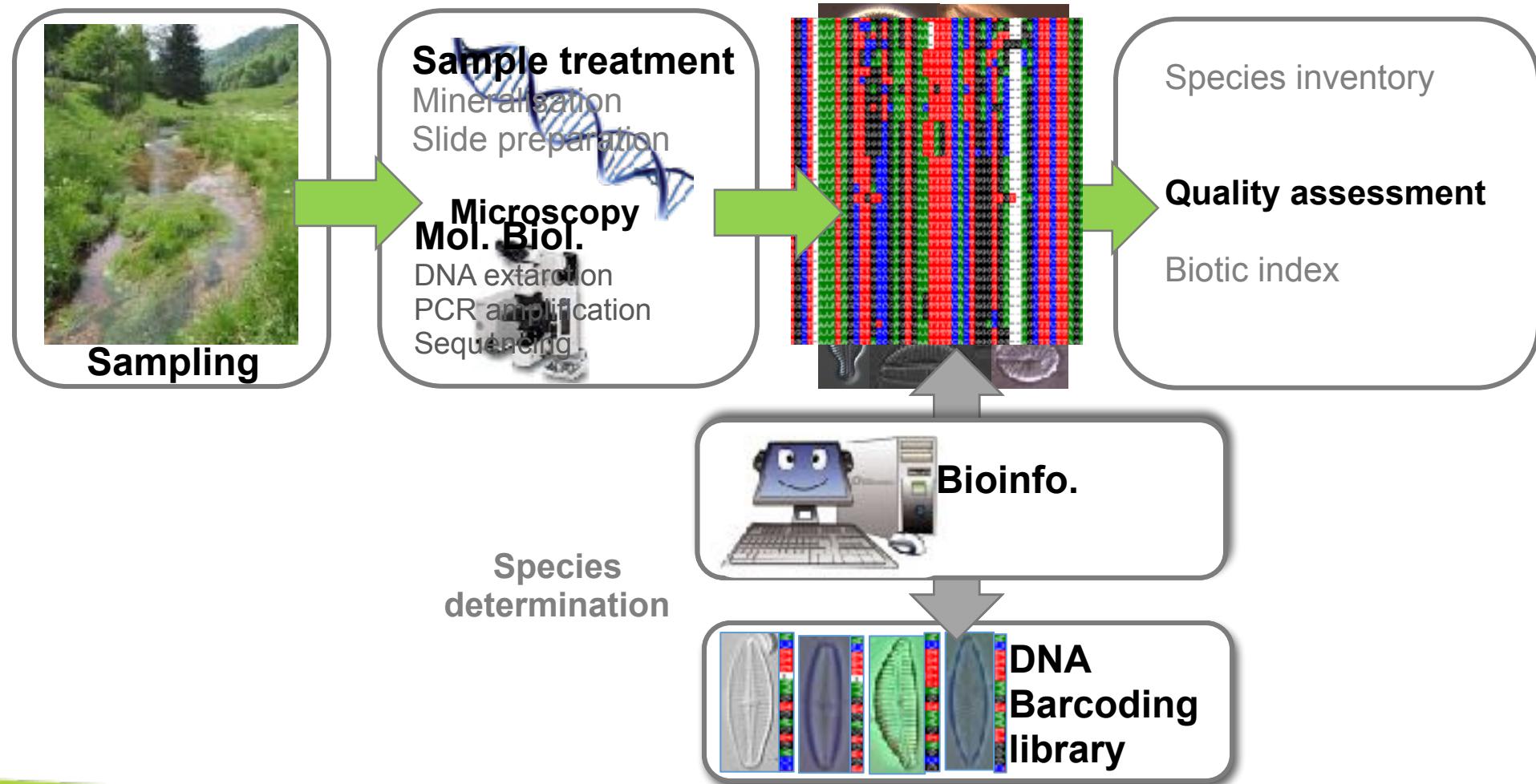


Ecological status of rivers in 2009

- No biomonitoring tools in 2009 in Mayotte
- We made a first proof of concept of diatom DNA metabarcoding for biomonitoring rivers in 2012 on 4 samples (Mainland France and tropical rivers)
- The french office in charge of biomonitoring asked us in 2013 to test alternative biomonitoring methods, including DNA metabarcoding in Mayotte



Objective: replace microscope determination and counting with DNA metabarcoding



Questions?



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