

Continuous and High Resolution Observation of the Mediterranean Sea



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Initiative d'excellence Aix-Marseille



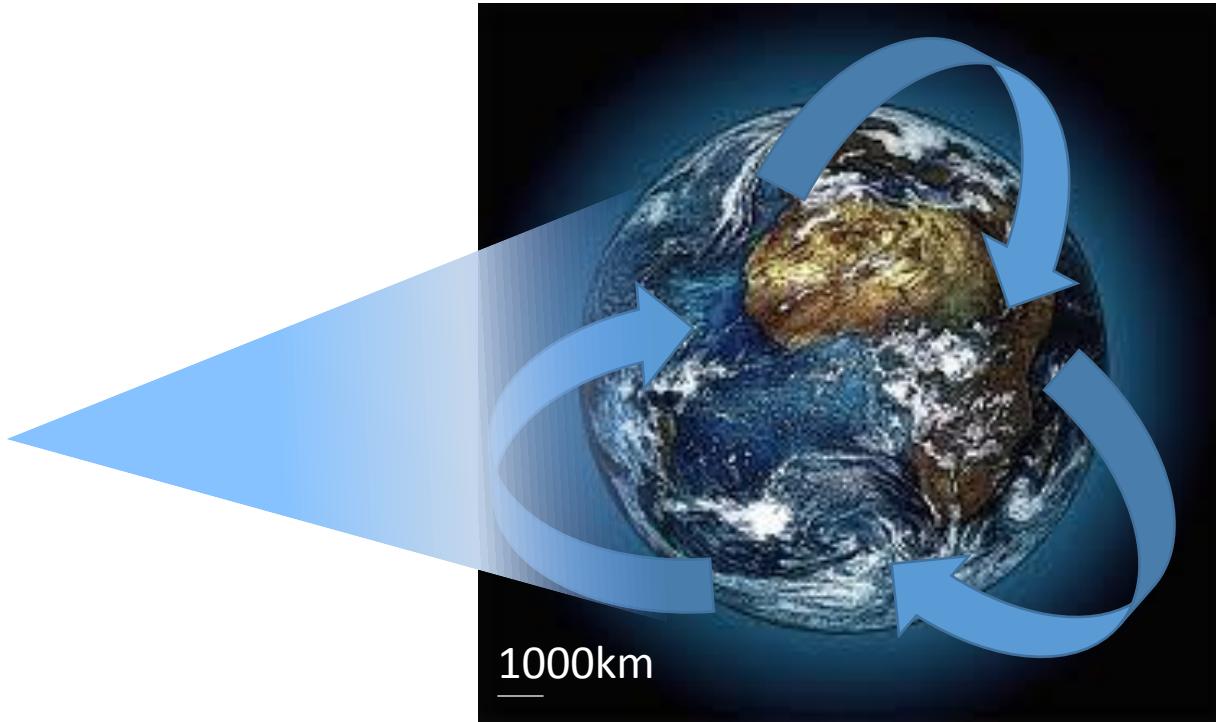
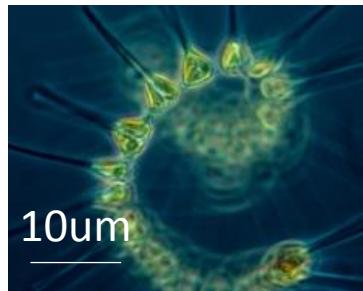
Deployment of automated flow cytometry on a FerryBox in the western Mediterranean Sea: Management and quality control of flow cytometry data

Soumaya Lahbib

M. Dugenne, M. Libes, M. Belhassen, A. Zouari, P. Marrec, N. Bhairy, G. Grégori , C. Sammari, S. Ben Ismail and M. Thyssen

Phytoplankton

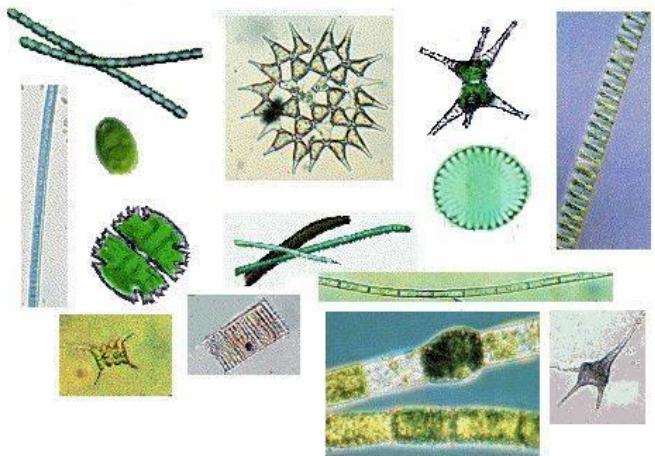
Thousands of species < 1000 μm catalyze the most important geochemical processes for sustaining life on earth AND at a minute scale.



Phytoplankton produces between 45 and 57 Pg C Yr^{-1} of the NPP on earth (~45%) but represents <2% of its biomass.
Very high turn-over rate !

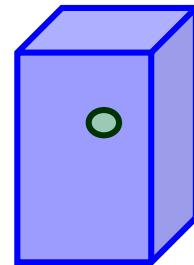
Phytoplankton observation is complex

Morphology and size

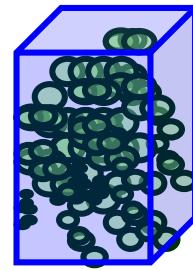


Abundances

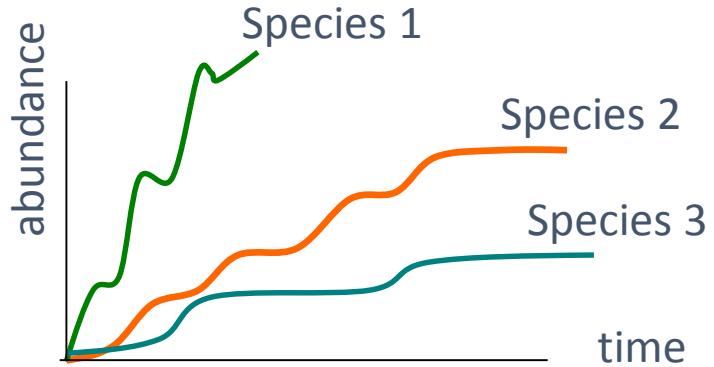
1 cell. cm^{-3}



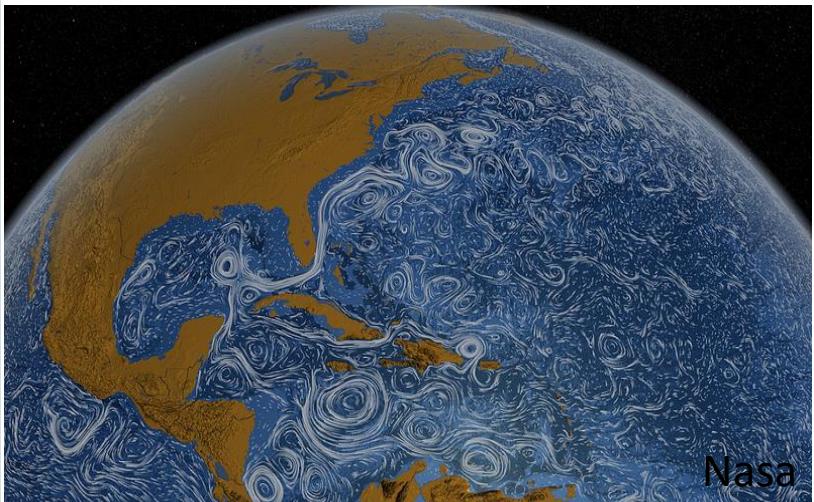
$10^6 \text{ cells. cm}^{-3}$



Growth rates



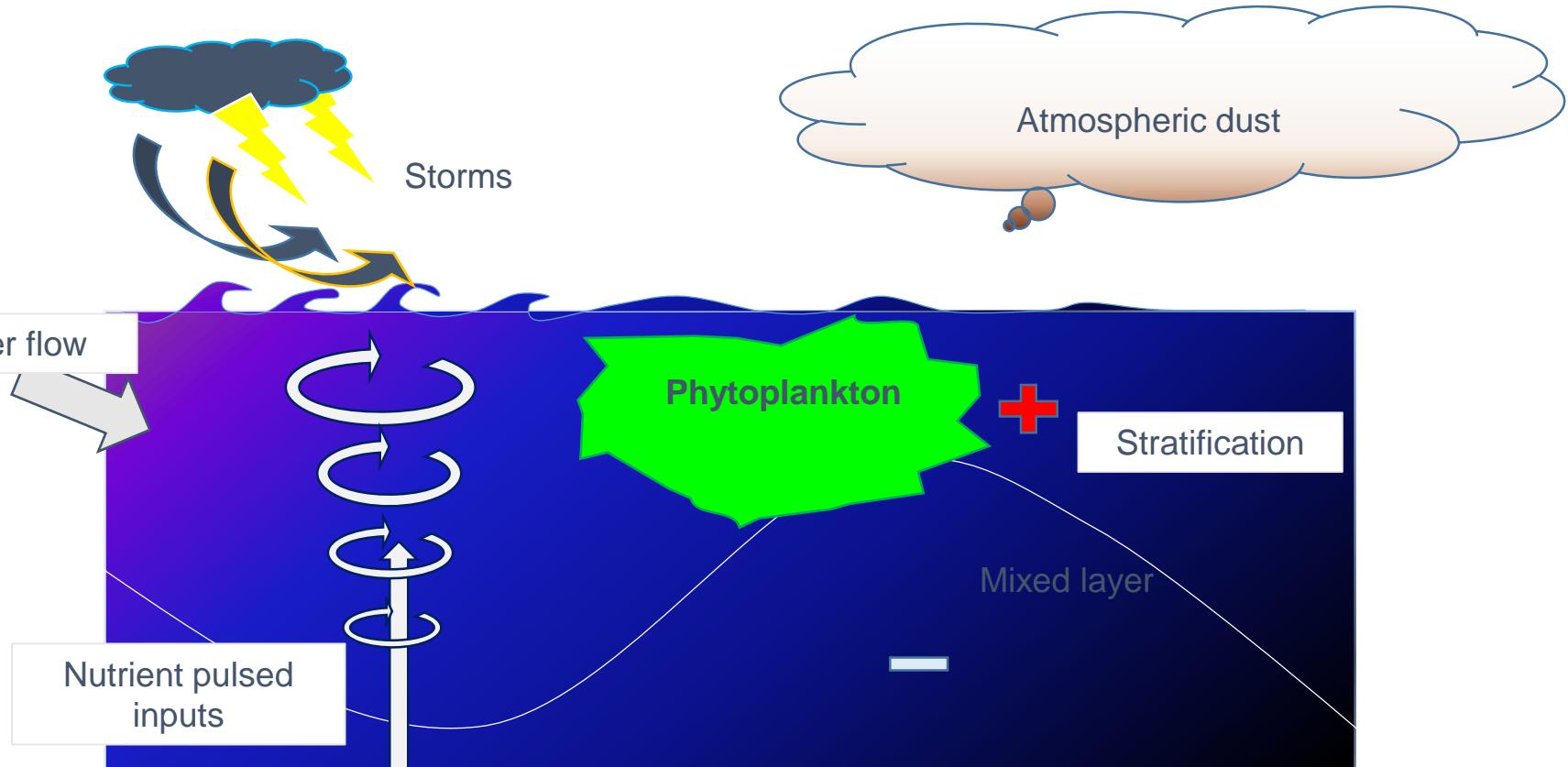
Turbulence



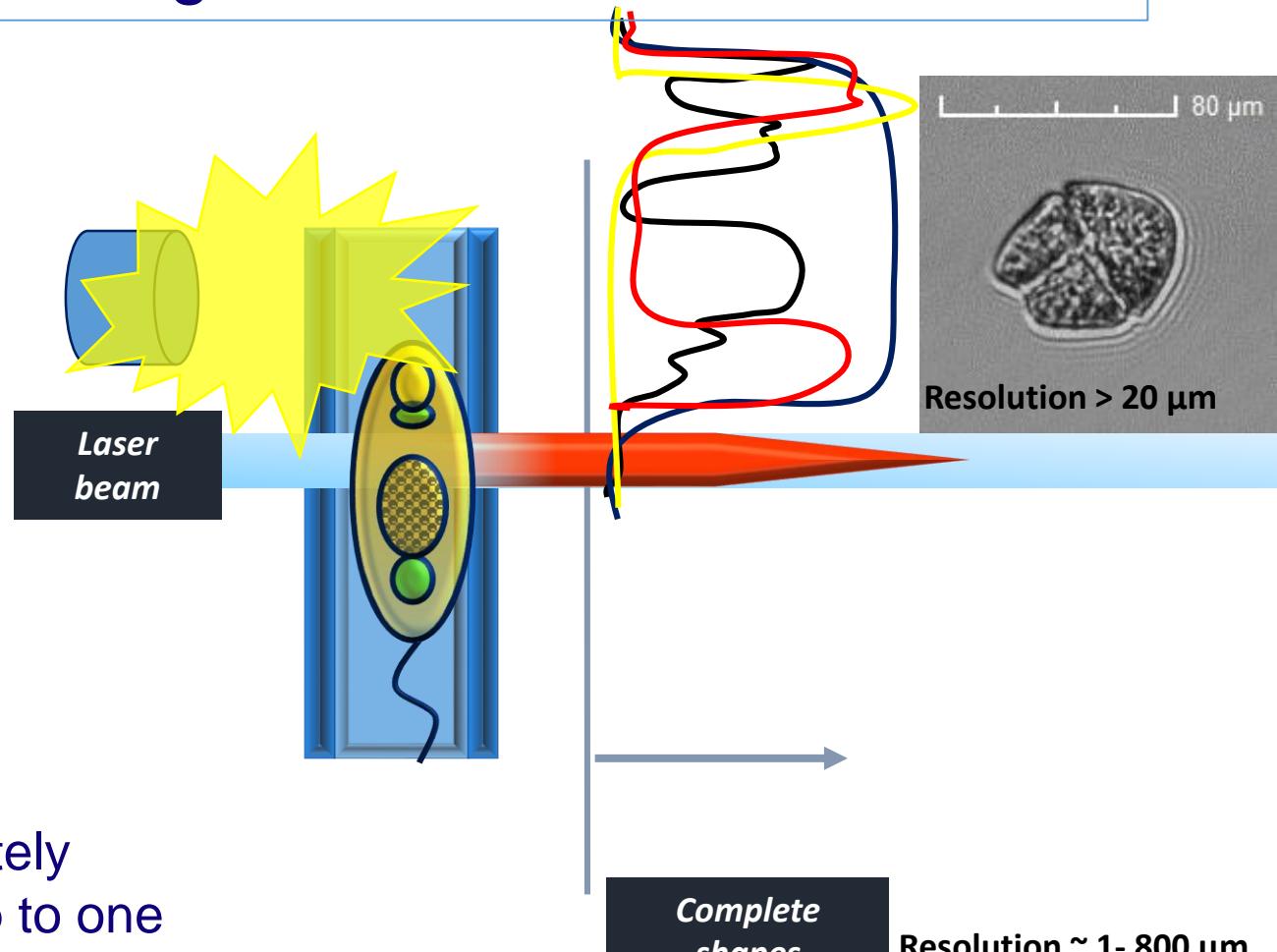
Nasa

Serious lack in understanding and
quantifying the role of phytoplankton in
the biogeochemical processes

Short term variation and sporadic events impacts are nearly unknown

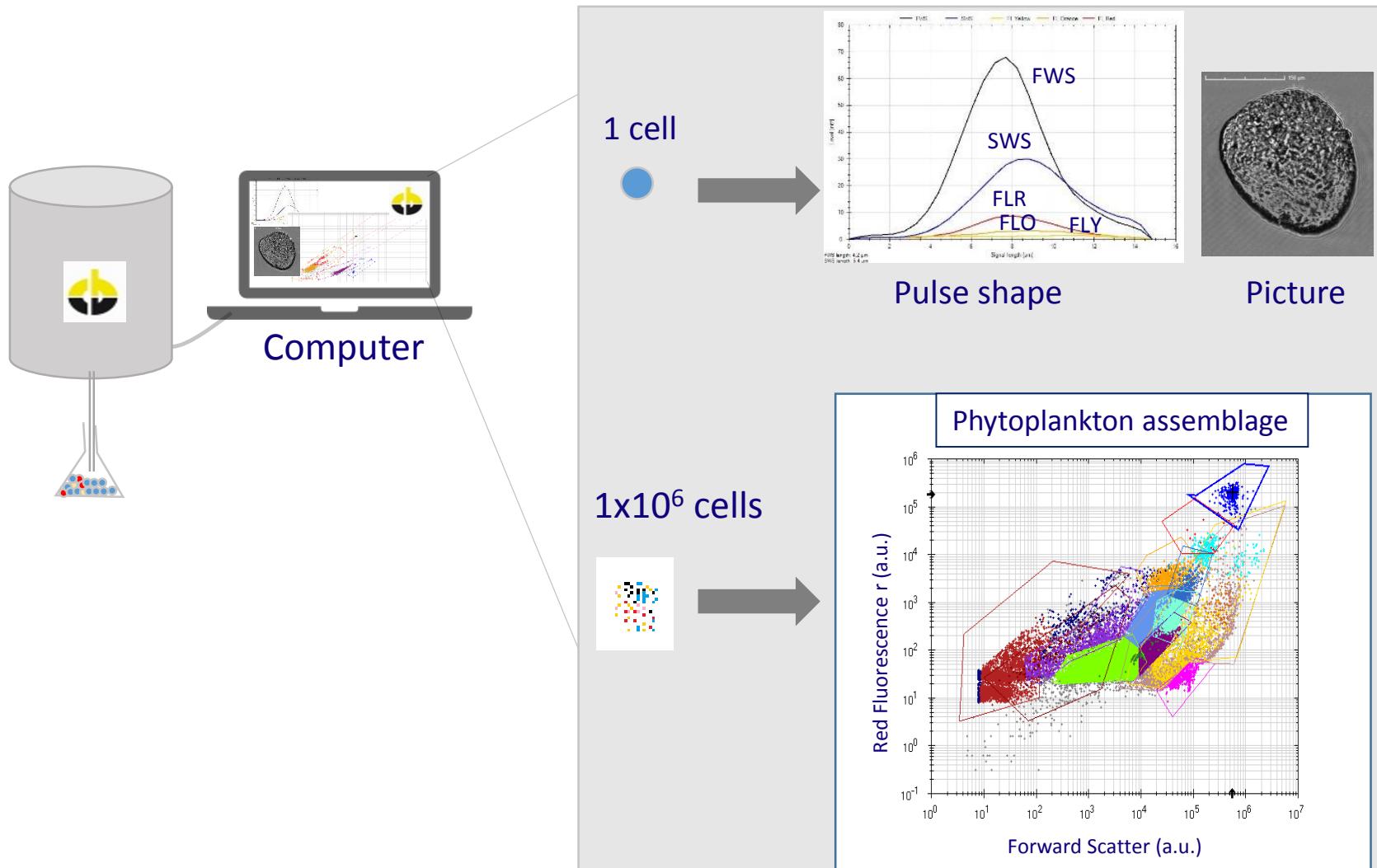


New technology for the resolution of phytoplankton functional diversity at hourly and regional scales



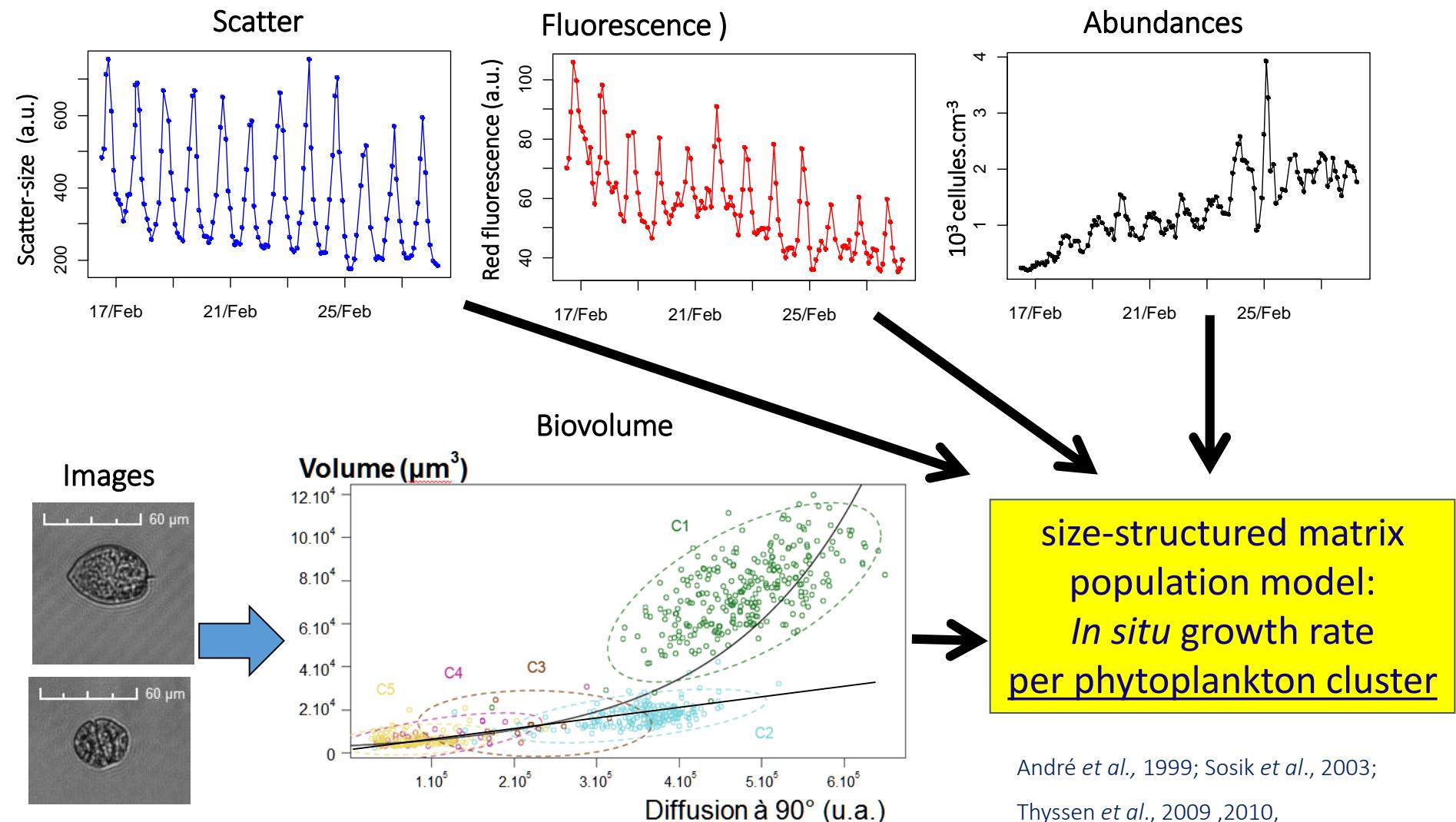
Automated and remotely controlled analysis up to one sample every 15 min.
Sample up to 5 cm³

Phytoplankton functional groups resolution



- Phytoplankton functional groups/Phytoplankton abundance per group
- Fluorescences/scatter per cell/Size estimation after calibration of scatter
- Phytoplankton images (taxonomical identification >20 μm)

Additional information extracted from the single cell approach:



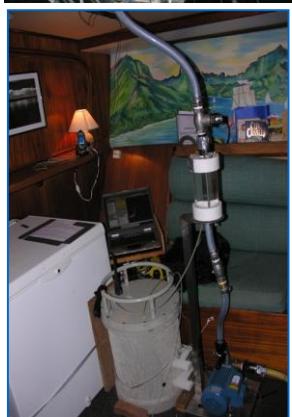
Several scientific experiences were conducted with a relative autonomie up to 6 months

Scientific vessels

Coastal platforms

Ships of opportunity

Buoys



Malkassian et al. 2011

Dugenne et al. 2014

Thyssen et al. 2008, 2009a,b, 2012, 2014, 2015





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Station Biologique
Roscoff

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de Marseille,
UMR 7373



CHROME

Continuous High Resolution Observation of the
Mediterranean Sea:

<https://chrome.mio.univ-amu.fr/>

*Understanding of the ecological and
biogeochemical functioning in relation to
meso-scale dynamics at the Mediterranean
sub-basin scale and weekly scale.*



Aix***Marseille**
université





Step 1: Data Acquisition= one analysis every 30 min.



— C/F CARTHAGE trajectories

DATASET 2

- Phytoplankton functional groups
- Phytoplankton abundance per group
- Fluorescences/scatter per cell
- Size estimation after calibration of scatter
- Phytoplankton images (taxonomical identification >20 µm)

Deck 1 – C/F CARTHAGE



Cherif Sammari



DATASET 1

- Temperature
- Salinity
- Fluorescence
- Turbidity
- pH
- pCO₂
- Oxygen



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Step 2: Data analysis

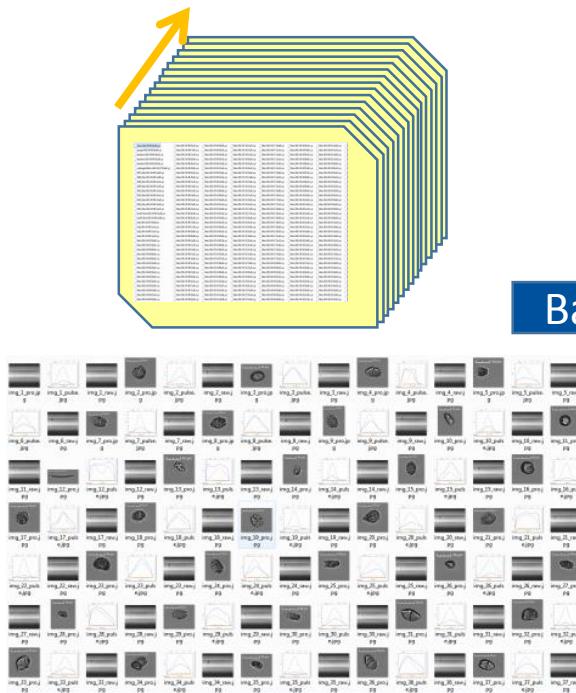


Measurements
by CytoSense



Output

Manual
clustering



Separate statistical CSV files:
Average values of optical
properties and Counts
+ Pictures

Backup



Issues:

1. Not a dedicated FCM database
2. Data memory size consuming

THE CYTOBASE DATABASE



Step 3 : Data management Workflow

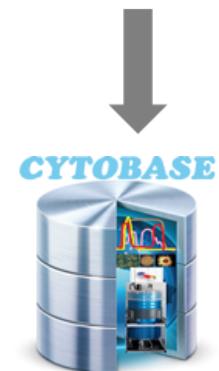


Data Acquisition

Batch Analyses

Data Consolidation

Expert QC



Data Integration



Step 4: Data consolidation

Output data files	Format	Number	Parameters
Average avgprops_process_xx-xx-xxxx avgprops_process_xx-xx-xxxx avgprops_process_xx-xx-xxxx etc..	csv	1 or many	Selection set, File name, Date, Time, Total Analyzed Volume [mL], Number of Particles Min. TOF, Max. TOF, Mean TOF, etc...
Count counts_process_xx-xx-xxxx counts_process_xx-xx-xxxx counts_process_xx-xx-xxxx counts_process_xx-xx-xxxx etc...	csv	1 or many	File name, Volume [mL], Trigger chn, Level [mV], Date, Particles, Beads 2 mu - Number, Beads 2 mu - pct/Tot, Beads 2 mu - N/mL, Microphytoplankton - Number, etc...



Cytobase input processor is a standalone software (R) by **Mathilde Dugenne** (PhD Student) in 2015. It aims to create raw data and metadata.

- ✓ Generate metadata about the project, the used instrument, the analysis methode, and the project operator
- ✓ Consolidate, aggregate and classify output data files
- ✓ Couple phytoplankton group names with standardised names
- ✓ Couple pictures with measured files
- ✓ Calculate size estimation from calibration procedure
- ✓ Generate picture and data table files interoperable with CYTOBASE



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Cytobase Input Processor

2015-08-18

Parcourir... 11 fichiers sélectionnés.

Upload complete

Warning:

Project and samples context

Raw data

Size conversion

Project

Enter project name

Project date

2015-08-18

PI

Enter PI name

Cytometer ID

Enter cytometer ID

Station

Depth

Latitude

Longitude

Metadata

Filename

BERRE_082013_3F_FLR9 2013-12-17 13l

Samples operator

Enter name of operator

Standards reference

Enter standards beads ref

Clustering method

Automated

Observation type

In situ

Data

2013-12-17T14:17:00Z	2013-12-17T14:17:00Z	16,056.60		Synechococcus	BERRE_082013_12S_FLR9 2013-12-17 14u17.cyz	1.99	FL Red	10
2013-12-17T14:17:00Z	2013-12-17T14:17:00Z	16,056.60		Cryptophytes	BERRE_082013_12S_FLR9 2013-12-17 14u17.cyz	1.99	FL Red	10
2013-12-17T14:43:00Z	2013-12-17T14:43:00Z	16,056.61		Beads 2 mu	BERRE_082013_16F_FLR9 2013-12-17 14u43.cyz	2.04	FL Red	10
2013-12-17T14:43:00Z	2013-12-17T14:43:00Z	16,056.61		Microphytoplankton	BERRE_082013_16F_FLR9 2013-12-17 14u43.cyz	2.04	FL Red	10
2013-12-17T14:43:00Z	2013-12-17T14:43:00Z	16,056.61		Picoplankton 2	BERRE_082013_16F_FLR9 2013-12-17 14u43.cyz	2.04	FL Red	10
2013-12-17T14:43:00Z	2013-12-17T14:43:00Z	16,056.61		Picoplankton 1	BERRE_082013_16F_FLR9 2013-12-17 14u43.cyz	2.04	FL Red	10

Please associate each selection set to trigger, PMT's amplification and standardized phytoplankton category

NB: All incompatible entries will be removed

Expert name

Cluster

Beads 2 mu

Trigger

Channel/Level

FL Red 10

PMT's amplification

SWS

50

FLO

50

100

Standardized name

Cluster

Standard beads

Associate



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Picture selection

Project and samples context Raw data Size conversion Image-In-Flow pictures Stations explorer

Add samples pictures

Select file: DEWEXL2FLR10 2013-04-05 15u04.cyz

Check all

Add pictures: Choisir les fichiers 15 fichiers Upload complete

Show 4 entries Search:

Stations Explorer

Project and samples context Raw data Size conversion Image-In-Flow pictures Stations explorer

Select station: S1

X axis: Mean Total FWS

Y axis: Mean Total FLR

Scatter plot:

Map view showing a route path from France to Italy. The path is highlighted in blue. An inset map shows the location of Lazio, Italy, with a 40 μm scale bar.

Legend for scatter plot:

- PicoRED
- PicoORG
- NonaSWS
- New Set 4
- New Set 3
- NanorORG1
- NanoRed
- NanoORG2

Download Table

Data Table
Picture Table



Step 5: Data integration into CYTOBASE

Info Tables Columns Indexes Triggers Views Stored Procedures Functions Grants Events

CytoBase_Local
cytobase_v6

Schema Details

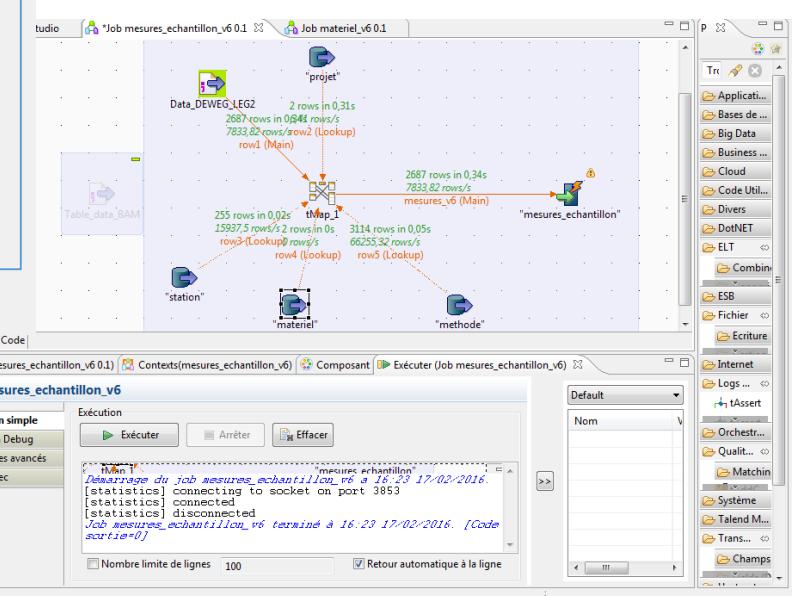
Default collation: latin1_swedish_ci

Default characterset: latin1

Table count: 7

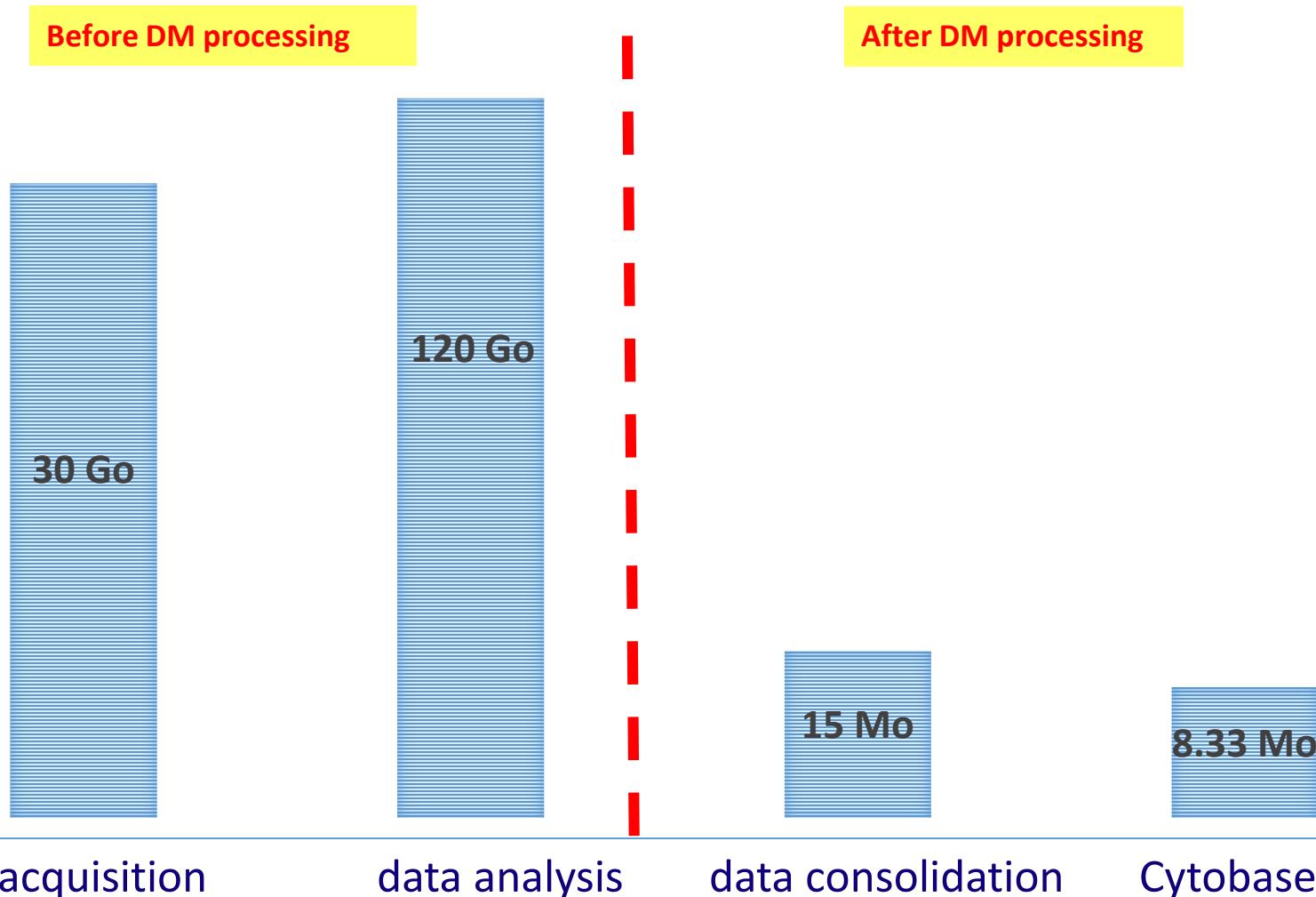
Database size (rough estimate): 5.8 MiB

- 7 tables
- 6 associations
- 69 columns





- Case of a 9 days Cruise : 1 sample / 20 min



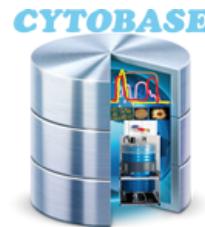


Step 6: Data retrieval and accessibility

- ✓ Web Developpement



- ✓ Access to CYTOBASE





V. Conclusion & Perspectives

- Smart storage and sustainability → Net decrease on file size
- Dynamic and user friendly web-based interface
- Deployment of CYTOBASE (*In progress*)
- Working on international standardization and interoperability with SeaDataNet Pan-European Infrastructure.



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Thank you for your attention

Any questions?