



1^{er} Atelier technique

« Bonnes pratiques techniques au service de l'interopérabilité »

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Campus Luminy, Marseille

Brest, 11 et 12 septembre 2017



Centralisation des jeux de données de l'OSU

En cours par Maurice Libes

Géo-cataloguage

<http://139.124.2.98/geonetwork/srv/fre/catalog.search#/home>

The screenshot shows a GeoNetwork catalog search interface. At the top, there are two search bars: 'Rechercher ...' and 'Rechercher ...'. Below them is a button 'Rechercher parmi 17 jeux de données, services et cartes, ...'. The interface includes filters for 'Parcourir par Thèmes INSPIRE topics': Conditions atmosphériques (1), Caractéristiques géographiques (2), Régions maritimes (17), Habitats et biotopes (1), and Hydrographie (1). There are also filters for 'Types de ressource': Jeu de données (17). A sidebar shows 'Nouveautés' and 'Les plus vues'. The main area displays a grid of dataset cards:

- ROMARIN – Réseau d'Observation du domaine MARIN** (*Jeu de données*)
Sites d'étude : Zone Proximale de la Baie de Marseille - SOLEMIO Baie de Marseille 05° 17' 30 E - 43° 14' 30 N (3 milles de la
- Météo Frioul** (*Jeu de données*)
- Suivis du panache rhône mer** (*Jeu de données*)
- SOLEMIO – Mesures haute fréquence paramètre physico-chimique Baie de Marseille** (*Jeu de données*)
- PhytoMed - Phytoplancton Baie de Marseille** (*Jeu de données*)
- Modèle océan** (*Jeu de données*)
- HTMNED Hydrodynamique et Transport de Matière en Suspension - Niveaux d'Eau et Températures** (*Jeu de données*)
- Quantification des apports rhodaniens à la mer** (*Jeu de données*)
- ANTARES** (*Jeu de données*)

Visualisation



ERDDAP > List of All Datasets

Pick a Dataset

1340 matching datasets, listed in alphabetical order. View page: 1 (current) 2 .

Grid DAP Data	Sub-set	Table DAP Data	Make A Graph	W M S	Source Data Files	Accessible	Title
set	data	graph				public	* The List of All Active Datasets in this ERDDAP *
data		graph				public	AMSRE Model Output, obs4MIPs NASA-JPL, Global, 1 Degree, 2002-2010
data		graph	M			public	AMSRE Model Output, obs4MIPs NASA-JPL, Global, 1 Degree, 2002-2010
		data	graph		files	public	AN EXPERIMENTAL DATASET: Underway Sea Surface Temperature and S

Equipe de Cytométrie en flux

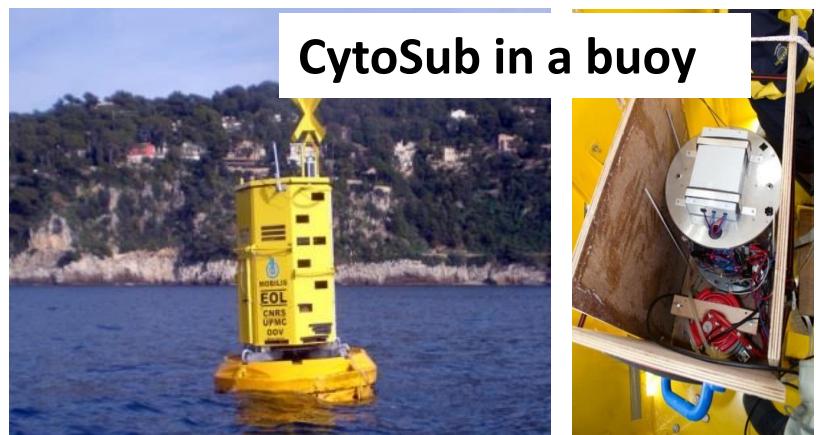


CytoPro

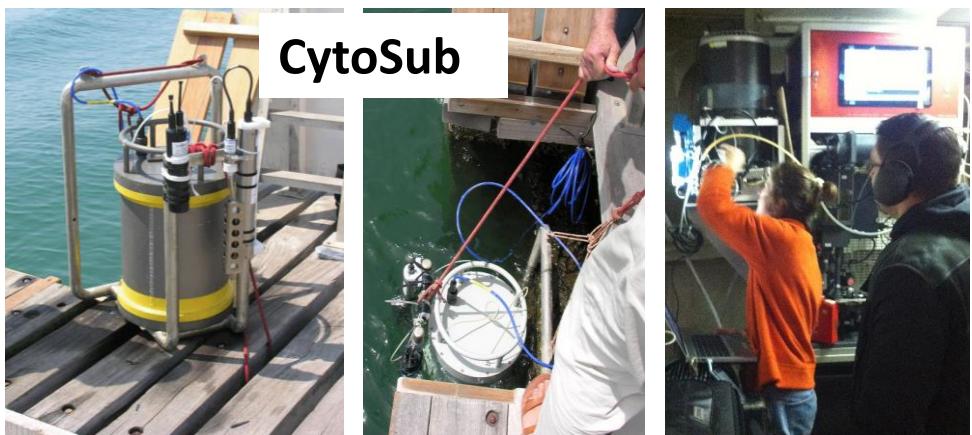


CytoSense

FerryBox

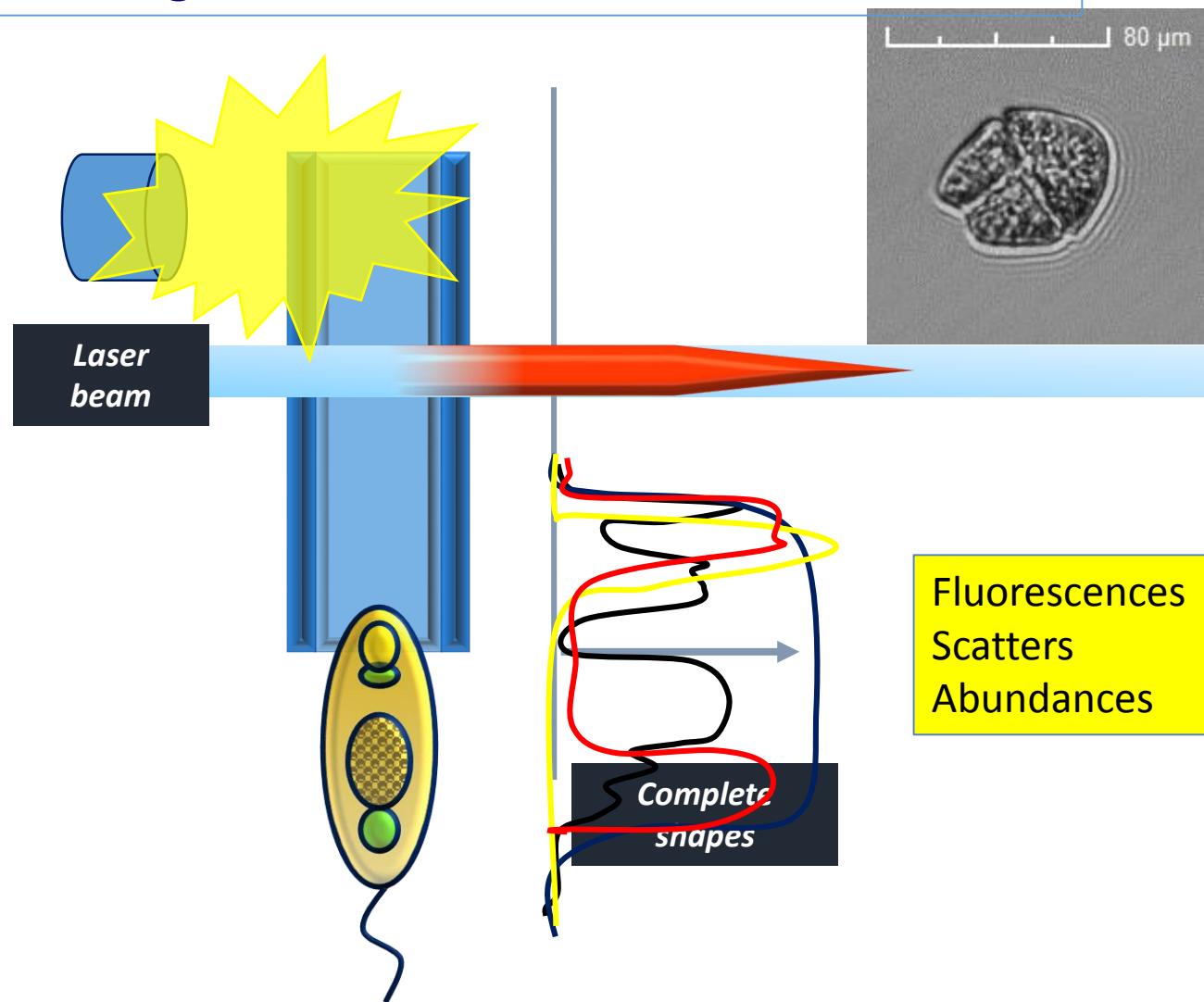
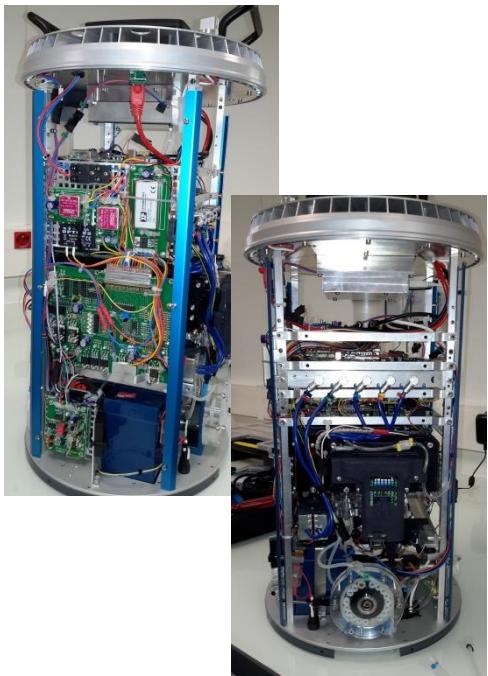


CytoSub in a buoy



CytoSub

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





CNRS UPMC INSU
Station Biologique
Roscoff

Institut de
Mathématiques
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



A*MIDEX CHROME Project

Data Acquisition = one analysis every 30 min.

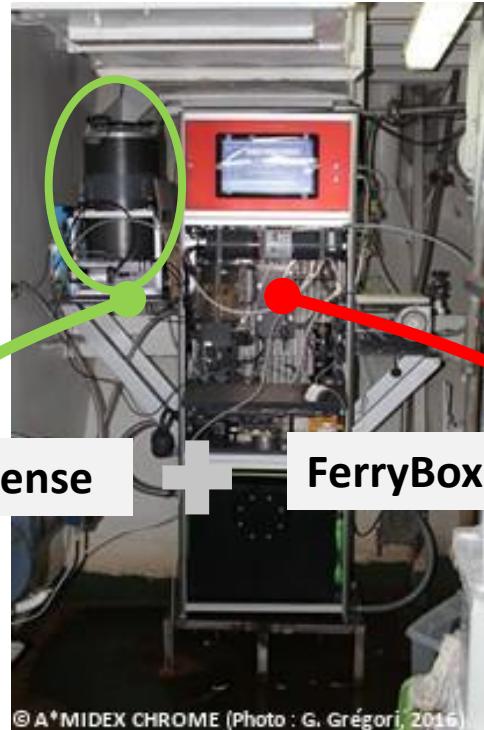


— C/F CARTHAGE trajectories

./30min

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

Pont 1 – C/F CARTHAGE



./1min

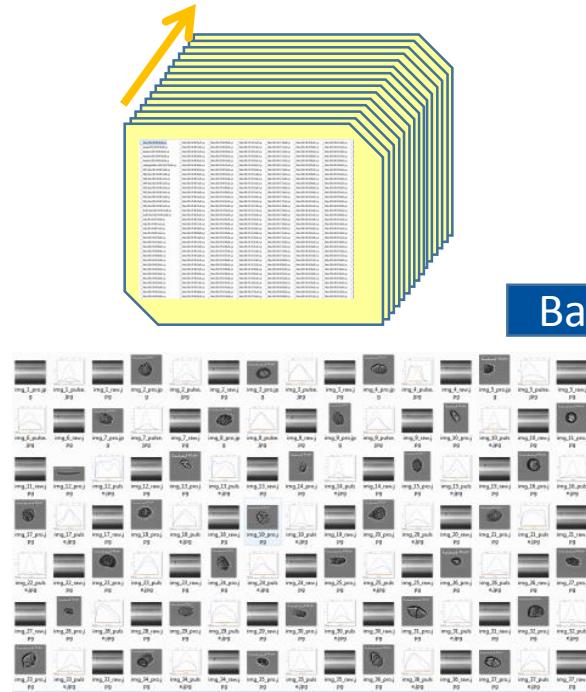
- Temperature
- Salinity
- Fluorescence
- Turbidity
- pH
- pCO₂
- Oxygen

Data acquisition & analysis



Output

Manual clustering



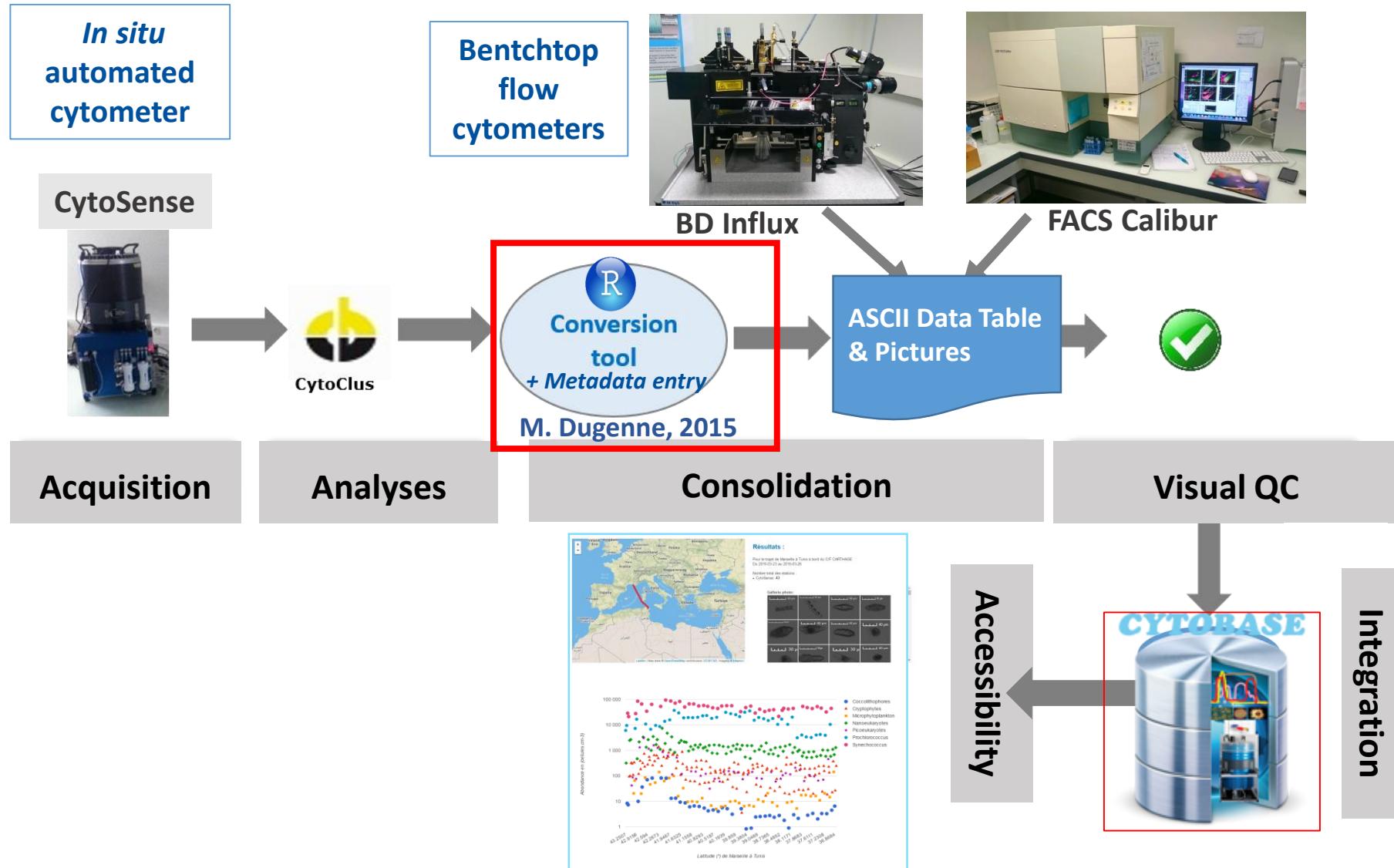
Measurements
by CytoSense



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

MIO FCM Data management

(working on interoperability with SeaDataNet)



→ Data Consolidation

Cytobase Input Processor (Mathilde Dugenne, 2015)

mathilde.dugenne@mio.osupytheas.fr



[Create Inputs folder](#)

Metadata

Data

Warning:

Upload successful

Project and samples context		Raw data	Size conversion																																																	
Project	Project date	PI	Cytometer ID																																																	
Enter project name	2015-08-18	Enter PI name	Enter cytometer ID																																																	
Station	Depth	Latitude	Longitude																																																	
Filename model																																																				
Filename	BERRE_082013_3F_FLR9 2013-12-17 13l																																																			
Samples operator	Standards reference	Clustering method	Observation type																																																	
Enter name of operator	Enter standards beads ref	Automated	In situ																																																	
<table border="1"> <tbody> <tr><td>2013-12-17T14:17:00Z</td><td>2013-12-17T14:17:00Z</td><td>16,056.60</td><td>Synechococcus</td><td>BERRE_082013_12S_FLR9 2013-12-17 14u17.cy</td><td>1.99</td><td>FL Red</td><td>10</td></tr> <tr><td>2013-12-17T14:17:00Z</td><td>2013-12-17T14:17:00Z</td><td>16,056.60</td><td>Cryptophytes</td><td>BERRE_082013_12S_FLR9 2013-12-17 14u17.cy</td><td>1.99</td><td>FL Red</td><td>10</td></tr> <tr><td>2013-12-17T14:43:00Z</td><td>2013-12-17T14:43:00Z</td><td>16,056.61</td><td>Beads 2 mu</td><td>BERRE_082013_16F_FLR9 2013-12-17 14u43.cy</td><td>2.04</td><td>FL Red</td><td>10</td></tr> <tr><td>2013-12-17T14:43:00Z</td><td>2013-12-17T14:43:00Z</td><td>16,056.61</td><td>Microphytoplankton</td><td>BERRE_082013_16F_FLR9 2013-12-17 14u43.cy</td><td>2.04</td><td>FL Red</td><td>10</td></tr> <tr><td>2013-12-17T14:43:00Z</td><td>2013-12-17T14:43:00Z</td><td>16,056.61</td><td>Picoplankton 2</td><td>BERRE_082013_16F_FLR9 2013-12-17 14u43.cy</td><td>2.04</td><td>FL Red</td><td>10</td></tr> <tr><td>2013-12-17T14:43:00Z</td><td>2013-12-17T14:43:00Z</td><td>16,056.61</td><td>Picoplankton 1</td><td>BERRE_082013_16F_FLR9 2013-12-17 14u43.cy</td><td>2.04</td><td>FL Red</td><td>10</td></tr> </tbody> </table>					2013-12-17T14:17:00Z	2013-12-17T14:17:00Z	16,056.60	Synechococcus	BERRE_082013_12S_FLR9 2013-12-17 14u17.cy	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.cy	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.cy	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.cy	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.cy	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.cy	2.04	FL Red	10
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Please associate each selection set to trigger, PMT's amplification and standardized phytoplankton category
NB: All incompatible entries will be removed

Expert name	Trigger	PMT's amplification	Standardized name
Cluster	Channel/Level	SWS	Cluster
Beads 2 mu	FL Red 10		Standard beads
		FLO	

→ Data Consolidation

Picture selection

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

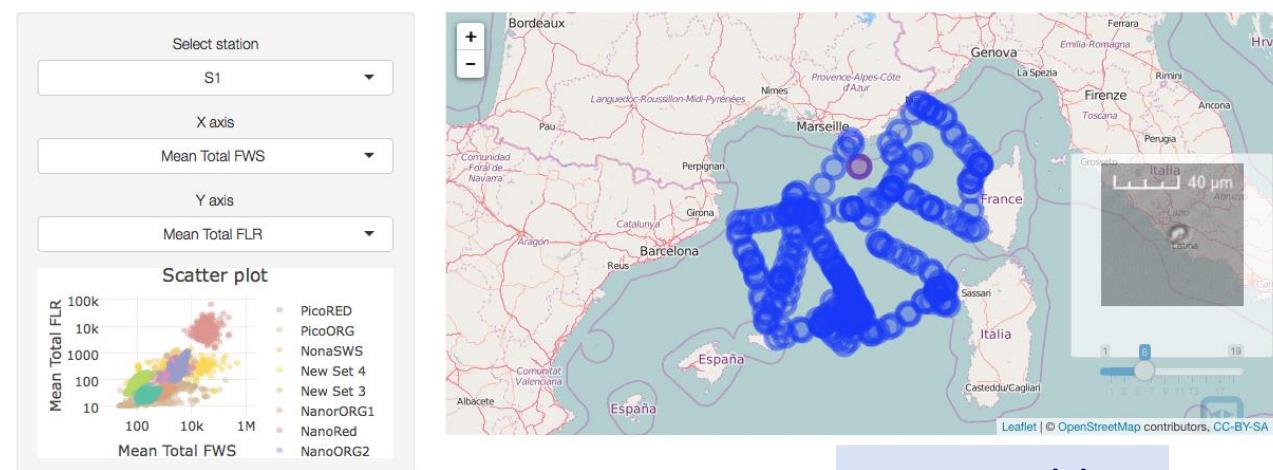
Add samples pictures
Select file
DEWEXL2FLR10 2013-04-05 15u04.cyz

Add pictures
Choisir les fichiers 15 fichiers Upload complete

Check all

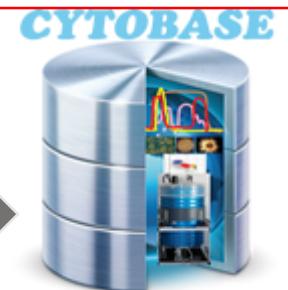
Show 4 entries Search:

Stations Explorer



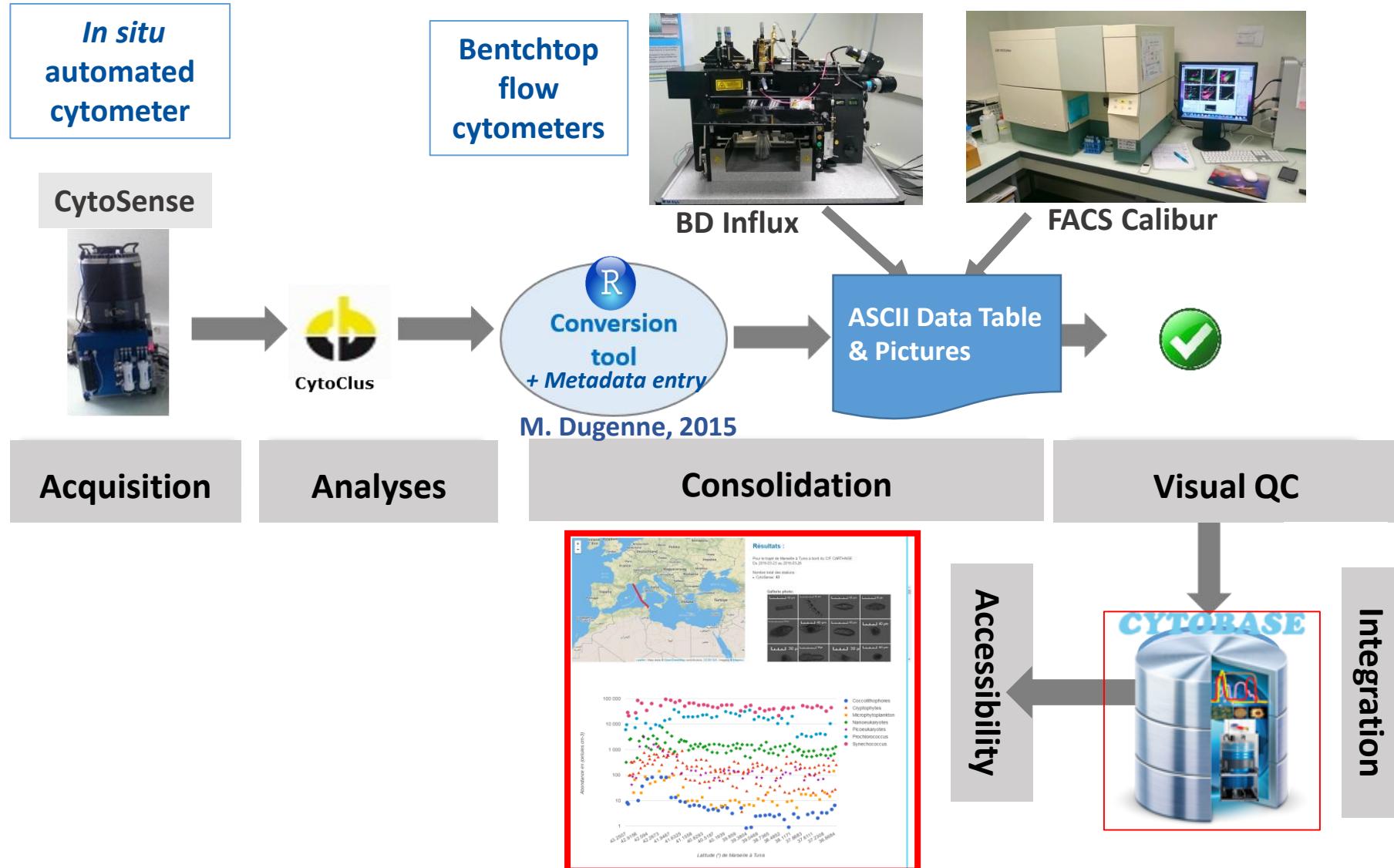
Download Table

Data Table
Picture Table



MIO FCM Data management

(working on interoperability with SeaDataNet)



Visualisation



FLOW CYTOMETRY DATABASE

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

CHROME

Continuous and High Resolution Observation of the Mediterranean Sea



Zone d'étude (Area of Study):
Mediterranean Sea - Western basin

Trajet (Transect):
MRS-TUN : du 24-03-2016 au 26-03-2016

Date de début (Start Date)
23/03/2016

Date de fin (End Date)
26/03/2016

Note: Please refer to the date mentioned in "Transect"

Instruments :

- CytoSense
- Photos cellulaires phytoplanctoniques
- FerryBox (Données pas disponibles)

Rechercher

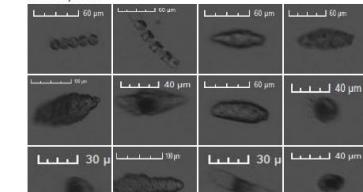


Résultats :

Pour le trajet de Marseille à Tunis à bord du C/F CARTHAGE
Du 2016-03-23 au 2016-03-26

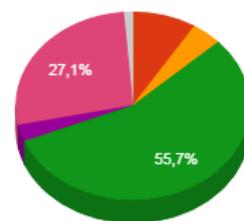
- Nombre total des stations :
- CytoSense : 43

Galerie photo:



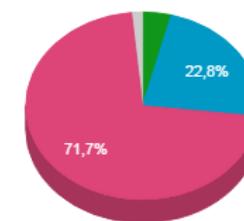
[Récupérer les données](#) [Imprimer](#)

Total Fluorescence Rouge (u.a.cm⁻³)



- Cryptophytes
- Microphytoplankton
- Nanoeukaryotes
- Picoeukaryotes
- Synechococcus
- Autres

Total Abundance (cell.cm⁻³)



- Nanoeukaryotes
- Prochlorococcus
- Synechococcus
- Autres



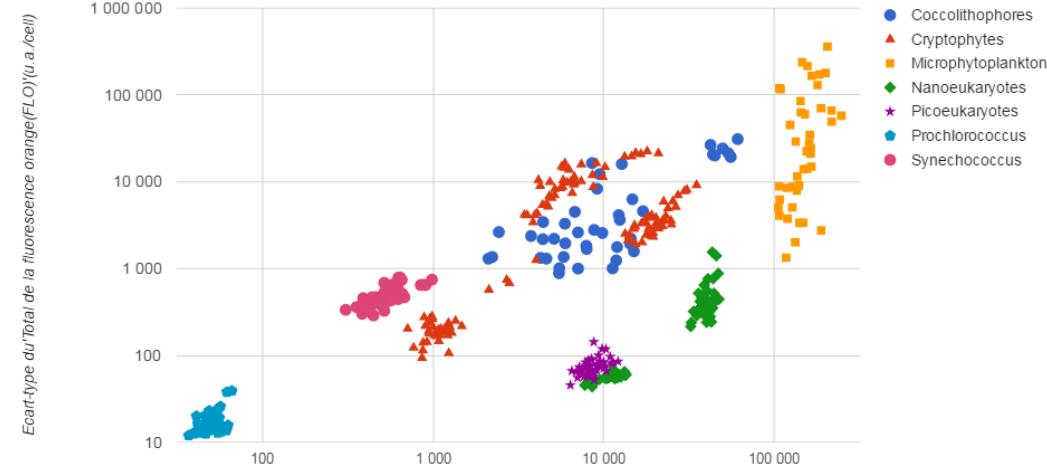
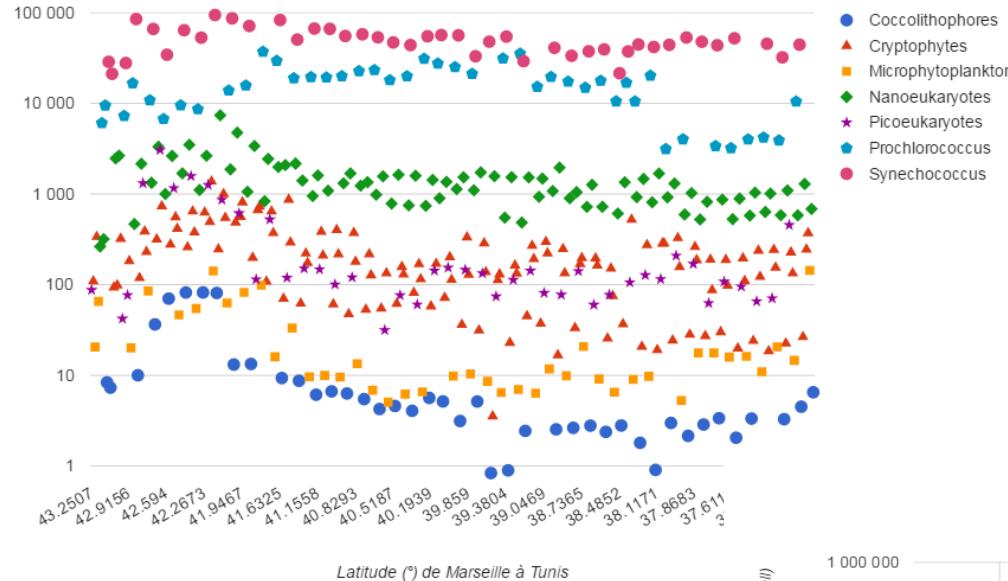
dygraphs



Visualisation



FLOW CYTOMETRY DATABASE



CYTOBASE data accessibility (Access within the MIO)

CYTOBASE

FLOW CYTOMETRY DATABASE



A world map showing data collection locations. The map includes zoom controls (+, -) in the top-left corner.

Zone d'étude: Mediterranean Sea - Western basin

Nom et date du Projet: DEWEX LEG2 : du 01-04-2013 au 30-04-2013

Date de début: 31/03/2013

Date de fin: 30/04/2013

Flow cytometer :

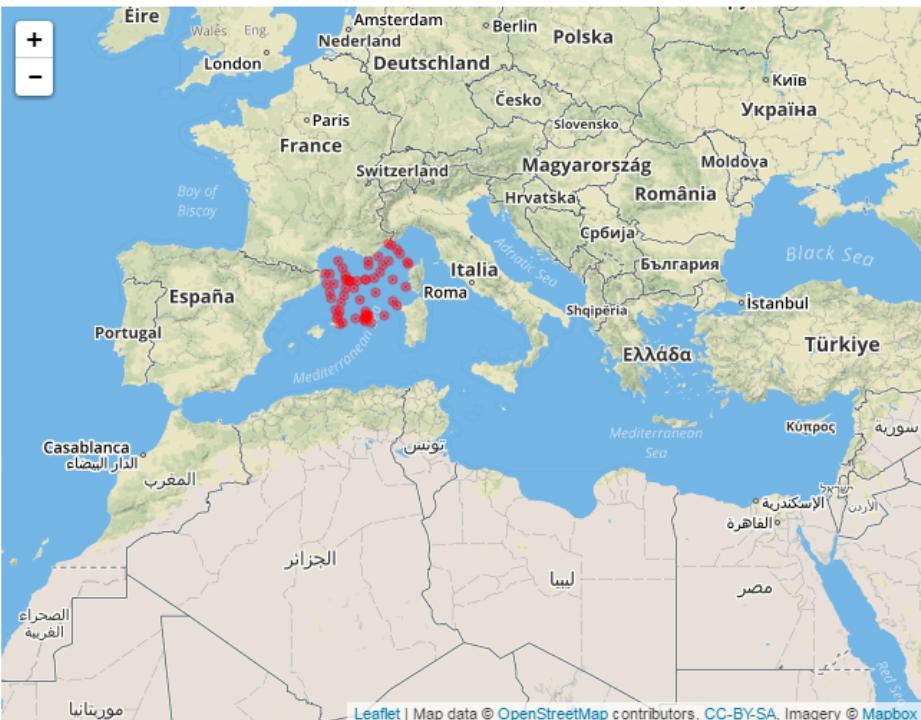
- CytoSense
- FACS Calibur
- BD Influx
- All instruments
- Phytoplankton groups
- Heterotrophic Bacteria

Search

<http://www.mio.univ-amu.fr/cytobase/>



FLOW CYTOMETRY DATABASE



Results :

For the period from 2013-03-31 to 2013-04-30

Project name DEWEX LEG2

Flow Cytometer : FACS Calibur

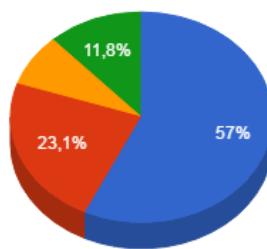
Station number : 59

No pictures

[Download Data](#)

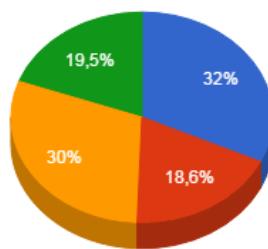
[Print](#)

Contribution to Green Fluorescence (u.a..cm⁻³)



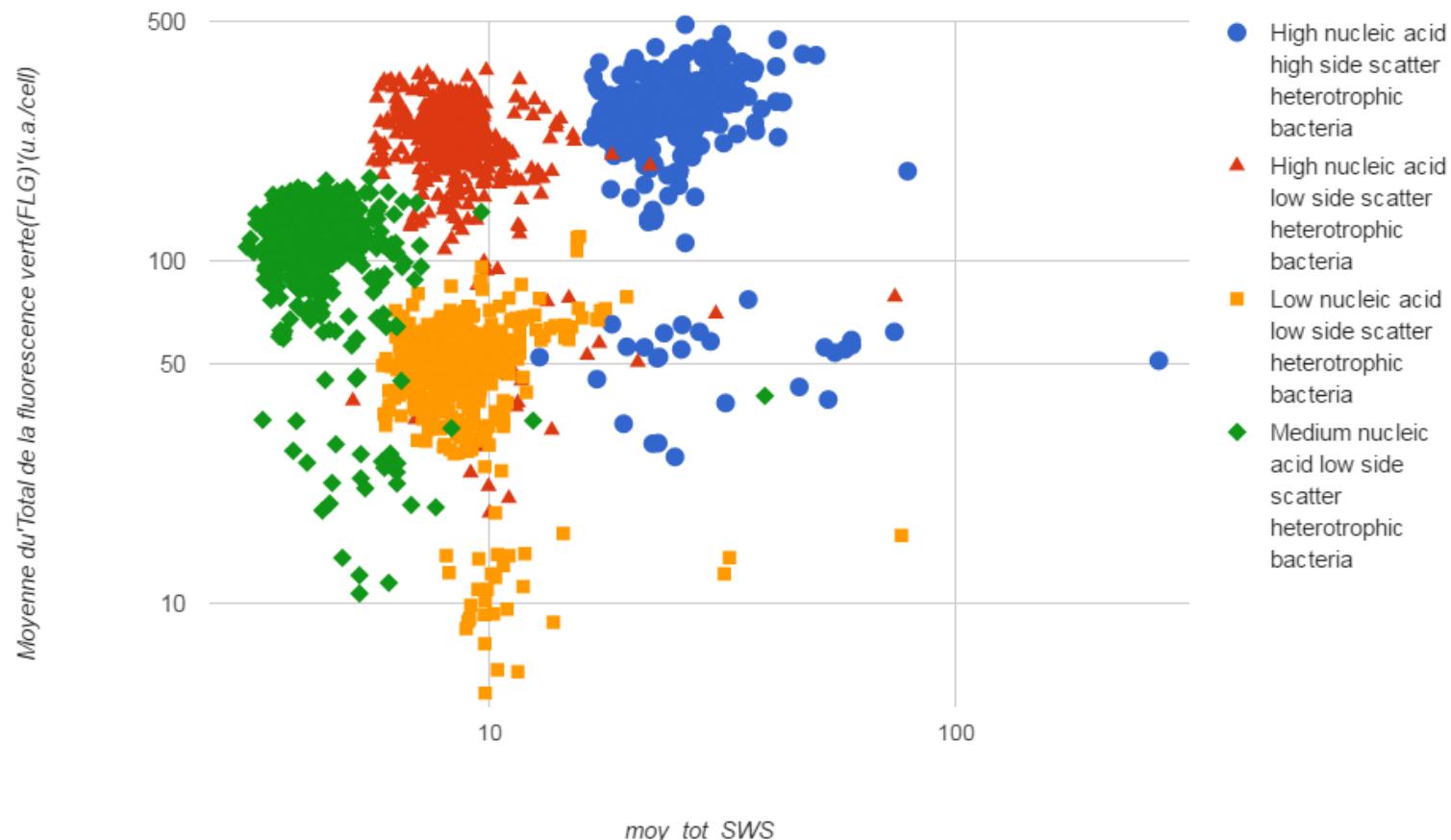
- High nucleic acid high side scatter heterotrophic bacteria
- High nucleic acid low side scatter heterotrophic bacteria
- Low nucleic acid low side scatter heterotrophic bacteria
- Medium nucleic acid low side scatter heterotrophic bacteria

Relative Abundance (cell.cm⁻³)



- High nucleic acid high side scatter heterotrophic bacteria
- High nucleic acid low side scatter heterotrophic bacteria
- Low nucleic acid low side scatter heterotrophic bacteria
- Medium nucleic acid low side scatter heterotrophic bacteria

Cytometric distribution parameters per group



Toward EU standardisation of FCM data



PI: Patrick FARCY (IFREMER)

Task 3.1: Automated platform
for the observation of
Phytoplankton diversity in
relation to ecosystem
services

Leader: Felipe ARTIGAS (CNRS-ULCO)



PI: Michèle FICHAUT (SISMER/IFREMER)



WP9.5.2: Ingesting, validating,
long-term storage and access of
Flow Cytometer data

Leaders: VLIZ, CNRS-MIO, NERC-BODC and ICES

Vocabulaire commun FCM

Toward EU standardisation of FCM data



PI: Patrick FARCY (IFREMER)

Task 3.1: Automated platform
for the observation of
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Leader: Felipe ARTIGAS (CNRS-ULCO)



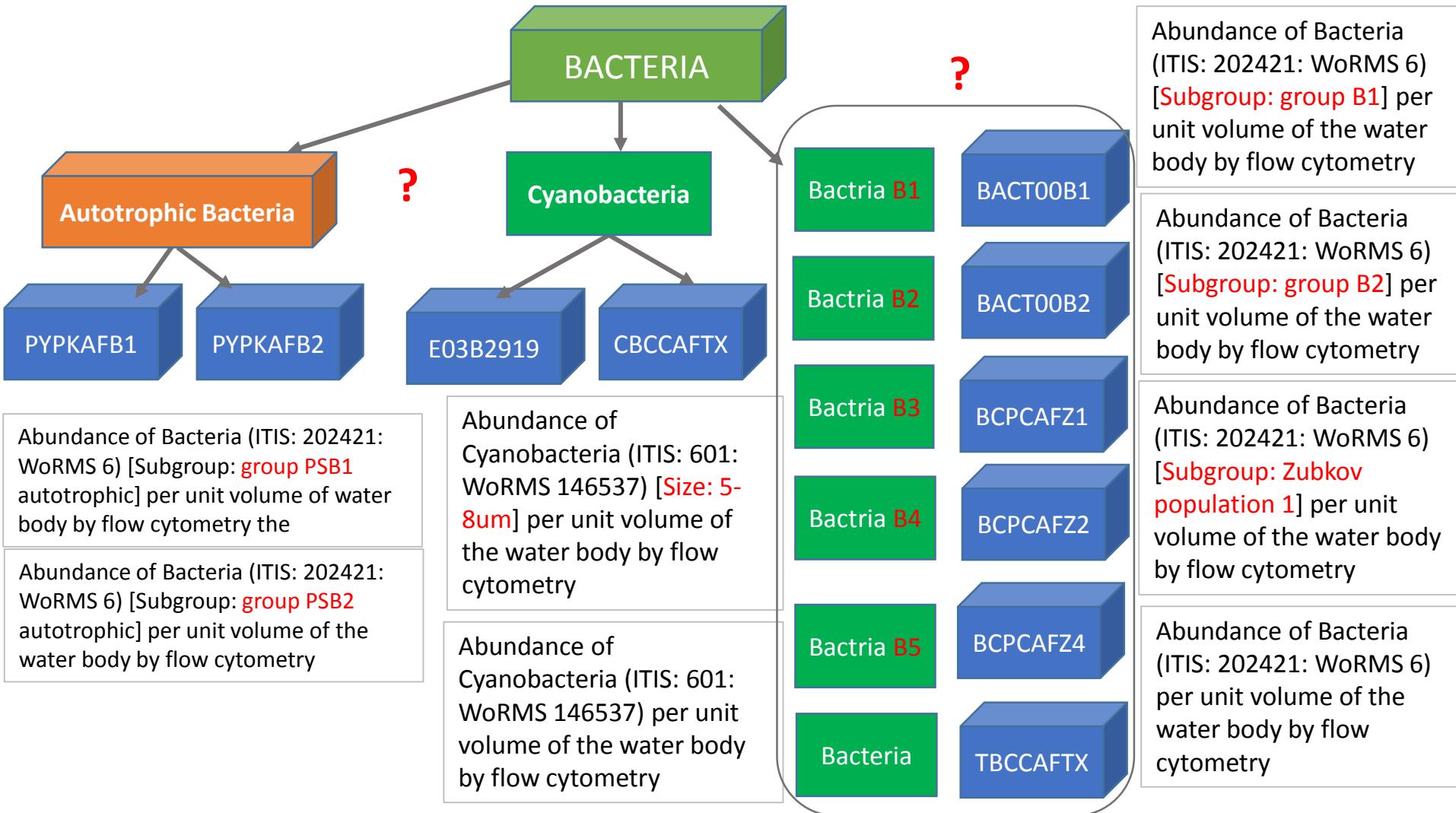
PI: Michèle FICHAUT (SISMER/IFREMER)

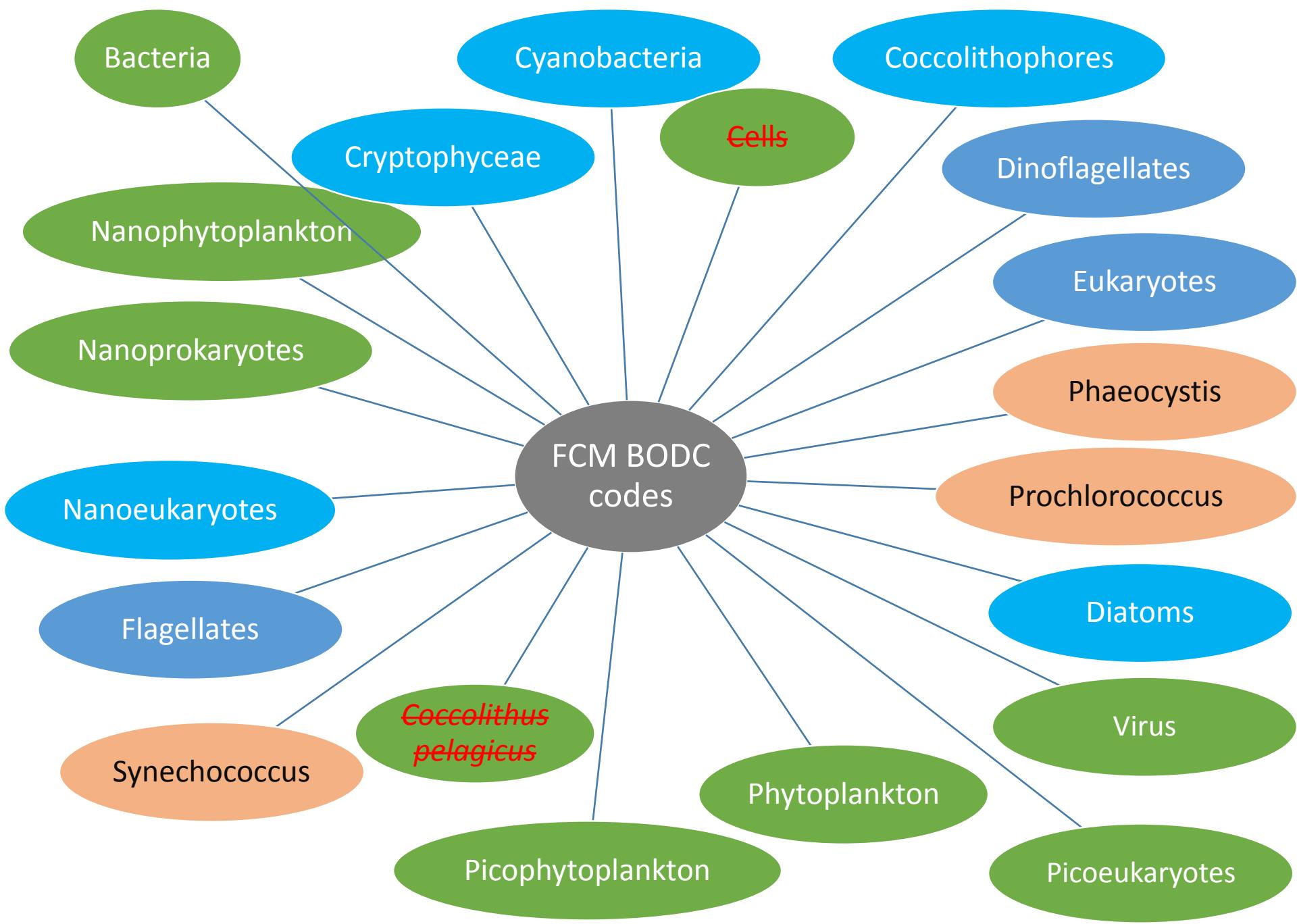


WP9.5.2: Ingesting, validating,
long-term storage and access of
Flow Cytometer data

Leaders: VLIZ, CNRS-MIO, NERC-BODC and ICES

Analyse critique des 34 codes existants au NERC-BODC





Etude bibliographique de 137 articles (1983 – 2017)

Titre	Créateur	Année
Flow cytometry and cell sorting: A technique for analysis and...	Yentsch et al.	1983
Marine phytoplankton distributions measured using shipboa...	Olson et al.	1985
Discrimination between types of pigments in marine <i>Syn...	Wood et al.	1985
Effects of Environmental Stresses on the Cell Cycle of Two M...	Olson et al.	1986
Light and dark control of the cell cycle in two marine phytopl...	Vaulot et al.	1986
Effect of light on the cell cycle of a marine <i>Synechococcus</i> st...	Armbrust et al.	1989
Kinetics of bacterial processes in natural aquatic systems bas...	Button et Robertson	1989
Using phytoplankton and flow cytometry to analyze grazing ...	Cucci et al.	1989
A flow cytometric approach to assessing the environmental a...	Demers et al.	1989
Rapid analytical technique for the assessment of cell metabol...	Dorsey et al.	1989
Optical plankton analyser: A flow cytometer for plankton ana...	Dubelaar et al.	1989
Use of a neural net computer system for analysis of flow cyto...	Frankel et al.	1989
Change in Photosynthetic Capacity over the Cell Cycle in Lig...	Gerath et Chisholm	1989
Photobiology of natural populations of zooxanthellae from t...	Lesser	1989
Discrimination of eukaryotic phytoplankton cell types from li...	Olson et al.	1989
Optical plankton analyser: A flow cytometer for plankton ana...	Peeters et al.	1989
Flow cytometry and phytoplankton	Phinney et Cucci	1989
Algorithm to estimate cell biovolume using image analyzed ...	Sieracki et al.	1989
Role of light and the cell cycle on the induction of spermato...	Armbrust et al.	1990
The Rapid Analysis of Single Marine Cells by Flow Cytometry ...	Burkhill et al.	1990
DNA polymorphism within the WH7803 serogroup of marine...	wood et Townsend	1990
Flow cytometric determination of phytoplankton DNA in cult...	Boucher et al.	1991
High-density photoautotrophic algal cultures: Design, constr...	Javanmardian et Palsson	1991
Survival of <i>Aeromonas salmonicida</i> in lake water.	Morgan et al.	1991
Phycoerythrins of Marine Unicellular Cyanobacteria. I.Bilin ty...	Ong et Glazer	1991
Growth and cell cycle of two closely related red tide-forming ...	Partensky et al.	1991
The picoplankton in Antarctic lakes of northern Victoria Land...	Andreoli et al.	1992
VERTEX: biological implications of total attenuation and chlo...	Broenkow et al.	1992
Interactions between marine bacteria and dissolved-phase an...	Button et al.	1992
Prochlorococcus marinus nov. gen. nov. sp.: an oxyphototro...	Chisholm et al.	1992
Analyzing multivariate flow cytometric data in aquatic scienc...	Demers et al.	1992
Evaluation of photosynthetic capacity in phytoplankton by fl...	Furuya et Li	1992
Quantifying heterogeneity: flow cytometry of bacterial cultur...	Kell et al.	1992

- 
1. Les groupes les plus identifiés
 2. Les définitions des groupes
 3. Les paramètres les plus mesurés
 4. Les capteurs les plus utilisés

Flow Cytometry vocabulary standardization Questionnaire

Flow Cytometry vocabulary standardization Questionnaire (FCM VSQ) is dedicated to identify the most common Flow Cytometry (FCM) Metadata and Data vocabulary used within the FCM community mainly involved in JERICO-NEXT (FP7 project) under the Task 3.1 on Automated platform for the observation of Phytoplankton diversity in relation to ecosystem services.

The aim of this questionnaire is to build a common standardized FCM vocabulary in order to better serve the marine research by understanding each other's data, sharing them and making them accessible through SeaDataNet which is a pan-European standardized infrastructure for managing a large and diverse marine data sets.

Part I: Flow Cytometer Metadata

What model of Flow Cytometer(s) do you use?

 (e.g.: CytoSense, FACS Calibur)

Does your instrument have an image in flow device?

Yes No

 (e.g.: pictures)

What is your sample inlet internal diameter (in micron)?

 (Separate multiple entries with commas)

Which lasers wavelengths do you use (in nanometer)?

 ((e.g.:488). In case of multiple entries, separate them with commas)

Laser beam powers (in mW)?

 ((e.g.:25). In case of multiple entries, please indicate the power of each of your laser separated by commas.)

Part II: Sample Metadata

• What Beads reference do you use ?

 (e.g.: brand, size, fluorescence, material)

• What beads diameters do you use ?

 (e.g.: 1, 2, 3, 6, 10 um, ...)

• What are your Beads Fluorescences ?

 (e.g.: Yellow)

• For what purpose do you use this instrument?

Check any that apply

- Research
- Monitoring
- Biotechnology
- Other:

• Where is your area of study?

Part III: Flow Cytometer Data

Do you use a fluorescent Dye?

Yes No

Which type of particles do you measure?

Check any that apply

- Phytoplankton
- Heterotrophic bacteria
- Virus
- Other:

What are the recurrent autotrophic functional groups of your area of study?

Check any that apply

- Synechococcus
- Prochlorococcus
- Picoeukaryotes
- Nanoeukaryotes
- Coccolithophore
- Cryptophytes
- Microphytoplankton
- Not concerned
- Other:

What are the recurrent Heterotrophic groups of your area of study?

Check any that apply

Part IV: Groups definition from the FCM point of view

Based on literature from 1983 to 2017, do you agree on these group definitions:

• Prochlorococcus

Prochlorococcus are defined as the smallest cyanobacteria found in marine environment. No staining is required to distinguish them by flow cytometry. FWS and FLR signatures are the smallest recorded up to now and require sensitive PMT or high powered lasers. The cluster, when well defined (often deep water communities) is below or may overlap that of Synechococcus group, and is often partially masked by the instrument background noise. In samples stained for Heterotrophic bacteria analysis, Prochlorococcus can be distinguished using Sideward Scatter (SWS) vs Chlorophyll Red Fluorescences (FLR) cytogram. They do not emit orange fluorescence because they lack phycoerythrin.

Check any that apply

- I agree
- I do not agree

• Synechococcus

They are unicellular photosynthetic Cyanobacteria with flow-cytometry forward-scatter (FWS) and sideward scatter (SWS) signatures that are larger than those of most of marine heterotrophic bacteria. No staining is required to distinguish them by flow cytometry. The related cluster has higher FWS and red fluorescence (FLR) signatures than Prochlorococcus and a distinct orange fluorescence (FLO) signature from their phycoerythrin accessory pigment when excited by lasers whose wavelength is below 533 nm. Cyanobacteria may contain phycocyanin, excited by a red laser and emitting above the chl a emission wavelength. The Synechococcus cluster is well resolved in red vs green (FLR/FLG) and in red vs orange fluorescences (FLR/FLO) cytograms. Due to their small size (0.8-1.2 μm) as reported in the literature, Synechococcus cells exhibit low intensity FWS, SWS and FLR signals.

Check any that apply

- I agree
- I do not agree

• Eukaryotes Picophytoplankton

The common definition of this group is size dependent; i/e <2-3 μm . No staining is required to distinguish them by flow cytometry. The smallest known eukaryotic picophytoplankton is Ostreococcus. Eukaryotic picophytoplankton exhibits a well defined flow cytometry signature, with FWS and FLR signals larger than that of Prochlorococcus and Synechococcus,

Merci pour votre intérêt

