



Data intensive research from genes to ecosystems in changing oceans

Damien Eveillard Stéphane Pesant & Guy Cochrane Tara Oceans Consortium

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Use Case: Tara Oceans Expedition 2009-2013



- Global study of Plankton
- 210 stations in 20 biogeo. provinces
- State of the art oceanographic equipment
- Consistent sampling methods





- State of the art plankton protocols
- across size spectra (0.02 µm 2 m)
- across taxonomic spectra (virus to jellyfish)

Context / Provenance is key to data integration



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Imaging data

- >4 million Images
- >30 terabytes
- State of the art automated methods adapted to organism from 0.1µm to 0.1m
- Semi-automatic recognition --Machine learning (supervised)
- Network approach for data sharing, expert annotation and training



Genomics data

- **40 million genes** --Largest-ever genomics effort in marine science
- **12 terabytes** -- equivalent to 135x human genomes
- >7000 data sets One of the richest molecular collection in the public domain



[De Vargas et. al. (2015)]

Cutting edge & innovative Science



- Plankton at **planetary scale**
- Eukaryotic plankton **diversity**
- Structure and function of the ocean **microbiome**
- Patterns and **ecological drivers** of ocean viruses
- Plankton community structure & "Interactome"
- Plankton transport across the Atlantic

nature Global Modelling - New approach to Biogeography

[Richter et. al., in revision]

Combining Lagrangian and genomic distances







New distribution theory for each size fraction



nature Global Modelling - New approach to Biogeochemistry [Guidi et. al. Nature 2016]

Predicting Carbon export based on Environmental, Imaging and Genomic Data







Key stone species as a community and marker genes

Metagenomic knowledge

Physical measurements

Few Species, or 51 genes predict quantitatively carbon export with 90% of accuracy

Toward a semi Automatic Computational Pipeline and design of Geochips

FUTURE PERSPECTIVE

Enabling grass-root innovation like Tara Oceans via existing European data infrastructures & services

Needs for a Blue Cloud