



# Demonstrator #2 Plankton Genomics





Blue-Cloud workshop 23rd of March 2021





#### **Plankton Genomics**

Genetic potential and diversity of plankton is unlocked by sequencing projects

However we are still only scratching the surface of microbial diversity in the Ocean



© Christian Sardet



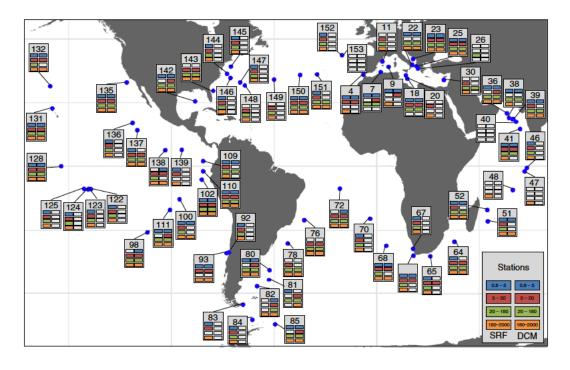
#### **Plankton Genomics**

No match in existing databases:

Marine Eukaryotes 50 %

(Carradec, Pelletier et al., 2018)

Common practice to discard any data that cannot be annotated through public databases



Tara Oceans: 68 different geographic locations across all the major oceanic provinces

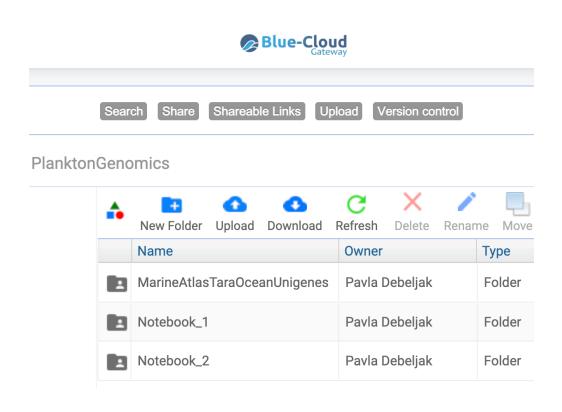


#### Plankton Genomics Demonstrator

#### Notebooks in the Blue-Cloud VRE

Notebook 1: Discovery of unknown biodiversity from unassigned genetic signals, co-occurence with known biodiversity, and environmental context

Notebook 2: Niche modelling and biogeography of as yet unknown biodiversity





#### **Plankton Genomics Datasets**

Exploration of the biggest
Eukaryotic plankton database
built from metagenomic and
metatranscriptomic sequences

Combining information on biogeographic distribution and providing data for global ocean modelling



#### Data available through



http://gs-service-production.geodab.eu/gs-service/search?view=elixir-ena



## **Workflow - Notebook 1: Discovery of Unknowns**

Annotated\_Taxa

Annotated\_Function

MetaG Genetic potential

MetaT
Gene expression

**Jupyter** Notebooks

- KNOWN
- UNKNOWN



Product Datasets
to be passed
on to **R** for
UNKNOWN/KNOWN
Analysis



Datasets published in the Blue-Cloud catalogue







NOTEBOOK 1



## **Workflow - Notebook 1: Discovery of Unknowns**

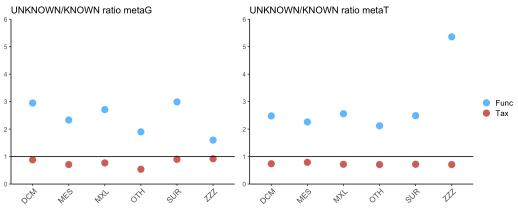
Annotated\_Taxa

Annotated\_Function

MetaG Genetic potential

MetaT
Gene expression

Ratio of UNKNOWN/KNOWN for MetaG and MetaT by depth layer



**NOTEBOOK 1** 



TARA

**OCEANS** 

# Workflow - Notebook 1: Discovery of Unknowns

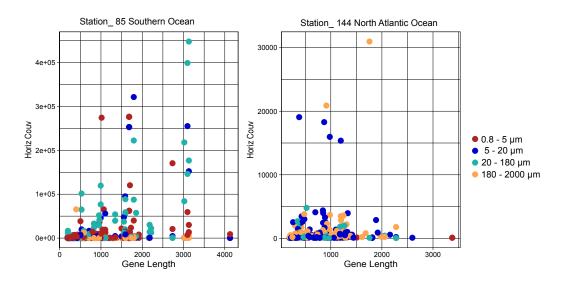
Annotated\_Taxa

Annotated\_Function

MetaG Genetic potential

MetaT Gene expression Horizontal coverage by sampling station and planktonic size class







**TARA** 

**OCEANS** 

# **Workflow - Notebook 1: Discovery of Unknowns**

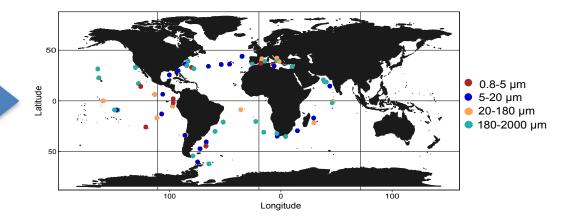
Annotated\_Taxa

Annotated\_Function

MetaG Genetic potential

MetaT
Gene expression

Distribution by oceanic region and planktonic size class

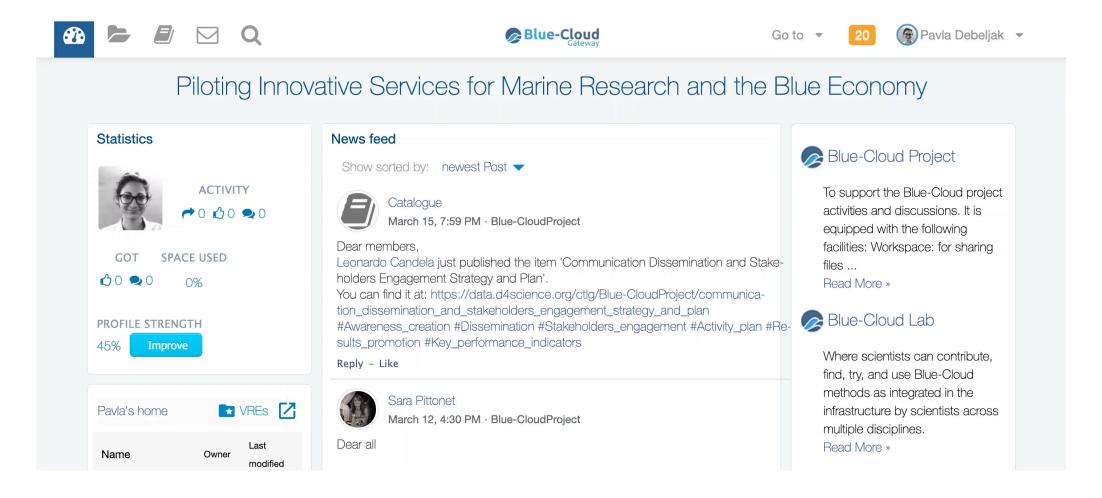


Published as catalogue in the VRE to be Passed on to Notebook 2 for niche modelling

NOTEBOOK 1



# Workflow - Notebook 1: Discovery of Unknowns





# Thank you for your attention!

# Demonstrator #2 Plankton Genomics

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