Summary of Available Tara Oceans Data

study title	data type	size fraction	study accession	reads_1e6	bases_1e9	runs	experiments	samples
Shotgun Sequencing of size fractions for prokaryotes and large DNA viruses	Metagenome	prokaryotes	ERP001736	24,764	4,892	248	248	135
Shotgun Sequencing of size fractions for prokaryotes and large DNA viruses	Metagenome	prokaryotes	ERP001737	8,972	1,801	85	85	75
Shotgun Sequencing of size fractions for protist	Metagenome	protists	ERP003628	23,518	4,660	243	243	159
Shotgun Sequencing of size fractions for DNA viruses	Metagenome	DNA viruses	ERP003708	4,489	863	115	114	92
Metatranscriptome sequencing of size fractions for protists.	Metatranscriptome	protists	ERP006152	284	52	27	27	22
Metatranscriptome sequencing of size fractions for protists.	Metatranscriptome	protists	ERP006156	1	0	29	29	24
MAST4 Single Cell Whole Genome Amplification	Single-cell		ERP003711	125	12	4	4	2
16S Sequencing of size fractions for prokaryotes	16S	prokaryotes	ERP003634	0	0	6	6	6
Amplicon sequencing of size fractions for large DNA viruses.	16S	DNA viruses	ERP006153	2	1	35	35	19
Amplicon sequencing of size fractions for protists.	16S	protists	ERP006157	993	293	370	370	334
Amplicon sequencing of size fractions for protists.	16S	protists	ERP007024	24	6	20	20	14

"This project will rely on samples collected during the scientific expedition Tara-Oceans (2009-2012). By March 2012, the schooner Tara, equipped with innovative systems for sampling of 11 organismal size-ranges covering entire planktonic communities from viruses to animals, has collected standardized genetic (total DNA/RNA), morphological, and physico-chemical (contextual) samples from 153 sites across the world oceans, locations carefully selected with input from near-real-time remote sensing and in-situ hydrographic criteria. Overall, a total of ~50,000 biological samples and ~13,000 contextual measures from 3 depths will be analysed. The metagenomics component of the project consists of size fractionated plankton samples that are submitted to barcoding and shotgun sequencing, as well as isolated single-cell amplified protists and single-organisms isolated metazoans that are sequenced as reference genomes."

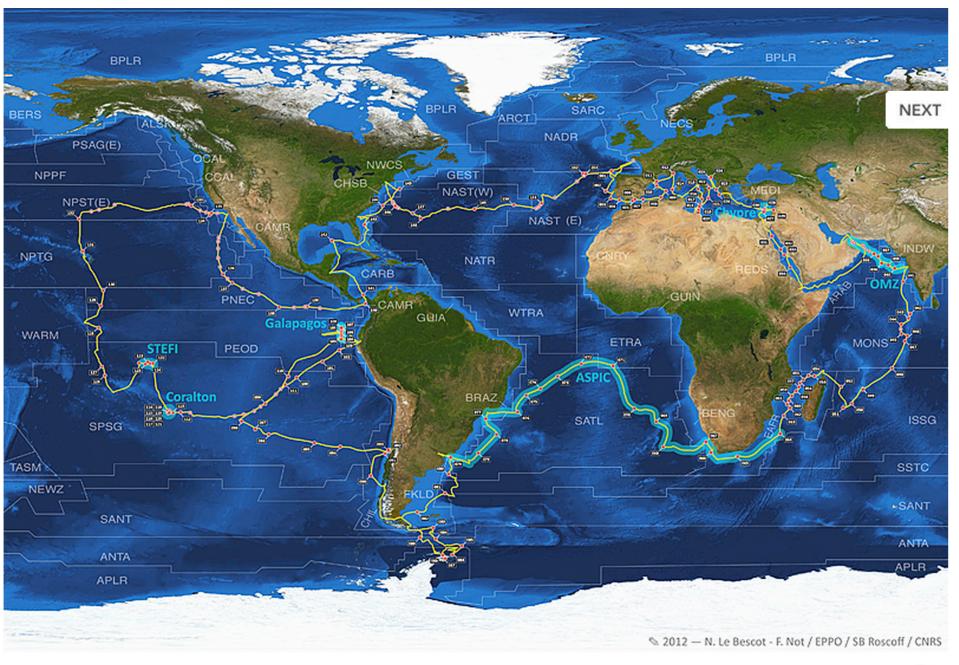
Data release policy

(http://www.embl.de/tara-oceans/start/facts figures/Tara-Oceans-Data-Release-Policy.pdf)

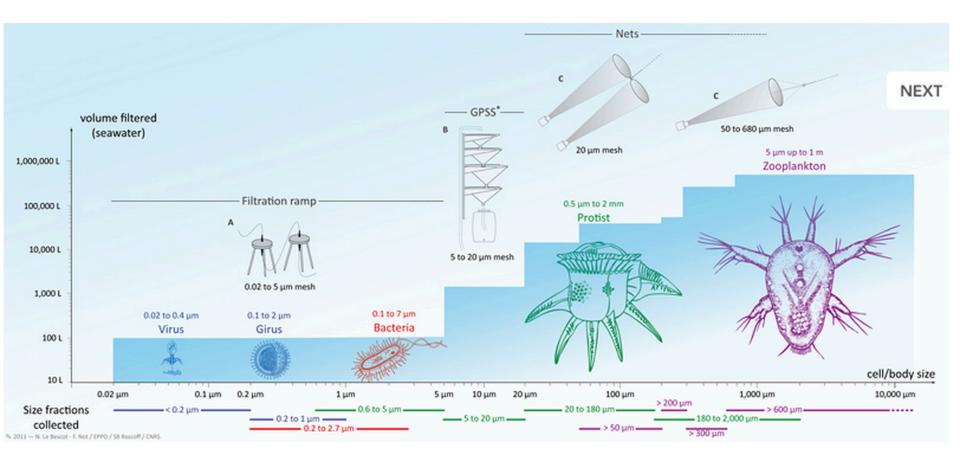
The Tara Oceans Consortium has established a policy for pre-publication data usage with the aim to make quickly and broadly available the global data generated during the project for advancing science, while maintaining a priority for analyses by the laboratories that participate in the data production effort. During one year after release in a public database, the data will be considered usable with the following restriction: specifically, intended analyses may not compete with those reserved for the Tara Oceans Consortium, including global analyses at the genome and community level, and comparisons between samples. Therefore, before starting any study on these reserved data sets, investigators are required to contact the Tara Oceans Consortium to obtain an agreement.

This restriction will be lifted after the publication of analyses on the considered data sets by the Tara Oceans Consortium, or automatically one year after the release of the data. If a publication is made from the released data sets, it is expected that the Tara Oceans Consortium is cited in the acknowledgment section. Raw data sets have the same limits for use than the advanced data sets derived from them. Download of data from the Tara Oceans project implies a full acceptance of the policy usage described in this paragraph.

- All runs in the SRA were updated at some point in 2014
- Run dates range from 3/19/2010 to 11/20/2013 (one in 2014)







TARA OCEANS organisms

Image 1 of 4



http://www.embl.de/tara-oceans/start/publications/index.html

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Acinas, S. G. et al. Validation of a new catalysed reporter deposition-fluorescence in situ hybridization probe for the accurate quantification of marine Bacteroidetes populations. Environ Microbiol, doi:10.1111/1462-2920.12517 (2014).

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Environmental data stored at Pangaea

http://www.pangaea.de/search?env=All&q=project%3Alabel%3ATara+Oceans

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Always quote citation when using data

More than 10000 datasets found on search for »project:label:Tara Oceans«

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1. Tara Oceans Expedition, P; Tara Oceans Consortium, C (2014): Registry of selected samples from the Tara Oceans Expedition (2009-2013)

Related to: Brum, J; Tara Oceans Consortium, C; et al.: Global patterns and ecological drivers of ocean viral communities. Science

de Vargas, C; Tara Oceans Consortium, C; et al.: Sea change in eukaryotic plankton diversity. Science

Lima Mendez, G; Tara Oceans Consortium, C; et al.: Top-down determinants of community structure in the global plankton interactome. Science

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Size: 10784 data points

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2. Morozov, NP (1968): Lithium and rubidium concentrations in waters of seas and oceans

Supplement to: Morozov, NP (1968): Geochemistry of rare alkaline elements in the oceans and seas. Oceanology

Size: 4 datasets

doi:10.1594/PANGAEA.753510 - Score: 71% - Similar datasets

3. Johnson, DA; Nigrini, CA (1985): Neogene radiolarian datum levels in the equatorial Indian and Pacific Oceans

Supplement to: Johnson, DA; Nigrini, CA (1985): Synchronous and time-transgressive Neogene radiolarian datum levels in the Equatorial Indian and Pacific Oceans. Marine Micropaleontology

Size: 9 datasets

doi:10.1594/PANGAEA.690483 - Score: 58% - Similar datasets

4. Schmiedl, G; Mackensen, A (1993): Observational data around large miliolid foraminifera from the polar oceans

Supplement to: Schmiedl, G; Mackensen, A (1993): Cornuspiroides striolatus (BRADY) and C. rotundus nov. spec.: Large miliolid foraminifera from Arctic and Antarctic Oceans. Journal of Foraminiferal

Research

Size: 2 datasets

doi:10.1594/PANGAEA.715971 - Score: 58% - Similar datasets

Questions

- Data embargos?
- Which studies?
- What's already been done?
- Analysis plan?
- Data QC?