

The KAUST Global Ocean Genome workshop provides an opportunity for scientists and students to discuss progress in the assessment of the Global Ocean Genome. The main goal of this conference is to bring together internationally recognized scientists who will introduce their research and students who will discuss the progress in the assessment of the Global Ocean Genome, including:

- Updating the current inventory of genes in the microbial communities of the Global Ocean and assessing the progress made towards providing an adequate representation of microbial communities and environments in the Global Ocean
- Exploring the functional capacities of the global ocean microbiome as represented by the Global Ocean Genome
- Examining the global distribution and sequence variability of key functional genes
- Developing new analytical approaches to mine the Global Ocean Genome for genes of industrial interest

Sunday 29 October level 0 Auditorium between Buildings 2 and 3

8:30 -8:40	Welcome address Pierre Magistretti, Dean, Biological and Environmental Sciences and Engineering Division, KAUST
8:40 -8:45	House keeping notes and review of workshop goals Workshop Chair, Carlos M. Duarte, RSRC & CBRC, KAUST
8:45 - 12:15	Session 1. Exploration of the global ocean genome Chair Chris Bowler, ENS, Paris
8:45- 9:15	Keynote lecture: Disclosing the deep ocean microbiome Silvia Acinas, ICM, CSIC, Spain
9:15 - 9:35	Power law distribution of marine OTUs: Implications for the richness of ocean microbes Victor M. Eguiluz, KAUST & IFISC, Spain
9:35 - 9:55	The Tara Oceans microbial reference gene catalog Guillem Salazar, ETHZ, Switzerland
9:55 - 10:15	Mesopelagic gene and microbial species enrichment across the global tropical and subtropical ocean Pablo Sanchez, ICM, CSIC, Spain
10:15 -10:45	Coffee break
10:45 - 11:15	Keynote lecture: Comparative marine metagenomics in the global ocean project Takashi Gojobori, BESE, KAUST

11:15 - 11:35	Sequencing effort dictates gene discovery in ocean microbial metagenomes Carlos M. Duarte, RSRC & CBRC, KAUST
11:35 - 11:55	High-throughput sequencing the global ocean eukaryotic plankton: promises and challenges Naiara Rodriguez, AZTI, Spain
11:55 - 12:15	Evidence of different community structuring patterns in eukaryotic vs. prokaryotic picoplankton inhabiting the sunlit global ocean Ramiro Logares, ICM, CSIC, Spain
12:15 - 13:45	Lunch (campus diner, building 13)
13:45	Session 1 (cont). Exploration of the global ocean genome
13:45 - 14:05	Curation of samples from the microbial group of MALASPINA (harmonisation of metadata, QA/QC and submission to EBI/ENA). Stephane Pesant, Marum, Germany
14:05 - 14:25	Metagenomic analyses using Dragon Metagenomic Analysis Platform (DMAP) Intikhab Alam, CBRC, KAUST
14:25 - 14:45	Seasonal dynamics of the Red Sea metagenomics Katsuhiko Mineta, CBRC, KAUST

14:45 - 15:15	Coffee break
15:15	Session 2. Metatranscriptomics Chair Silvia Acinas, ICM, CSIC, Spain
15:15- 15:45	Keynote Lecture: Exploration of Tara Oceans metatranscriptomics data Chris Bowler, ENS, Paris
15:45 - 16:05	Changes on a Synechococcus population during a mesocosm experiment on the Red Sea Alexia Coello, RSRC, KAUST
16:05 - 16:25	Widespread distribution of salvage pathways in oceanic microbes: implications for autotrophs David Ngugi, RSRC, KAUST
16:25 - 16:45	EMOSE (2017) inter-comparison of marine plankton 'Omics methods (filtration, sequencing and analysis pipelines) Stephane Pesant, Marum, Germany
16:45 - 17:05	Elucidating the genome of a deep sea fungus Ramiro Logares, ICM, CSIC, Spain
19:30	END OF SESSION Workshop dinner: Al Marsa Restaurant, Yacht Club

Monday 30 October level 0 Auditorium between Buildings 2 and 3

9:00	Session 3. The ocean virome Takashi Gojobori, BESE, KAUST
9:00 – 9:30	Keynote Lecture: vignettes of virus variation in the oceans Curtis Suttle, UBC, Canada
9:30 - 9:50	Ecogenomics of deep sea viral communities Elena Lara, Istituto di Scienze Marine, Venice, Italy (presented by Carlos Duarte)
9:50 - 10:10	RNA viruses: The underestimated entities in the marine environment Marli Vlok, UBC, Canada
10:10 - 10:30	Out of the blue: atmospheric inputs to the global ocean genome Nojood Aalismail, RSRC, KAUST
10:30 - 11:00	Coffee break
10:30 - 11:00 11:00	Coffee break Session 4. Applications of the Open ocean genome
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11:00	Session 4. Applications of the Open ocean genome Keynote Lecture: Global trends in the use of marine genetic resources
11:00 11:00 – 11:30	Session 4. Applications of the open ocean genome Keynote Lecture: Global trends in the use of marine genetic resources Txetxu Arrieta, IEO, Spain Global diversity of marine animals based on metagenomes

13:30	Session 5. Top, & bottom boundaries and computational issues Session Chair, Vlad Bajic, Director CBRC, KAUST
13:30 - 14:00	Keynote lecture: Microbial community assembly and evolution in subseafloor sediment Kasper Kjeldsen, Aarhus University, Denmark
14:00 - 14:20	Computational resources at KAUST Saber Feki, KSL, KAUST and Greg Wickham, ACI, KAUST
14:20 – 14:40	Multi-omics approach to assess viral impact on marine plankton Hiroyuki Ogata, Kyoto, Japan
14:40 - 15:10	The extreme gene and metabolic process repertoires of prokaryotes inhabiting the deep sea hypersaline anoxic basins. Daniele Daffonchio, BESE, KAUST
15:10 - 15:40	Coffee break
15:40 – 16:00	Diazotroph dynamics in tropical and subtropical oceans Roslinda Mohammed, RSRC, KAUST
16:00 - 16:40	Proposal for working groups (Oct 31st - Nov 1st)
16:40 - 17:40	 Parallel demonstrations Single Cell Sorting and Genomics Platform Takashi Gojobori, BESE, KAUST
	DMAP Hands on Demonstration Intikhab Alam, CBRC, KAUST
	 Tour to the BIO Core Lab Facilities Abdallah M. Abdallah, Bioscience Core Lab, KAUST
19:30	END OF SESSION Dinner at Thuwal Traditional Fish Restaurant

(this is outside KAUST, so you need to bring the KAUST

ID card, passport, and appropriate clothing).

Tuesday 31 October

8:45 - 13:00 Informal conversation snorkeling trip to Red Sea coral reefs (breakfast and lunch provided - only for registered participants)

13:00 – 14:00 Prepare for writing sessions

14:00 - 18:00 Writing session 1

Rooms available: 3318, campus library sea view (24 seats) 2286, building 2, open area (12 seats) 2290, building 2, open area (16 seats)

Writing Groups

- 1. Resources and infrastructure for advancing the exploration of the Global Ocean genome (what are the bottle necks, how can the community organize themselves to overcome these?)
- Best practices in the assemblage and analyses of Ocean Metagenome Data (we have been exploring how different choices impact on the outcome, and these are quite considerable, other metagenomic communities, eg. Human micro biome, are developing benchmarks and standards to test procedures, which is something we seem to be lacking).
- 3. Advancing the Exploration of the Global Ocean Genome (where do we stand?, what are the key research questions, challenges and opportunities?).

END OF SESSION

19:30 Dinner: PURE restaurant (Island Recreational Club)

Wednesday 1 November

9:00 – 10:00	Report to the plenary on progress by writing groups
10:00 – 12:30	Writing Session 2 Rooms available: 3318, campus library sea view (24 seats) 2286, building 2, open area (12 seats) 2290, building 2, open area (16 seats)
12:30 - 13:30	Lunch (Campus diner, building 13)
13:30 – 16:00	Writing Session 3 Rooms available: 3318, campus library sea view (24 seats) 2286, building 2, open area (12 seats) 2290, building 2, open area (16 seats)
16:00 – 17:00	Report to the plenary on progress by writing groups
17:00 – 17:30	Action items and closing Carlos M. Duarte, KAUST, workshop chair

Participants

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