Fluorescent in situ hybridization data from B/O Hermano Gines cruises for the CARIACO Basin Time Series Station from May to November 2014 (CariacoMetaOmics project)

Website: https://www.bco-dmo.org/dataset/652438
Data Type: Cruise Results
Version: Final
Version Date: 2016-07-22

Project
» Genetic and Metabolic Signatures of Marine Microorganisms in Oxygen Depleted and Varying Geochemical Seascapes (CariacoMetaOmics)

Contributors	Affiliation	Role
Edgcomb, Virginia P.	Woods Hole Oceanographic Institution (WHOI)	Principal Investigator
Taylor, Gordon T.	Stony Brook University - SoMAS (SUNY-SB SoMAS)	Principal Investigator
Taylor, Craig	Woods Hole Oceanographic Institution (WHOI)	Co-Principal Investigator
Ake, Hannah	Woods Hole Oceanographic Institution (WHOI BCO-DMO)	BCO-DMO Data Manager

Table of Contents

Dataset Description

Fluorescent in situ hybridization (FISH) data enumerated from microscopic samples on 2 legs of each of the CAR212 and CAR216 cruises.

Methods & Sampling

All samples were collected via Niskin bottles.

Seawater samples were fixed with 2% borate-buffered formaldehyde, frozen overnight, then filtered the next day onto 0.2-um polycarbonate membrane filters. Filters were refrozen at -20 degrees celsius until hybridization. All FISH hybridization procedures were performed according to Pernthaler et al. (2001) with fluorescent probes labeled with CY3 or fluorescein. Filters were counterstained with DAPI and probe-positive cells were enumerated as a percentage of the total community (DAPI-positive). FISH probes were:

Probe	Target Group	References
Alf968	α-*proteobacteria*	Neef 1997
Gam42a	γ-*proteobacteria*	Manz et al. 1992
Eps682	ε-*proteobacteria*	Lin et al. 2008

Data Processing Description

All data were normalized to volume of seawater.

DMO Notes:

-transferred FISH data section into separate a data object.
-reformatted the date to comply with BCO-DMO standards.
-reformatted some column names to comply with BCO-DMO standards.

[table of contents | back to top]

Data Files

File
Fluor_insitu_Hybrid.csv (Comma Separated Values (.csv), 3.80 KB) MD5:9d7fb7ee76bc2ab58ce6531bcc4abdd7 Primary data file for dataset ID 652438

[table of contents | back to top]

Parameters

Parameter	Description	Units
cruise_id	cariaco cruise number and leg number	unitless
date_start	start date; mm/dd/yyyy	unitless
depth	depth	meters
delta495	percent Delta495 cells of total	dimensionless
delta495_sd	standard deviation of the percent Delta495 cells of total	dimensionless
gam42a	percent Gam42a cells of total	dimensionless
gam42a_sd	standard deviation of the percent Gam42a cells of total	dimensionless
eps682	percent Eps682 cells of total	dimensionless
eps682_sd	standard deviation of the percent Eps682 cells of total	dimensionless
alf968	percent Alf968 cells of total	dimensionless
alf968_sd	standard deviation of the percent Alf968 cells of total	dimensionless

[table of contents | back to top]

Instruments

Dataset-specific Instrument Name	Niskin bottle
Generic Instrument Name	Niskin bottle
Dataset-specific Description	All samples were collected via Niskin bottles.
Generic Instrument Description	A Niskin bottle (a next generation water sampler based on the Nansen bottle) is a cylindrical, non-metallic water collection device with stoppers at both ends. The bottles can be attached individually on a hydrowire or deployed in 12, 24, or 36 bottle Rosette systems mounted on a frame and combined with a CTD. Niskin bottles are used to collect discrete water samples for a range of measurements including pigments, nutrients, plankton, etc.

[table of contents | back to top]

Deployments

CAR212_2

Website	https://www.bco-dmo.org/deployment/652493
Platform	B/O Hermano Gines
Start Date	2014-05-07
End Date	2014-05-09
Description	These deployments are part of the MetaOmics studies in the Cariaco Basin

CAR212_3

Website	https://www.bco-dmo.org/deployment/652571
Platform	B/O Hermano Gines
Start Date	2014-05-10
End Date	2014-05-12
Description	part of the MetaOmics Cariaco Basin Project

CAR216_2

Website	https://www.bco-dmo.org/deployment/652494
Platform	B/O Hermano Gines
Start Date	2014-11-05
End Date	2014-11-07
Description	These deployments are part of the MetaOmics studies in the Cariaco Basin.

CAR216_3

Website	https://www.bco-dmo.org/deployment/652574
Platform	B/O Hermano Gines
Start Date	2014-11-08
End Date	2014-11-10
Description	part of MetaOmics Project in the Cariaco Basin

[table of contents | back to top]

Project Information

Genetic and Metabolic Signatures of Marine Microorganisms in Oxygen Depleted and Varying Geochemical Seascapes (CariacoMetaOmics)

Coverage: Southern Caribbean Sea - 10" 30' N, 64" 40' W (CARIACO Ocean Time Series Station)

Oxygen depleted water columns (ODWCs) appear to be expanding in response to global climate change. This alters trophic structure, compresses habitat and modifies geochemical cycles of major elements. Oxygen depletion can vary in intensity and duration from seasonal hypoxia to permanent anoxia. The focus of this study is a classic example of the anoxic end-member, the Cariaco Basin. The overall goal is to examine how microbial functional potential (metagenomic), activity (metatranscriptomic), taxonomic diversity (based on SSU rRNA) and the ecological/geochemical consequences (in terms of measured rates of key processes) relate along vertical oxygen/geochemical gradients and between seasons in the Cariaco Basin. This will reveal relationships between expression of particular sets of genes, environmental differences in nutrients, energy substrates and oxidant availabilities.

The objectives are to: (1) Integrate hydrographic, geochemical and microbial ecological data with metagenomic and metatranscriptomic profiles to understand regulatory and metabolic networks defining microbial community responses to environmental forcing during high and low productivity periods. This will help to understand the importance of processes, such as anaerobic oxidation of methane, utilization of redox-sensitive metals, the cryptic sulfur cycle in this ODWC, and the impacts of oxygen depletion on nitrogen transformations. (2) Determine the importance of associations between microbial eukaryotes (mEuks) and prokaryotes in this ODWC. (3) Identify "indicator" genes of known or unknown function that may be relevant to major elemental and trace gas cycling as targets for further biochemical characterization and molecular probe development, and quantify a key subset of these genes and transcripts across redox gradients using qPCR. (4) Provide a basis for developing monitoring tools using expressed genes indicative of important elemental transformations and fluxes for diagnosing the health status of natural and human engineered ecosystems. (5) Compare results with recent and ongoing studies of other ODWCs to discern shared and unique attributes of these systems.

Intellectual Merit: Previous studies of ODWCs have underscored the need for more data on microbial community structure and functionality in ODWCs, particularly biochemical rate measurements and other data on community responses to changing conditions. Better predictive models of responses of marine microbial communities and biogeochemical processes to global climate change are essential for informing future policy and management decisions. Data from an anoxic end-member ODWC like Cariaco Basin are critically needed to compare with data from other recent and ongoing studies of seasonally-depleted coastal systems and permanently-depleted deep basin and western boundary oxygen minimum zones (OMZs) to construct more skillful models. This study will advance the understanding of impacts of expanding ODWCs around the world, moving beyond assessments based only on taxonomic diversity, to yield new insights into the ecology and physiology of major microbial groups in these environments and interactions among Bacteria, Archaea and microbial eukaryotes.

Broader Impacts: The PIs and their collaborators will train one Research Associate, one postdoctoral investigator, a graduate student, and numerous undergraduates from SBU. All personnel will be trained in various aspects of microbial ecology and oceanography, with an emphasis on both traditional (e.g., microscopy) and "cutting edge" (e.g. metagenomics/transcriptomics) techniques. The PIs will also involve the Zephyr Education Foundation's marine science literacy and education program, located in Woods Hole, MA.

The PIs will work with this organization to educate inner city K-12 students using local boat field trips organized by Zephyr, and lectures, and classroom laboratory exercises designed by the PIs. Additionally, this project will have broad implications for understanding how ODWCs affect marine ecosystems, and may influence future management strategies and models describing the cycling of C and N between the ocean and atmosphere.

[table of contents | back to top]

Funding

Funding Source	Award
NSF Division of Ocean Sciences (NSF OCE)	OCE-1335436
NSF Division of Ocean Sciences (NSF OCE)	OCE-1336082

[table of contents | back to top]