

Cruise track from R/V Atlantic Explorer AE1012 in the BATS site from May 2010 (Ocean Microbial Observatory project)

Website: <https://www.bco-dmo.org/dataset/543646>

Data Type: Cruise Results

Version: 1

Version Date: 2014-12-22

Project

» [Transitions in the Surface Layer and the Role of Vertically Stratified Microbial Communities in the Carbon Cycle - An Oceanic Microbial Observatory](#) (Ocean Microbial Observatory)

Contributors	Affiliation	Role
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Abstract

Cruise track from R/V Atlantic Explorer AE1012 in the BATS site from May 2010.

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Coverage

Spatial Extent: N:-63.586302 E:32.504323 S:-64.7248 W:31.233993

Temporal Extent: 2010-05-12 - 2010-05-16

Dataset Description

Cruise track from R/V Atlantic Explorer AE1012 in the BATS site from May 2010.

Methods & Sampling

Control point navigation was obtained from the R2R catalog: <http://www.rvdata.us/catalog/AE1012>

Data Processing Description

BCO-DMO Processing Notes

- Data obtained from R2R site
- Added parameter header

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Data Files

File
cruise_track_AE1012.csv (Comma Separated Values (.csv), 2.20 KB) MD5:07f285519f18c9927fb4fbd55a0f82b
Primary data file for dataset ID 543646

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Parameters

Parameter	Description	Units
lat	Latitude component of geographic position; north is positive.	Decimal degrees
lon	Longitude component of geographic position; east is positive.	Decimal degrees
ISO_DateTime_UTC	Date/Time (UTC) ISO formatted. This standard is based on ISO 8601:2004(E).	YYYY-MM-DDTHH:MM:SS[.xx]Z

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Instruments

Dataset-specific Instrument Name	GPS
Generic Instrument Name	Global Positioning System Receiver
Generic Instrument Description	The Global Positioning System (GPS) is a U.S. space-based radionavigation system that provides reliable positioning, navigation, and timing services to civilian users on a continuous worldwide basis. The U.S. Air Force develops, maintains, and operates the space and control segments of the NAVSTAR GPS transmitter system. Ships use a variety of receivers (e.g. Trimble and Ashtech) to interpret the GPS signal and determine accurate latitude and longitude.

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Deployments

AE1012

Website	https://www.bco-dmo.org/deployment/543344
Platform	R/V Atlantic Explorer
Start Date	2010-05-12
End Date	2010-05-16

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Project Information

Transitions in the Surface Layer and the Role of Vertically Stratified Microbial Communities in the Carbon Cycle - An Oceanic Microbial Observatory (Ocean Microbial Observatory)

Website: <http://www.bios.edu/research/projects/oceanic-microbial-observatory/>

Coverage: Bermuda Atlantic Time-Series study site

(Adapted from the NSF award abstract)

The premise of this project is that stratified bacterioplankton clades engage in specialized biogeochemical activities that can be identified by integrated oceanographic and microbiological approaches. Specifically, the objective of this project is to assess if the mesopelagic microbial community rely on diagenetically altered organic matter and subcellular fragments that are produced by microbial processes in the euphotic zone and delivered into the upper mesopelagic by sinking or mixing. In past efforts this microbial observatory had greater success cultivating members of the euphotic zone microbial community, and revealed an unanticipated growth requirement for reduced sulfur compounds in alphaproteobacteria of the SAR11 clade. Genomic information showed that intense competition for substrates imposes trade-offs on bacterioplankton - there are regions of N dimensional nutrient space where specialists win. We postulate that specific growth requirements may explain some the regular spatial and temporal patterns that have been observed in upper mesopelagic bacterioplankton communities, and the difficulties of culturing some of these organisms.

The specific objectives of this project are: 1) to produce ^{13}C and ^{15}N labeled subcellular (e.g., soluble, cell wall, and membrane) and DOM fractions from photosynthetic plankton cultures and use stable isotope probing to identify specific clades in the surface and upper mesopelagic microbial community that assimilate fractions of varying composition and lability. 2) to use fluorescence in situ hybridization approaches to monitor temporal and spatial variability of specific microbial populations identified from the SIP and HTC experiments. To increase resolution we will use CARD-FISH protocols. 3) to measure the proteomes of bacterioplankton communities to identify highly translated genes in the surface layer and upper mesopelagic, and community responses to seasonal nutrient limitation. 4) and, to cultivate these organisms via high throughput culturing (HTC) by pursuing the hypothesis that they require specific nutrient factors and/or diagenetically altered organic substrates. Complete genome sequences from key organisms will be sought and used as queries to study patterns of natural variation in genes and populations that have been associated with biogeochemically important functions.

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Funding

Funding Source	Award
NSF Division of Ocean Sciences (NSF OCE)	OCE-0802004

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