

One-minute navigation points from R/V Cape Hatteras CH0912 in the Blake Ridge, Cape Fear Diapir from 2012-2012 (SEEP project)

Website: <https://www.bco-dmo.org/dataset/537481>

Version: 2014-11-05

Project

» [Connectivity in western Atlantic seep populations: Oceanographic and life-history processes underlying genetic structure](#) (SEEP)

Contributors	Affiliation	Role
Van Dover, Cindy	Duke University	Chief Scientist
Ball, Bernard	Duke University (Duke - Bio)	Technician
Copley, Nancy	Woods Hole Oceanographic Institution (WHOI BCO-DMO)	BCO-DMO Data Manager

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Dataset Description

Retrieved from R2R: <http://www.rvdata.us/catalog/CH0912>

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Data Files

File
CH0912_track.csv (Comma Separated Values (.csv), 144.98 KB) MD5:f39053a22ee308e9d89a4dcc86e2bdf5
Primary data file for dataset ID 537481

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Parameters

Parameter	Description	Units
ISO_DateTime.UTC	UTC time and date	YYYY-MM-DDTHH:MM:SS[.xx]Z
lon	longitude; east is positive	decimal degrees
lat	latitude; north is positive	decimal degrees
sog	Instantaneous speed-over-ground	meters/second
cog	Instantaneous course-over-ground	degrees clockwise from north

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.

Deployments

CH0912

Website	https://www.bco-dmo.org/deployment/521433
Platform	R/V Cape Hatteras
Report	http://dmoserv3.who.edu/data_docs/SEEPC/CH0912_cruise_report.pdf
Start Date	2012-11-01
End Date	2012-11-03
Description	SEEPC project cruise. Cruise information and original data are available from the NSF R2R data catalog.

Project Information

Connectivity in western Atlantic seep populations: Oceanographic and life-history processes underlying genetic structure (SEEPC)

Coverage: Western Atlantic, Gulf of Mexico, Intra-American Sea

This project will evaluate connectivity on spatial scales that match those at which vent systems are being studied (3500 km), with a set of nested seeps (within the Barbados system) within which connectivity can be explored at more local spatial scales (30 to 130 km), and with species that span depth (600 m to 3600 m) and geographic ranges (30 km to 3500 km) and that have diverse life-history characteristics. Five deep-sea seep systems in the Intra- American Sea (IAS) are targeted: Blake Ridge, Florida Escarpment, Alaminos Canyon, Brine Pool, Barbados (El Pilar, Orenoque A, Orenoque B). The primary objective is to advance our general knowledge of connectivity in the deep sea. The focus is on species and processes occurring in the IAS, with attention to oceanographic circulation, life histories, and genetics. Questions that apply in shallow-water systems motivate this study:

1. What phylogeographic breaks occur in the system? It is important to distinguish between phylogeographic history and connectivity. A phylogeographic break with no shared alleles between populations implies a long

history of isolation or possibly cryptic speciation.

2. Are populations connected by ongoing migration? This is the fundamental question about connectivity and the scale of genetic variation in marine species with planktonic larvae.

3. What biophysical processes underlie observed connectivities? Biological processes (e.g., larval distributions in the water column, timing of reproduction, and planktonic larval duration) and physical processes of transport and dispersion interact to determine connectivity.

The oceanographic model for the IAS will be improved and coupled to a Lagrangian larval transport model. The field program includes time-series sampling of larvae at seeps with records of current velocities, water column sampling to determine larval distribution potential, shipboard studies of larval biology and behavior, and sampling of benthic target species. Phylogenetic and population genetic tools will be used to explore historical and contemporary gene flow. Iterative interactions among the science teams will advance our understanding of connectivity in the deep sea and to develop effective and best methods for hypothesis testing under the constraints of working in a relatively inaccessible environment. Since their discovery, deep-sea chemosynthetic ecosystems have been novel systems within which to test the generality of paradigms developed for shallow-water species. This study will explore scale-dependent biodiversity and recruitment dynamics in deep-sea seep communities, and will identify key factors underlying population persistence and maintenance of biodiversity in these patchy systems.

[Google Earth map](#) showing positions of stations, CTD, XBT, multibeam locations (KMZ file download)

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Funding

Funding Source	Award
NSF Division of Ocean Sciences (NSF OCE)	OCE-1031050

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