

CTD data from casts before and after larval vertical distribution sampling from R/V C-Hawk day cruises in the Eastern Gulf of Maine from 2012 to 2014

Website: <https://www.bco-dmo.org/dataset/783736>

Data Type: Cruise Results

Version: 1

Version Date: 2019-12-30

Project

» [Intertidal community assembly and dynamics: Integrating broad-scale regional variation in environmental forcing and benthic-pelagic coupling](#) (GOMEPRO)

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Abstract

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Coverage

Spatial Extent: N:44.6609108 E:-67.3390943 S:44.411259 W:-67.6526958

Temporal Extent: 2012-08-01 - 2014-08-07

Dataset Description

These data were presented in Weinstock et al., 2018 (see Fig 1 and 2). Only downcast density data were analyzed and presented in Weinstock et al., 2018. Water density data were converted to % of water column maximum in Figure 2 of Weinstock et al., 2018.

These data are available in two formats. The version in the BCO-DMO data system contains data from all casts concatenated together with added columns `cast_name`, `Cast_time_UTC`, `start_latitude`, `start_longitude`, `end_latitude`, and `end_longitude` which was originally contained in comment and header lines in 35 cast csv files. The 35 individual cast csv files are available in the "Data Files" section as CTD.zip: "CTD csv files with seabird headers."

Related Datasets: CTD casts were conducted immediately before and after the associated larval vertical distribution sampling (separate dataset description) and on both flood and ebb tides.

* CTD cast log for mussel study: <https://www.bco-dmo.org/dataset/783749>

* Mussel Larvae Vertical Distribution: <https://www.bco-dmo.org/dataset/783755>

Methods & Sampling

Conductivity, temperature, and depth were sampled via an instrument cast from a small boat. Casts were conducted immediately before and after the associated larval vertical distribution sampling (see Mussel Larvae Vertical Distribution <https://www.bco-dmo.org/dataset/783755>) and on both flood and ebb tides.

See Weinstock et al., 2018, for more methodology details.

Problem report:

The second cast data for the flood tide on 7/31/13 are missing.

Data Processing Description

BCO-DMO Data Manager Processing Notes:

- * 35 csv files for each cast combined into one tabular dataset. "cast_name" column added which contains information in the original filenames which included "mm_dd_yy_tidephase_cast.csv"
- * The 35 original csv files submitted to BCO-DMO were bundled into CTD.zip and is available to download in the "Data Files" section as "CTD csv files with seabird headers." The only change made to these files was to remove blank rows that were in-between every data and header row in a subset of the 35 files.
- * added a conventional header with dataset name, PI name, version date
- * modified parameter names to conform with BCO-DMO naming conventions (spaces, +, and - changed to underscores). Units in parentheses removed and added to Parameter Description metadata section.
- * blank values in this dataset are displayed as "nd" for "no data." nd is the default missing data identifier in the BCO-DMO system.
- * Cast timestamp and start/end latitude and longitudes added to this dataset from parsed information contained in comment and header lines in the originally submitted cast csv files. Other information parsed from these comment lines was turned into a second tabular dataset "CTD cast log" (<https://www.bco-dmo.org/dataset/783749>).
- * Data sorted by Cast_time_UTC, Time, Depth columns

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Data Files

File	
CTD csv files with seabird headers filename: CTD.zip	(ZIP Archive (ZIP), 323.58 KB) MD5:c093e4db408e3e153bdcbfd1e1e31e65
Data are structured in a separate pair of files for each sampling date/tidal phase combination, with file names formatted mm_dd_yy_tidephase_cast. The first cast was conducted immediately before the larval vertical distribution sampling described in the accompanying dataset file, and the second cast was conducted immediately after larval sampling. Coordinates, start and end times, and other pertinent metadata are included in the header of each file. An additional descriptor stating the site name is included in the files from 8/16/12, when two different sites were sampled on the same date.	
The CTD.zip file contains 35 .csv files.	
ctd_larvae.csv	(Comma Separated Values (.csv), 499.60 KB) MD5:223a6b7d3ece231b5670486f55ccfac9
Primary data file for dataset ID 783736	

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Related Publications

Weinstock, J. B., Morello, S. L., Conlon, L. M., Xue, H., & Yund, P. O. (2018). Tidal shifts in the vertical distribution of bivalve larvae: Vertical advection vs. active behavior. *Limnology and Oceanography*, 63(6), 2681–

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Parameters

Parameter	Description	Units
cast_name	Cast identifier. Contains cast date (in format mm_dd_yy), tidal phase (e.g. Ebb) and site identifier (e.g. MB2) where more than one cast was conducted during o mm_dd_yy_tidephase_cast	unitless
Pressure	Pressure	Decibars (dbar)
Depth	Depth	Meters (m)
Density	Density	Kilograms per Cubic Meter (kg/m ³)
Conductivity	Conductivity	MicroSiemens per Centimeter (uS/cm)
Temperature	Temperature	Celsius (C)
Specific_conductance	Specific conductance	MicroSiemens per Centimeter (uS/cm)
Salinity	Salinity	Practical Salinity Scale
Sound_velocity	Sound velocity	Meters per Second (m/s)
Time	Seconds elapsed from the start of the cast.	seconds (s)
Cast_time_UTC	Cast date and time (UTC) in ISO 8601 format yyyy-mm-ddTHH:MM:SSZ	unitless
Start_latitude	Cast start latitude	Decimal degrees
Start_longitude	Cast start longitude	Decimal degrees
End_latitude	Cast end latitude	Decimal degrees
End_longitude	Cast end longitude	Decimal degrees

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Deployments

Yund_GoME

Website	https://www.bco-dmo.org/deployment/699509
Platform	R/V C-HAWK
Start Date	2012-08-01
End Date	2016-07-29
Description	<p>The C-Hawk is a 22 ft. fiberglass modified V-hull. These were multiple single-day deployments for GOMEPRO project. Eastern Gulf of Maine Sampled with single-day cruises on: 8/1/12 8/16/12 8/22/12 7/31/13 7/22/14 8/5/14 8/6/14 8/7/14</p> <p>Methods & Sampling Date Site 8/1/12 MB1 8/16/12 MB1 & MB2 8/22/12 WB1 7/31/13 WB2 7/22/14 WB3 8/5/14 WB3 8/6/14 WB3 8/7/14 WB3</p>

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Project Information

Intertidal community assembly and dynamics: Integrating broad-scale regional variation in environmental forcing and benthic-pelagic coupling (GOMEPRO)

Coverage: Rocky intertidal shores and nearshore coastal waters throughout the Gulf of Maine

Rocky intertidal habitats in the Gulf of Maine (GoM) provide a model system to examine the structure and dynamics of natural communities. Throughout the Gulf of Maine, the same species are often found in these habitats but community structure, dynamics and productivity differ markedly among 3 distinct regions (southern, central and northern GoM). Past influential work, conducted primarily in the southern and central GoM, focused on the local processes driving intertidal community structure but produced very different conceptual models of how these communities are structured. This project examines whether regional differences in rocky shore community processes are driven by differences in recruitment that are shaped by regional variation in temperature and food availability and nearshore coastal oceanography. This project will improve the understanding of how large-scale environmental forces interact with local processes to control the distribution of species and the structure and dynamics of these communities. Understanding the interaction between processes operating at different scales is fundamentally important to developing more reliable models that can be used to predict community dynamics. In addition, data resulting from this project will have important implications for regional dynamics in commercially important species and for ecosystem and fisheries management within the GoM.

The overarching hypothesis of this project is that regional differences in community-level processes are driven by very different patterns of population connectivity and recruitment in a few key species, and that these differences are ultimately caused by regional variation in temperature and food availability and mediated by physical larval transport processes. Hence, the project will test the following hypotheses with manipulative field experiments, field sampling, connectivity estimates, and integrative modeling:

- 1) Locally-dispersing species dominate dynamics in regions with a net export of planktonic larvae (Northern GoM), while species with planktonic larvae dominate the dynamics in regions with high settlement and extensive connectivity among populations (Southern GoM).
- 2) Settlement density of species with planktonic larvae increases from northern to southern regions in accord with regional variation in food availability.
- 3) Population connectivity varies greatly among regions, with regions differing in the degree to which they are self-seeded or serve as larval sources vs. sinks; self-seeding leads to relatively localized population dynamics in the middle portion of the GoM.
- 4) Patterns of population connectivity are driven by physical transport processes and can be represented by coupling basic larval behavior models with circulation models.

At 18 different sites in the GoM across ~ 600 km, surveys will evaluate variation in recruitment, food availability and secondary productivity and experiments will assess community processes in wave-exposed and sheltered habitats. We will use hydrographic, current profile, and larval vertical distribution surveys to collect data for coupled larval/circulation models. Population connectivity will be both modeled and empirically evaluated (for

one species) using elemental fingerprinting. A spatially explicit metacommunity model will integrate across all project components and test the relative importance of regional and local processes in controlling community organization and dynamics.

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Funding

Funding Source	Award
NSF Division of Ocean Sciences (NSF OCE)	OCE-1458188

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