CTD data collected on R/V Lowell Weicker in the Long Island Sound in 2008

Website: https://www.bco-dmo.org/dataset/3656 Version: 31 May 2012 Version Date: 2012-05-31

Project

» <u>Testing hypotheses about diversity, gene flow, and effective population size in marine planktonic ciliates</u> (CiliateDivGenePop)

| Contributors | Affiliation | Role |
|------------------------|---|---------------------------|
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Dataset Description

CTD data are provided for 6 stations sampled in Long Island Sound on 13 August 2008. Data include temperature, density, salinity, fluorescence, and oxygen saturation along a depth gradient.

Methods & Sampling

Sample stations were selected to represent a range of variation in depth, salinity, and vertical stratification. At each station, measurements were made with a SeaBird CTD for the purpose of creating density profiles.

Stations and methods are further described in the following publication: Tamura, M, LA Katz, and GB McManus. 2011. Distribution and diversity of Oligotrich and Choreotrich ciliates in a large temperate estuary. Aquat. Microb. Ecol. 64:51-67. doi:10.3354/ame01509 (<u>PDF</u>)

Data Processing Description

Data were processed with the SeaSoft program from SeaBird and smoothed by binning at 0.1 m intervals. BCO-DMO made the following modifications to the dataset: format of parameter names was changed to conform to BCO-DMO conventions; blanks and values of '-9.99E-29' were replaced with 'nd'; lat and lon values were converted from degrees/decimal minutes to decimal degrees. Note: lat and lon values are approximate for Station 2.

Data Files

File

CTD_profiles.csv(Comma Separated Values (.csv), 58.70 KB) MD5:0caabe47f4b3042d8483018adcea31d1

Primary data file for dataset ID 3656

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Parameters

| Parameter | Description | Units |
|------------------|--|-----------------------------|
| cruiseid | Unique identifier for the cruise. | dimensionless |
| year | Year in YYYY format. | dimensionless |
| month_gmt | Month in which the samples were collected, in MM format (GMT). | dimensionless |
| day_gmt | Day on which the samples were collected, in DD format (GMT). | dimensionless |
| station | Numeric identifier of the sample station. | dimensionless |
| lat | Latitude in decimal degrees. Positive values indicate North. Values have been converted from degrees and decimal minutes to decimal degrees. | decimal degrees |
| lon | Longitude in decimal degrees. Negative values indicate West. Values have been converted from degrees and decimal minutes to decimal degrees. | decimal degrees |
| time_start_local | Local time at which the CTD cast began, in HHMM format. (Local time = GMT - 5 hours). | dimensionless |
| depth | Sample depth in meters. | meters |
| density | Density, measured in kg per cubic meter. | kg/m^3 |
| temp | Water temperature in Celsius degrees. | degrees Celsius |
| sal | Salinity. | PSU |
| fluor | Fluorescence measured in mg per cubic meter. | mg/m^3 |
| O2_mg | Oxygen saturation measured in mg/liter. | mg/L |
| time_elapsed | Time elapsed (in seconds) since the first CTD sample at the station. | seconds |
| day_local | Day on which the samples were collected, in DD format (local time). | dimensionless |
| month_local | Month in which the samples were collected, in MM format (local time). | dimensionless |
| time_start_gmt | GMT time at which the CTD cast began, in HHMM format. | dimensionless |
| ISO_DateTime_UTC | Date and time formatted to ISO8601 standard. T indicates start of time string; Z indicates UTC. | YYYY-MM- DDTHH:MM:SS.ssZ |

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Instruments

| Dataset- specific Instrument Name | CTD Sea-Bird SEACAT 19 |
|--|---|
| Generic Instrument Name | CTD Sea-Bird SEACAT 19 |
| Dataset- specific Description | A SeaBird SBE 19 CTD was used at each station to record temperature, salinity, and other physical data. |
| Generic Instrument Description | The Sea-Bird SBE 19 SEACAT Recorder measures conductivity, temperature, and pressure (depth). The SEACAT is self-powered and self-contained and can be deployed in profiling or moored mode. The SBE 19 SEACAT was replaced in 2001 by the 19plus. more information from Sea-Bird Electronics |

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Deployments

LW_LIS08

| Website | https://www.bco-dmo.org/deployment/58823 |
|-------------|---|
| Platform | R/V Lowell Weicker |
| Start Date | 2008-08-13 |
| End Date | 2008-08-13 |
| Description | Samples were collected at 6 stations in Long Island Sound on 13 August 2008 as part of the project titled "Collaborative Research: Testing hypotheses about diversity, gene flow, and effective population size in marine planktonic ciliates". |

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Project Information

Testing hypotheses about diversity, gene flow, and effective population size in marine planktonic ciliates (CiliateDivGenePop)

Website: <u>http://microzooplankton.uconn.edu</u>

Coverage: Coastal Northwest Atlantic, from Long Island Sound to Maine

The microbial ecologist Tom Fenchel recently said, "The decoupling of molecular and classical (including experimental) approaches to environmental microbiology has not been fruitful and it represents one of the most important challenges for the field in the coming years." (Fenchel 2005). Classical approaches center on the centuries-old tradition of describing individual species via meticulous observation and analysis to generate monographs, such as is done for plants and animals. Unfortunately, the rush to new molecular techniques has sometimes ignored this tradition, with claims about new lineages never seen before and reports of staggering diversity of microbial eukaryotes based on environmental DNA samples not backed up by even the most elementary microscopic observations.

In the face of this disconnect between the traditional and the molecular, we propose a marriage of the two approaches in the study of marine ciliate diversity and gene flow. Our own data show that in some clades of planktonic ciliates (Strombidiidae) there is indeed a high level of molecular diversity underlying a relatively small number of morphospecies. In other clades (some choreotrichs), the opposite appears to be true, with morphological heterogeneity underlain by apparently clonal lines, based on molecular data. Currently, we do not understand what sustains diversity in some clades; nor do we know why other clades show low diversity. But this problem is amenable to both experimental and observational approaches.

This proposal uses a two-pronged approach, combining molecular (clone libraries, DGGE,FISH) and traditional (light microscopy) techniques to address three broad questions:

i. What are the most important physical and biological factors that affect distribution and diversity of planktonic marine ciliates?

ii. What is the effective population size for marine ciliate populations, and how does this compare to census population sizes?

iii. How well do traditional morphological descriptions of ciliate species fare when compared with molecular characterizations?

Using a combination of molecular and microscopy methods, we will address these questions in coastal planktonic ciliates. Analyses of the resulting data will yield insights into the nature of ciliate species and patterns of gene flow within the North Atlantic.

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Funding

| Funding Source | Award |
|--|--------------------|
| NSF Division of Ocean Sciences (NSF OCE) | <u>OCE-0648711</u> |

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