# ADCP surface current velocities and directions from R/V Atlantic Explorer cruise AE2207 to the Bermuda Atlantic Time Series (BATS) station in April 2022

Website: https://www.bco-dmo.org/dataset/877170

Data Type: Cruise Results

Version: 1

Version Date: 2022-07-18

#### **Project**

» Collaborative Research: Inferring Cellular Lysis and Regeneration of Organic Matter by Marine Viruses (InVirT)

Contributors	Affiliation	Role
Wilhelm, Steven W.	University of Tennessee Knoxville (UTK)	Principal Investigator
Sullivan, Matthew	Ohio State University	Co-Principal Investigator
<u>Weitz, Joshua</u>	Georgia Institute of Technology (GA Tech)	Co-Principal Investigator
Muratore, Daniel	Georgia Institute of Technology (GA Tech)	Student
Rauch, Shannon	Woods Hole Oceanographic Institution (WHOI BCO-DMO)	BCO-DMO Data Manager

#### **Abstract**

This dataset was gathered from ADCP for surface current velocities and directions with ship position, time, and surface temperature along with some derived variables useful for plotting. Data are from R/V Atlantic Explorer cruise AE2207 to the Bermuda Atlantic Time Series (BATS) station in April 2022.

## **Table of Contents**

- Coverage
- Dataset Description
  - Methods & Sampling
  - Data Processing Description
- Data Files
- Parameters
- Instruments
- <u>Deployments</u>
- Project Information
- Funding

## Coverage

Spatial Extent: N:32.381 E:-64.156 S:31.012 W:-64.683

Temporal Extent: 2022-04-22 - 2022-04-25

# Methods & Sampling

ADCP readings were taken approximately every 10 minutes over the course of the cruise. ADCP readings were derived from .mat files during the cruise.

#### **Data Processing Description**

#### **Data Processing:**

Data were retrieved from ADCP all\_bins .mat files. After collecting surface U and V velocity readings, the total current magnitude and direction were calculated using vector algebra. ADCP UTC time for readings were then

converted to local time.

# **BCO-DMO Processing:**

- converted date/time fields to ISO8601 format.

[ table of contents | back to top ]

# **Data Files**

File

**AE2207\_ADCP\_surface.csv**(Comma Separated Values (.csv), 165.71 KB)

MD5:df10623558d4bbce2cceab2e94b46ff7

Primary data file for dataset ID 877170

[ table of contents | back to top ]

# **Parameters**

Parameter	Description	Units
Lon	Longitude	degrees East
Lat	Latitude	degrees North
Julian_day	Julian day from reading	days
Surf_Temp	ADCP-derived surface temperature	degrees Celsius
U	Horizontal surface velocity (positive is east)	meters per second (m/s)
V	Meridional surface velocity (positive is north)	meters per second (m/s)
endpoint	Current magnitude	meters per second (m/s)
angle	Velocity vector angle in radians from due north	radians
endpoint_scaled	Current magnitude rescaled to a maximum of 0.5 for plotting	unitless
yland	Y-coordinate of endpoint for quiver plotting	degrees Noth
xland	X-coordinate of endpoint for quiver plotting	degrees East
direction	Data for determining which orientation to plot arrows for quiver plot	unitless
ISO_DateTime_UTC	Date of observation from ADCP in UTC; format is ISO8601: YYYY-MM-DDThh:mm:ssZ	unitless
rounded_time	String containing month:day:hour:minute in UTC time	unitless
ISO_DateTime_Local	Date of observation from ADCP in local time UTC+4 time; format is ISO8601: YYYY-MM-DDThh:mm:ss	unitless

[ table of contents | back to top ]

# Instruments

specific Instrument Name	ADCP using osnb75
Generic Instrument Name	Acoustic Doppler Current Profiler
Generic Instrument Description	The ADCP measures water currents with sound, using a principle of sound waves called the Doppler effect. A sound wave has a higher frequency, or pitch, when it moves to you than when it moves away. You hear the Doppler effect in action when a car speeds past with a characteristic building of sound that fades when the car passes. The ADCP works by transmitting "pings" of sound at a constant frequency into the water. (The pings are so highly pitched that humans and even dolphins can't hear them.) As the sound waves travel, they ricochet off particles suspended in the moving water, and reflect back to the instrument. Due to the Doppler effect, sound waves bounced back from a particle moving away from the profiler have a slightly lowered frequency when they return. Particles moving toward the instrument send back higher frequency waves. The difference in frequency between the waves the profiler sends out and the waves it receives is called the Doppler shift. The instrument uses this shift to calculate how fast the particle and the water around it are moving. Sound waves that hit particles far from the profiler take longer to come back than waves that strike close by. By measuring the time it takes for the waves to bounce back and the Doppler shift, the profiler can measure current speed at many different depths with each series of pings. (More from WHOI instruments listing).

## [ table of contents | back to top ]

# **Deployments**

#### **AE2207**

Dataset-

Website	https://www.bco-dmo.org/deployment/873946
Platform	R/V Atlantic Explorer
Start Date	2022-04-22
End Date	2022-04-27
Description	See additional cruise information at the Rolling Deck to Repository (R2R): https://www.rvdata.us/search/cruise/AE2207

# [ table of contents | back to top ]

# **Project Information**

Collaborative Research: Inferring Cellular Lysis and Regeneration of Organic Matter by Marine Viruses (InVirT)

**Coverage**: Bermuda Atlantic Time Series

#### NSF Award Abstract:

Viral infections of marine microbes can transform the fate of microbial populations that fuel global ocean biogeochemical cycles. For example, viral infections of microbes lead to the release of carbon and nutrients back into the environment. This regeneration of carbon and nutrients stimulates the activity of other microbes and diverts carbon and nutrients from larger organisms in marine food webs. Because virus-microbe infections are relatively specific, it is critical to identify those pairs of viruses and microbes that may disproportionally contribute to the turnover of carbon and nutrients in the ocean. This project will develop quantitative approaches and tools to quantify which viruses infect which microbes and to use these data to

quantify how viral infections of microbes collectively shape nutrient and carbon cycles in the North Atlantic Ocean. The project will analyze virus-microbe interactions in mesocosms at the Bigelow Laboratory for Ocean Sciences in mid-coast Maine and during open ocean expeditions to the Bermuda Atlantic Time-Series Study (BATS) site. An interdisciplinary team will leverage recent advances in molecular biology, computational biology, and mathematical modeling to identify virus-host partners and their impact on the movement of elements through marine systems. This project will support three graduate students, six undergraduate students and one postdoctoral researcher in an interdisciplinary context. Research advances will be translated into reproducible software methods to be disseminated via the community cyberinfrastructure platform iVirus, with additional training materials presented as part of a viral methods and informatics workshop held at The Ohio State University. The translation of discoveries to the public will be furthered by the involvement of journalism undergraduate students at the University of Tennessee-Knoxville.

This project builds upon advances in the molecular toolkit of viromics to develop an integrated approach to characterize lineage-specific rates of infection, lysis, and nutrient release induced by marine viruses in open ocean ecosystems. It will combine theory, in vitro experiments, and in situ sampling to (i) extend a robust inference method for estimating virus-microbe cross-infection networks from time-series data; (ii) establish and characterize in-vitro protocols for inferring cross-infectivity in complex communities using cultureindependent methods; (iii) estimate lineage-specific rates of lysis and regeneration of nutrients in marine systems, including applications to coastal and open ocean ecosystems. Project aims focus on quantifying the extent to which virus-induced lysis and regeneration of carbon and nutrients is heterogeneously distributed across microbial populations. To do so, the project will incorporate time series measurements of abundance information (via metagenomes) and activity information (via metatranscriptomes). In so doing, it will advance efforts to understand community-scale interactions rather than those amongst a single virus-host pair. Theoretical methods and in vitro protocols will directly infer lineage-specific infection, lysis, and nutrient release rates in coastal- and open-ocean ecosystems in the North Atlantic Ocean. Results will be used to identify key links that disproportionately influence bulk nutrient release. A novel PCR-based approach will augment and validate the core inference approach. Overall, the project aims to enhance our understanding of how viruses contribute to marine ecosystem function.

This award reflects NSF's statutory mission and has been deemed worthy of support through evaluation using the Foundation's intellectual merit and broader impacts review criteria.

# [ table of contents | back to top ]

## **Funding**

Funding Source	Award
NSF Division of Ocean Sciences (NSF OCE)	OCE-1829636
NSF Division of Ocean Sciences (NSF OCE)	OCE-1829640
NSF Division of Ocean Sciences (NSF OCE)	OCE-1829641

[ table of contents | back to top ]