

# CTD data from R/V Atlantic Explorer X0606, X0705, AE0810 in the Western Sargasso Sea roughly 38-20N and 66-43W from 2006-2008 (ATP3 project)

**Website:** <https://www.bco-dmo.org/dataset/3355>

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## Project

» [DOP Utilization in the Sargasso Sea: Quantifying Taxon-specific Rates of Hydrolysis and Uptake](#) (ATP3)

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## Table of Contents

- [Dataset Description](#)
  - [Methods & Sampling](#)
  - [Data Processing Description](#)
- [Data Files](#)
- [Parameters](#)
- [Instruments](#)
- [Deployments](#)
- [Project Information](#)
- [Funding](#)

## Dataset Description

DOP Utilization (ATP3, DOP) CTD data  
Biogeochemical data collected on transect cruises studying  
Dissolved Organic Phosphorus throughout the western Sargasso Sea.  
Data are from several cruises over the span 2006 to 2008.

Note the cruise identifiers for the Atlantic Explorer were originally formatted as XYY## (e.g. X0806 was the 6th cruise in 2008). The data files include cruise IDs of this type. The vessel operator changed the cruise ID syntax several years after the cruise and the official cruise ID syntax was changed to AEYY##. For example, AE0810 should be the same cruise as X0810. One exception for this dataset is that X0804 is cruise ID AE0810.

### Related files and references:

*Detailed information on analyses:*

Lomas, M.W., Burke, A., Lomas, D.A., Bell, D.W., Shen, C., Ammerman, J.W., Dyhrman, S.T. 2010. Sargasso Sea phosphorus biogeochemistry: An important role for dissolved organic phosphorus (DOP). *Biogeosciences* 7: 695-710.

*Other published work:*

Michelou, V.K., Lomas, M.W., Kirchman, D.L. Phosphate and ATP uptake by cyanobacteria and heterotrophic bacteria in the Sargasso Sea. *Limnology and Oceanography*, in press.

McLaughlin, K., Sohm, J.A., Cutter, G.A., Lomas, M.W., Paytan, A. 2010. Phosphate cycling in the Sargasso Sea: Investigation using the oxygen isotopic composition of phosphate, enzyme labeled fluorescence, and turnover times.

Longnecker, K., Lomas, M.W., and Van Mooy, B.A.S. 2010. Characterizing the abundance and diversity of heterotrophic bacterial cells assimilating phosphate in the sub-tropical North Atlantic Ocean. *Environmental Microbiology*, in press.

Orchard, E.D., Ammerman, J.W., Lomas, M.W., Dyhrman, S.T. 2010. Dissolved inorganic and organic phosphorus uptake in *Trichodesmium* and the microbial community: The importance of phosphorus ester in the Sargasso Sea. *Limnology and Oceanography*, 55:1390-1399.

Casey, J., Lomas, M.W., Michelou, V., Orchard, E.D., Dyhrman, S.T., Ammerman, J.W., and Sylvan, J. 2009. Phytoplankton taxon-specific orthophosphate (Pi) and ATP uptake in the northwestern Atlantic subtropical gyre. *Aquatic Microbial Ecology*, 58:31-44.

Van Mooy, B.A.S., Fredricks, H.F., Pedler, B.F., Dyhrman, S.T., Karl, D.M., Koblizek, M., Lomas, M.W., Moore, L.R., Moutin, T., Rappé, M.S., and Webb, E.A. 2009. Phytoplankton in the oligotrophic ocean use non-phosphorus lipids in response to phosphorus scarcity. *Nature Geosciences*, doi:10.1038/nature07659.

## **Methods & Sampling**

### **Sampling and Analytical Methodology:**

Detailed methods for all data collected as part of this study can be found in the publications arising from this study (references given below). This contains information on analytical machines and certified standards where applicable.

Sample QA/QC procedures followed those of the Bermuda Atlantic Time-series Study (BATS). At the point of collection, any leaking niskin bottles were noted on the master cast sheets and samples were taken from a different niskin fired at the same depth as the leaking bottle. No data are reported for leaking Niskin bottles. During sample analysis standard curves and/or certified standards were carefully examined to ensure that they were consistent with expectations and accurate. If nothing was found, then we examined other data from that niskin to see if other samples are in question. If no obvious error or problem was found, the data were considered OK and in the range of environmental data that this study hoped to observe.

### **Sample accuracy and precision:**

Sample accuracy was assessed by using certified standards, for those measurements where standards are available (dissolved oxygen, nutrients, salinity). Certified standards were run with each analytical run and compared to long term control charts for respective analyses. Samples were not run until certified standards were shown to be accurate for that analytical run. Sample precision was determined by analyzing replicate samples (not replicate analyses on the same sample) and therefore is higher than analytical precision due to the inclusion of sampling error. At the concentrations observed during this study, sample precision was <5% for stock measurements and <10% for rate measurements. Some analyses, namely dissolved oxygen and salinity, were much better and often had a sample precisions <1%. These precision estimates are consistent with the long term QA/QC seen with the BATS program.

The provided data files are complete matrices and therefore not every sample (columns) will be taken from every Niskin fired (rows). Data that were either not collected, or were associated with leaking Niskins, or were found to be in error for other reasons are denoted by "nd" in the spreadsheets.

## Data Processing Description

### Data Processing:

Most of the data given in this dataset are not derived variables and are calculated using reasonably standard equations as given in the appropriate reference above. The one exception is CTD data. Raw CTD data were processed using SBE-Data Processing software using configuration and calibration files provided by the Shipboard Science technician. Sensors were calibrated every 6 months. Manual determinations of dissolved oxygen, salinity and HPLC Chlorophyll a, once passing the above QA/QC steps, were taken as correct. CTD sensor data was regressed against the appropriate manual variable. In all cases, regression statistics were very strong and linear. CTD data were corrected to manual measurements using the regression data and it is this corrected data that is given in the associated data files.

Only nutrient analyses were close to analytical method detection limits (MDL). MDLs were estimated as 3x the standard deviation of the lowest standard used for the analysis and are ~30nM for nitrate and phosphate using a standard autoanalyzer. We used the MAGIC co-precipitation method for phosphate which lowered our MDL to ~1.5nM. Samples below the MDL are reported as calculated for the reason that they are somewhere between the MDL and a true zero; we consider listing them as either to be incorrect.

### BCO-DMO Edits

- Parameter names modified to conform to BCO-DMO convention
- date reformatted to YYYYMMDD
- time reformatted to HHMM
- lat/lon padded to 7 decimal places
- added CruiseId column and combined all CTD data into one dataset
- X0606 station 7 year corrected from 2007 to 2006
- Lat/Lons for X0606 standardized to positions from BGC data

[ [table of contents](#) | [back to top](#) ]

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## Data Files

File
<b>CTD.csv</b> (Comma Separated Values (.csv), 4.53 MB) MD5:c36a0e2523cdbee021112c536c5713d0 Primary data file for dataset ID 3355

[ [table of contents](#) | [back to top](#) ]

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## Parameters

Parameter	Description	Units
CruiseId	Cruise Id	text
Station	Station Id	integer
Cast	CTD drop number	integer
date	date of operation (GMT)	YYYYMMDD

time	time of operation (GMT)	HHMM
lon	Longitude position West is negative	decimal degrees
lat	Latitude position South is negative	decimal degrees
Depth	depth	meters
Depth_actual	depth retrieved from CTD pressure sensor	meters
Temp_1	primary temperature sensor	degrees celsius
Temp_2	secondary temperature sensor	degrees celsius
Salinity_1	primary conductivity sensor	dimensionless
Salinity_2	secondary conductivity sensor	dimensionless
Sigma_theta_1	density derived from primary sensors	kg/m3
Sigma_theta_2	density derived from secondary sensors	kg/m3
Fluorescence	CTD Fluormeter	relative
DO_1	primary dissolved oxygen sensor	umol/kg
DO_2	secondary dissolved oxygen sensor	umol/kg

[ [table of contents](#) | [back to top](#) ]

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## Instruments

<b>Dataset-specific Instrument Name</b>	CTD Sea-Bird 911
<b>Generic Instrument Name</b>	CTD Sea-Bird 911
<b>Dataset-specific Description</b>	1.0 CTD 1.1 Seabird Electronics SBE 9/11+ Max Depth is 6800m for all CTD sensors except Chelsea Fluorometer, Wetlabs Transmissometer and Altimeter which are all 6000m. Aluminum frame holds 24 12 liter water samplers. 1.2 Sensors include Seabird SBE 9/11+, Dual pumped Temperature, conductivity and Dissolved Oxygen. Chelsea Aquatracka III Fluorometer, Wetlabs SeaStar 25cm/660nm Transmissometer, Benthos PSA9000 Altimeter. All data logged with Seabird Software. 1.3 Water Samplers - 24x12 liter Ocean Test Equipment (OTE) Niskin Water samplers and 4x10 liter OTE Go-Flo's.
<b>Generic Instrument Description</b>	The Sea-Bird SBE 911 is a type of CTD instrument package. The SBE 911 includes the SBE 9 Underwater Unit and the SBE 11 Deck Unit (for real-time readout using conductive wire) for deployment from a vessel. The combination of the SBE 9 and SBE 11 is called a SBE 911. The SBE 9 uses Sea-Bird's standard modular temperature and conductivity sensors (SBE 3 and SBE 4). The SBE 9 CTD can be configured with auxiliary sensors to measure other parameters including dissolved oxygen, pH, turbidity, fluorescence, light (PAR), light transmission, etc.). More information from Sea-Bird Electronics.

[ [table of contents](#) | [back to top](#) ]

## Deployments

### X0606

<b>Website</b>	<a href="https://www.bco-dmo.org/deployment/58060">https://www.bco-dmo.org/deployment/58060</a>
<b>Platform</b>	R/V Atlantic Explorer
<b>Start Date</b>	2006-05-19
<b>End Date</b>	2006-05-27
<b>Description</b>	One in a series of transect cruises to study the biological and biogeochemical aspects of the marine phosphorus cycle.

### X0705

<b>Website</b>	<a href="https://www.bco-dmo.org/deployment/58061">https://www.bco-dmo.org/deployment/58061</a>
<b>Platform</b>	R/V Atlantic Explorer
<b>Start Date</b>	2007-06-02
<b>End Date</b>	2007-06-14
<b>Description</b>	One in a series of transect cruises to study the biological and biogeochemical aspects of the marine phosphorus cycle.

### AE0810

<b>Website</b>	<a href="https://www.bco-dmo.org/deployment/58062">https://www.bco-dmo.org/deployment/58062</a>
<b>Platform</b>	R/V Atlantic Explorer
<b>Start Date</b>	2008-05-03
<b>End Date</b>	2008-05-25
<b>Description</b>	One in a series of transect cruises to study the biological and biogeochemical aspects of the marine phosphorus cycle. Note the cruise identifiers for the Atlantic Explorer were originally formatted as XYY## (e.g. X0806 was the 6th cruise in 2008). The data files include cruise IDs of this type. The vessel operator changed the cruise ID syntax several years after the cruise and the official cruise ID syntax was changed to AEYY##. For example, AE0810 should be the same cruise as X0810. One exception for this dataset is that X0804 is cruise ID AE0810 (unclear how the cruise numbering scheme got so confused). Database validation showed that AE0804 was not the correct cruiseid based on information at R2R. The cruiseid was then updated to reflect the corrected information (the May 2008 cruise was AE0810. Additional Information from R2R Site

[ [table of contents](#) | [back to top](#) ]

## Project Information

### DOP Utilization in the Sargasso Sea: Quantifying Taxon-specific Rates of Hydrolysis and Uptake (ATP3)

**Website:** [http://www.bios.edu/Labs/pel/Research%20Pages/Research\\_DOP.html](http://www.bios.edu/Labs/pel/Research%20Pages/Research_DOP.html)

**Coverage:** Western Sargasso Sea roughly 38-20oN and 66-43oW. <br>Water depths always exceeded 4200m.

Photosynthetic uptake of CO<sub>2</sub> by oceanic phytoplankton and the export of the resulting organic carbon to the deep sea comprise a 'biological pump' capable of extracting globally significant amounts of CO<sub>2</sub> from the atmosphere. Mounting evidence suggests that primary production in two of the larger subtropical ocean gyres, the Western Tropical/Subtropical Atlantic and the North Pacific Subtropical Gyre (NPSG), may be controlled by the availability of inorganic phosphorus. This conclusion is based on vanishingly low inorganic phosphorus (SRP) concentrations, sub-nanomolar in some locales, ratios of inorganic nutrient availability that greatly exceed the canonical Redfield Ratio, and high rates of dissolved organic phosphorus (DOP) hydrolysis.

Indeed, data collected in the Sargasso Sea shows a 30% decline in DOP inventories during summer stratification. Moreover, several studies have documented significant taxonomic variability in the ability to hydrolyze and to assimilate phosphorus from organic sources. We hypothesize that despite rapid turnover times, chronically low and seasonally invariant SRP concentrations at BATS cannot support measured rates of primary production without utilization of additional P from the DOP pool. Moreover, we hypothesize that inherent physiological differences among microbial taxa represents a significant source of temporal and spatial variability in DOP utilization rates that is yet neither understood nor constrained.

Our specific research objectives are:

1. To quantify temporal and spatial variability in DOP hydrolysis in the Sargasso Sea with measures of whole-community and taxon-specific alkaline phosphatase
2. To quantify temporal and spatial variability in taxon-specific SRP and DOP uptake rates by combining flow cytometry and radioisotope methodologies.
3. To quantify whole-community total P uptake rates through BAP (biologically available phosphorus) assays, as well as SRP and model compound DOP uptake and regeneration rates.
4. To identify factors regulating rates of DOP hydrolysis and assimilation using experimental nutrient manipulations, and to evaluate the role of DOP in supporting primary production in the Sargasso Sea.

To successfully meet our objectives, we propose to employ three cruise sampling strategies: CORE, PROCESS, and CRUISES OF OPPORTUNITY. The CORE cruises and CRUISES OF OPPORTUNITY will be conducted in conjunction with the BATS biogeochemical time-series program. The PROCESS cruises are principal-use cruises

that are designed to allow a more intensive study on the mechanisms of and controls on DOP hydrolysis and utilization in the Sargasso Sea.

An understanding of ocean ecosystem function is important on a broad scale. This project will provide information critical for successful modeling efforts constrain predictions of the strength of the oceanic biological pump, as well as provide information of interest to students, teachers and the general public. If in fact DOP supports a significant, and previously unquantified, fraction of the annual primary production in the Sargasso Sea, then diversity in biological metabolic processes in the central oceans plays a greater role in the global carbon cycle - including regulation of atmospheric CO<sub>2</sub> - than we recognize at present. The overall goal of the student teaching/training programs at BBSR, WHOI and Rutgers is to expose students to oceanographic research, its global significance, and its impact on their daily lives. As such, we will incorporate data on DOP cycling in the Sargasso Sea into a problem-based learning module for courses taught by the PIs and submit our curriculum to the appropriate digital repository (e.g. [www.dlese.org](http://www.dlese.org)). The PIs have a strong commitment to direct mentoring, and they will also sponsor a minimum of three undergraduate researchers each year in their laboratories, and support the research and training of MIT/WHOI Joint Program and Rutgers University graduate students.

[ [table of contents](#) | [back to top](#) ]

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## Funding

Funding Source	Award
<a href="#">NSF Division of Ocean Sciences (NSF OCE)</a>	<a href="#">OCE-0453023</a>

[ [table of contents](#) | [back to top](#) ]