# Concentrations of particulate phosphate in treatments amended with dissolved organic phosphate compounds in bioassay experiments (incubation 48h) with seawater from R/V Savannah cruise SAV-19-02 in the NW Atlantic Ocean in Spring of 2019

Website: https://www.bco-dmo.org/dataset/864225

Data Type: Cruise Results, experimental

Version: 1

Version Date: 2021-11-02

#### **Project**

» <u>Collaborative Research: Assessing the role of compound-specific phosphorus hydrolase transformations in the marine phosphorus cycle</u> (P-hydrolase)

Contributors	Affiliation	Role
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#### **Abstract**

Concentrations of particulate phosphate in treatments amended with dissolved organic phosphate compounds, over an incubation period of 48 h during bioassay experiments with seawater collected during R/V Savannah cruise SAV-19-02 from March to April of 2019 in the Northwestern Atlantic from the surface to 50 m depth.

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## Coverage

**Spatial Extent**: N:31.7635 E:-79.8421 S:31.0175 W:-80.7965

**Temporal Extent**: 2019-03-30 - 2019-04-10

# Methods & Sampling

## Sampling and analytical procedures:

Bioassay experiments were conducted at station 1 and stations 3. At each station, inorganic and organic phosphate amendments were performed on seawater with and without nitrogen enrichment (NH4Cl, NaNO3). Bioassay Experiments consisted in incubating, over an incubation period of 48h, surface seawater (5m) with inorganic or organic phosphate compounds (20  $\mu$ M; final concentration of P) including, polyphosphate (polyp), inorganic phosphate (Pi), nucleotides (ATP or AMP) and methylphosphonate (Mepn). In each incubation experiment, a control treatment (surface seawater without amendment) was included.

In bioassay experiments, subsamples were taken for particulate phosphate (PP, 50 mL) from each experimental incubation bottle and then filtered through pre-combusted ( $450^{\circ}$ C, 4.5 h) and acid washed (1N HCl) GF/F syringe filters (25 mm). PP filters were wrapped in pre-combusted ( $450^{\circ}$ C, 4.5 h) aluminum foil, and kept frozen (- $20^{\circ}$ C) until analysis. PP concentrations were determined after high-temperature combustion ( $450^{\circ}$ C, 4.5 h) and HCl extraction (0.5 N) (Strickland and Parsons, 1972). This method relies on the release and the subsequent extraction of organically bound P compounds as SRP. SRP was then analyzed spectrophotometrically using a conventional spectrophotometer ( $10^{\circ}$ C cm cuvette). The efficiency of the procedure was tested on apple leaves (standard reference material 1515) and resulted in a yield of  $83\pm19\%$  (n=20).

**Instrument**: Measurements of PP were performed using a conventional spectrophotometer (Genesys®,10 cm cuvette) after a high temperature combustion (450°C, 4.5h).

**Location:** Northwestern Atlantic surface waters. Depth: surface-50 m.

## **Data Processing Description**

Data were organized using MATLAB and output as .mat files. Gaps in data were filled with NaN in the .mat files.

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## **Related Publications**

Strickland, J. D. H. and Parsons, T. R. (1972). A Practical Hand Book of Seawater Analysis. Fisheries Research Board of Canada Bulletin 157, 2nd Edition, 310 p. *Methods* 

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#### **Parameters**

Parameters for this dataset have not yet been identified

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#### Instruments

Dataset-specific Instrument Name	Genesys®,10 cm cuvette
Generic Instrument Name	Spectrophotometer
Dataset-specific Description	Measurements of PP were performed using a conventional spectrophotometer (Genesys®,10 cm cuvette) after a high temperature combustion (450°C, 4.5h).
Generic Instrument Description	An instrument used to measure the relative absorption of electromagnetic radiation of different wavelengths in the near infra-red, visible and ultraviolet wavebands by samples.

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# **Deployments**

#### SAV-19-02

Website	https://www.bco-dmo.org/deployment/864191
Platform	R/V Savannah
Start Date	2019-03-30
End Date	2019-04-11
Description	Cruise synonym: Zephyr (Zooming in on Enzymatic PhosphoHYdrolysis Reactions)

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## **Project Information**

Collaborative Research: Assessing the role of compound-specific phosphorus hydrolase transformations in the marine phosphorus cycle (P-hydrolase)

## NSF Award Abstract:

Phosphorus (P) is an essential building block for life. Because P is in short supply over vast areas of the ocean, P availability may control biological productivity, such as photosynthesis and carbon fixation, which has implications for uptake of the greenhouse gas carbon dioxide and thus climate regulation. Marine microorganisms must satisfy their nutritional requirement for P by obtaining it from seawater, where P is present in a variety of chemical forms, from simple phosphate ions (Pi) to complex dissolved organic phosphorus (DOP) molecules. The concentration of DOP vastly exceeds Pi over most ocean areas, therefore DOP is a critically important source of P for marine microbial nutrition and productivity. However, much remains unknown about the contribution of specific DOP compounds to the P nutrition, productivity, and structure of marine microbial communities. In this project, the investigators will conduct field experiments in the Atlantic Ocean and perform a series of controlled laboratory studies with pure enzymes and microbial cultures to determine how and to what extent different DOP compounds are degraded to Pi in the marine environment. Furthermore, the contribution of these compound-specific DOP molecules to microbial P nutrition, carbon fixation, and community structure will be determined, thus advancing the current state of knowledge regarding the factors that control the activity and distribution of microbial species in the ocean, and the ocean?s role in the climate system. This project will support two female junior investigators, a postdoctoral researcher, and graduate and undergraduate students. The undergraduate students will be recruited from the Marine Sciences program at Savannah State University, an Historically Black Colleges and Universities. In addition, results will be incorporated into new hands-on K-12 educational tools to teach students about microbial P biogeochemistry, including a digital game and formal lesson plans with hands-on demos. These tools will be validated with K-12 educators and will be widely accessible to the public through various wellknown online platforms. These activities will thus reach a broad audience including a significant fraction of underrepresented groups.

P is a vital nutrient for life. Marine microorganisms utilize P-hydrolases, such as alkaline phosphatase (AP), to release and acquire phosphate (Pi) from a wide diversity of dissolved organic P (DOP) compounds, including Pesters (P-O-C bonds), phosphonates (P-C), and polyphosphates (P-O-P). Compound-specific DOP transformations have the potential to exert critical and wide-ranging impacts on marine microbial ecology (e.g. variable DOP bioavailability among species), biogeochemistry (e.g. P geologic seguestration via formation of calcium Pi), and global climate (e.g. aerobic production of the greenhouse gas methane by dephosphorylation of methylphosphonate). However, the mechanisms and comparative magnitude of specific DOP transformations, in addition to their relative contributions to microbial community-level P demand, productivity, and structure, are not completely understood. This study will fill these knowledge gaps by tracking the fate of specific DOP pools in the marine environment. Specifically, this project will test four hypotheses in the laboratory using recombinant enzymes and axenic cultures representative of marine eukaryotic and prokaryotic plankton from high and low nutrient environments, and in the field using observational and experimental approaches along natural Pi gradients in the Atlantic Ocean. In particular, the investigators will reveal potential differences in the hydrolysis and utilization of specific DOP compounds at the community- (bulk enzymatic assays), taxon- (cell sorting of radiolabeled cells in natural samples), species- (axenic cultures) and molecular-levels (pure enzyme kinetic studies and cell-associated proteomes and exoproteomes). Results from our proposed work will provide a robust understanding of the enzymatic basis involved in the transformation of specific forms of DOP and create new knowledge on the relative contribution of these specific P sources to

Pi production, marine microbial nutrition, community structure, primary productivity, and thus global carbon cycling and climate. In particular, our refined measurements of the concentration of bioavailable DOP and our unique estimates of DOP remineralization fluxes will provide critical new information to improve models of marine primary production and P cycling.

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# **Funding**

Funding Source	Award
NSF Division of Ocean Sciences (NSF OCE)	OCE-1736967
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