

# Bioavailability factor of polyphosphate, nucleotides (ATP and AMP) and phosphonate in seawater collected during R/V Savannah cruise SAV-19-02 in the NW Atlantic Ocean in the Spring of 2019

**Website:** <https://www.bco-dmo.org/dataset/864280>

**Data Type:** Cruise Results

**Version:** 1

**Version Date:** 2021-11-02

## Project

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

Contributors	Affiliation	Role
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## Abstract

Bioavailability factor of polyphosphate, nucleotides (ATP and AMP) and phosphonate in 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:

Seawater samples for bioavailability factor determinations were collected from Niskin bottles (12 L), at station 1 (two measurements) and 3 (one measurement) along a transect from coastal Georgia to offshore waters. The bioavailability factor was calculated based on changes in PO<sub>4</sub><sup>3-</sup> turnover time. The bioavailability factor (BF) is calculated as follows:  $BF = TE - TN / TP - TN$  (Björkman and Karl, 1994), where TE reflects PO<sub>4</sub><sup>3-</sup> turnover time in the DOP amended treatment, TN the PO<sub>4</sub><sup>3-</sup> turnover time in the control (no additions) and TP the PO<sub>4</sub><sup>3-</sup> turnover time in the treatment amended with Pi. In this study, treatments were amended with polyphosphate (Polyp), adenosine triphosphate (ATP), adenosine monophosphate (AMP) or methyl phosphonate (Mepn).

BF ranges from 0 for an unavailable substrate, to a value of 1 for a DOP model substrate having a bioavailability equal to that of +Pi.

**Instruments:** Sampling was performed using Niskin bottles (12 L) mounted on a rosette. Radioactivity was assayed on a Packard Tri-Carb liquid scintillation counter.

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

**Problem report:** *BF in station 1 (first determination) were determined in duplicates instead of triplicate in station 1 (second determination) and station 3. Therefore, gaps are filled with NaN in the .mat files.*

## 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

Björkman, K., & Karl, D. M. (1994). Bioavailability of inorganic and organic phosphorus compounds to natural assemblages of microorganisms in Hawaiian coastal waters. *Marine Ecology Progress Series*, 111(3), 265–273. <http://www.jstor.org/stable/24849565> <https://www.jstor.org/stable/24849565>  
*Methods*

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## Parameters

*Parameters for this dataset have not yet been identified*

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## Instruments

<b>Dataset-specific Instrument Name</b>	
<b>Generic Instrument Name</b>	Liquid Scintillation Counter
<b>Dataset-specific Description</b>	Radioactivity was assayed on a Packard Tri-Carb liquid scintillation counter.
<b>Generic Instrument Description</b>	Liquid scintillation counting is an analytical technique which is defined by the incorporation of the radiolabeled analyte into uniform distribution with a liquid chemical medium capable of converting the kinetic energy of nuclear emissions into light energy. Although the liquid scintillation counter is a sophisticated laboratory counting system used to quantify the activity of particulate emitting ( $\beta$ and $\alpha$ ) radioactive samples, it can also detect the auger electrons emitted from $^{51}\text{Cr}$ and $^{125}\text{I}$ samples. Liquid scintillation counters are instruments assaying alpha and beta radiation by quantitative detection of visible light produced by the passage of rays or particles through a suitable scintillant incorporated into the sample.

<b>Dataset-specific Instrument Name</b>	
<b>Generic Instrument Name</b>	Niskin bottle
<b>Dataset-specific Description</b>	Sampling was performed using Niskin bottles (12 L) mounted on a rosette.
<b>Generic Instrument Description</b>	A Niskin bottle (a next generation water sampler based on the Nansen bottle) is a cylindrical, non-metallic water collection device with stoppers at both ends. The bottles can be attached individually on a hydrowire or deployed in 12, 24, or 36 bottle Rosette systems mounted on a frame and combined with a CTD. Niskin bottles are used to collect discrete water samples for a range of measurements including pigments, nutrients, plankton, etc.

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## Deployments

### SAV-19-02

<b>Website</b>	<a href="https://www.bco-dmo.org/deployment/864191">https://www.bco-dmo.org/deployment/864191</a>
<b>Platform</b>	R/V Savannah
<b>Start Date</b>	2019-03-30
<b>End Date</b>	2019-04-11
<b>Description</b>	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 well-known 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 P-esters (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 sequestration 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
<a href="#">NSF Division of Ocean Sciences (NSF OCE)</a>	<a href="#">OCE-1736967</a>
<a href="#">NSF Division of Ocean Sciences (NSF OCE)</a>	<a href="#">OCE-1737083</a>
<a href="#">NSF Division of Ocean Sciences (NSF OCE)</a>	<a href="#">OCE-2001212</a>
<a href="#">NSF Division of Ocean Sciences (NSF OCE)</a>	<a href="#">OCE-1948042</a>

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