

Reconstructing Seasonal Changes in Surface Particulate Organic matter using Carbon Isotopes of Amino Acids

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INTRODUCTION:

Background:

- How climate change will affect biological carbon pump is not well understood (Henson et al. 2022)
- There is a lack of understanding on historical carbon trends and marine primary productivity
 - Paleo-proxy data can be used to fill in historic data gap

Carbon Isotopes:

- ¹²C & ¹³C are stable isotopes
- ¹²C 98.9% Abundant
- ¹³C 1.1% Abundant

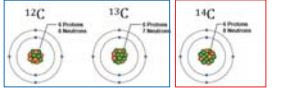


Figure 1. Diagram of Carbon Isotopes ¹²C, ¹³C, ¹⁴C adapted from Schmittner 2021

Amino Acids:

- Amino Acids (AAs) make up the largest characterizable molecular fraction of organic material at all depths
- Essential Amino Acids (EAAs) are not modified up the food chain

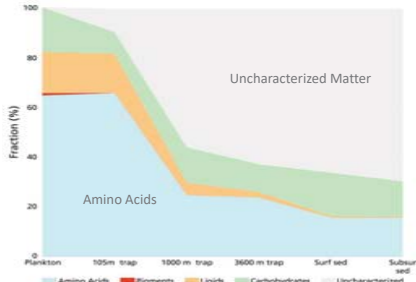


Figure 2. Biochemical composition of organic matter by depth adapted from Middleburg 2019.

Compound-specific Stable Isotope Analysis of Amino Acids (CSIA-AA)

- Delta 13 carbon ($\delta^{13}C$) can be used to create carbon fingerprints due to carbon fractionation
 - Different taxa have unique, ¹²C to ¹³C ratio: $\delta^{13}C$ values



$$\delta^{13}C = \frac{^{13}C/^{12}C_{sample} - ^{13}C/^{12}C_{standard}}{^{13}C/^{12}C_{standard}} \times 1000\text{‰}$$

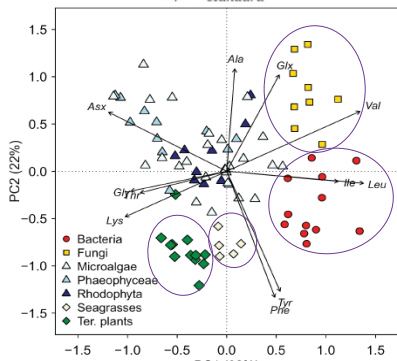


Figure 3. Principal Component Analysis of ¹³C-CAA modified Larsen et al. 2013, purple circles show major phylogenetic groupings.

Hypothesis:

- That the seasonal transition in carbon source from large-cell, diatom-based production in spring to small-cell, flagellate-based production in summer is reflected in $\delta^{13}C$ -AA signatures of seawater residue.

Objectives:

- Sample local coastal water for $\delta^{13}C$ bi-weekly following a spring to fall seasonal cycle
- Compare the found $\delta^{13}C$ values to those in the literature to validate carbon source composition
- Compare the values of $\delta^{13}C$ -AA to the bulk $\delta^{13}C$ values

METHODS:

Sample Site: Northwest Arm, Nova Scotia

- Oakland Park Road Dock (Halifax NS)
- Bi-weekly at High tide
- Surface seawater collected

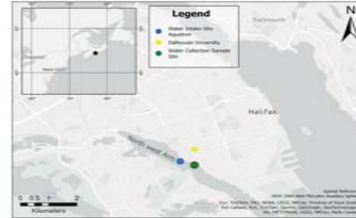
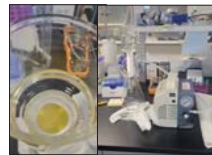


Figure 4. Map of Study site, green dot shows Oakland Park Dock made by Rori Mulholland July 2024.



Field collection for water samples



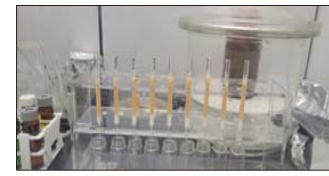
Water Filtered



Samples frozen and stored

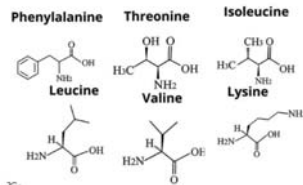


Samples freeze dried



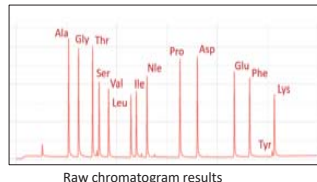
Samples Purified and Processed via Wet Chemistry and Derivatization

Essential Amino Acid's:

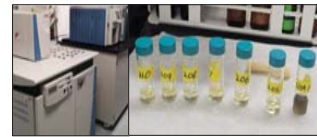


$$\delta^{13}C_{DER} = X\delta^{13}C_{AA} + (1 - X)\delta^{13}C_{ISO,TFAA}$$

Formula to remove Additive Carbon



Raw chromatogram results



Samples run through GC-IRMS



Samples Purified and Processed via Wet Chemistry and Derivatization

Workflow: N-trifluoroacetyl i-propyl ester method

- Hydrolysis
- Column & Syringe Filtration
- Cation Exchange Cleanup
- Esterification
- Acylation and Derivatization

RESULTS:

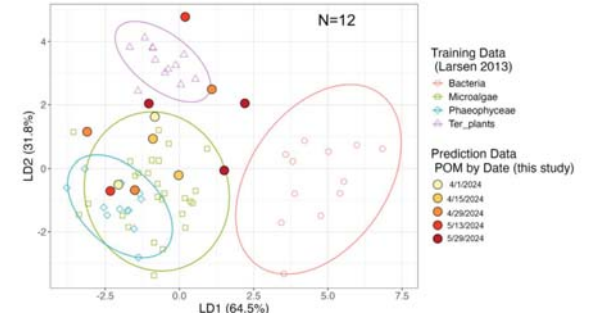


Figure 5. Linear Discrimination Analysis (LDA) of 5 spring sample dates using Larsen et al. 2013 training data.

- April samples fall into Microalgae and Phaeophyceae range
- May samples have increase pull to Bacteria and Terrestrial plants
 - Suggest higher terrestrial carbon input in later spring

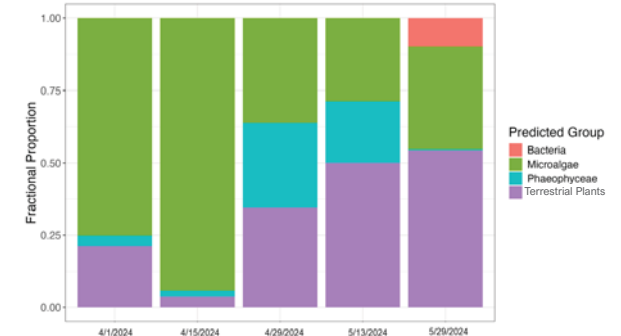


Figure 6. Time Series of Spring bloom sample dates showing fractional proportional of ¹³C signatures.

- Time series of fractional proportions support same findings as LDA
- Increase in Bacterial proportion in later May

CONCLUSIONS & NEXT STEPS:

- $\delta^{13}C$ values appear to identify carbon sources of seawater by major phylogenetic groups for 5 sample dates
- Analyze remaining 2024 sample dates
- Analyze bulk carbon and nitrogen data
- Expand training dataset

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Image of large particulate matter at sample site, Photo by R. Mulholland