Biogeosciences Discussions, 2, S673–S675, 2005 www.biogeosciences.net/bgd/2/S673/ European Geosciences Union © 2005 Author(s). This work is licensed under a Creative Commons License.



BGD

2, S673-S675, 2005

Interactive Comment

Interactive comment on "Bacterial carbon sources in coastal sediments: a review based on stable isotope data of biomarkers" by S. Bouillon and H. T. S. Boschker

Anonymous Referee #1

Received and published: 8 November 2005

The authors have brought together both previously published and new stable isotope data from various coastal environments, from mangrove systems to C3 and C4 marshes to seagrasses and unvegetated sites, to better generalize the source of carbon that is being remineralized by bacteria in these coastal systems. Phospholipid fatty acids (PLFA) were used as bacterial biomarkers; the isotopic composition of i+a 15:0 was compared to the isotopic composition of both bulk sedimentary organic carbon and macrophyte biomass. It was determined that in most settings, bacteria depend on carbon from many different sources, and not just the dominant macrophyte present. The bacterial PLFA d13C values tend to be more 13C-enriched in C3 marshes (and mangrove sites) and 13C-depleted in C4 marshes (and seagrasses and macroalgae sites) relative to the dominant d13C of the macrophytes present.

Full Screen / Esc

Print Version

Interactive Discussion

Discussion Paper

FGU

This is a nicely organized and written manuscript. The authors have also done a good job in describing how representative their data is of the various coastal environments and the veracity of the conclusions drawn. The main suggestion that I have would be to better incorporate the possibility that the bacteria are using specific components of the macrophyte biomass, which can have either 13C-enriched or 13C-depleted d13C values relative to the bulk plant material. Overall, this manuscript is a nice addition to field of estuarine/marine isotope biogeochemistry. I have only a few specific comments to add that the authors might want to take into consideration.

Specific comments:

Page 1628, lines 11-16: The isotopically-enriched source that the bacteria appear to be using could be either proteins or carbohydrates. Both components tend to have 13C-enriched isotope values relative to the bulk. Proteins and carbohydrates also tend to be more labile than other plant components.

Page 1628, lines 23-24: I am not familiar with a "cumulative sum analysis." Could the authors describe this?

Pagae 1630, lines 3 on: Only if the isotopic composition of the input organic matter is different would differences in lability and degradability be seen in the isotopic composition of the bacterial fatty acids. Are the authors assured that the algae and the macrophyte derived materials are different at the various sites?

Figure 3: When average values of isotope data were used to assign plant values, how many points made up these averages (what was the n?)? What was the error on the values?

Figure 5: I'm surprised that the standard deviation of Dd around the low %TOC values does not appear to be any larger than around the high %TOC values, yet the scatter in the data (5a) is so much greater. Is the standard deviation depicted the s.d. overall or was it binned?

BGD

2, S673-S675, 2005

Interactive Comment

Full Screen / Esc

Print Version

Interactive Discussion

Discussion Paper

EGU

Technical corrections:

Table 1: The abbreviation for Delaware is DE, so Canary Creek should read Canary Creek (DE, USA).

Interactive comment on Biogeosciences Discussions, 2, 1617, 2005.

BGD

2, S673-S675, 2005

Interactive Comment

Full Screen / Esc

Print Version

Interactive Discussion

Discussion Paper

EGU

S675