A ¹³C DOC tracer approach to estimate the contribution of semilabile dissolved organic carbon to stream ecosystem metabolism

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Introduction

Field Methods: ¹³C Release and Metabolism





Fig. 3. (A) Bioreactors kept in the dark and fed stream water in the upflow mode develop gradients of bacterial densities, activity, and species composition. (B) Residence time influences the amount of DOC metabolized. Uptake of BDOC of lower lability increases in larger bioreactors with increasing empty bed contact time (EBCT) and leads to effluents that are increasingly refractory. A series of bioreactors with increasing residence times from 0.5 min to 74 min. were used to separate the stream water DOC and the ¹³C -DOC tracer into different biological lability classes.



dominated by labile and semi-labile lability classes, respectively. Mass transfer coefficients (V_f) or uptake (mg/L) Average heterotrophic metabolism: 1.22 g C/m²/d Uptake of stream DOC: (mass transfer coefficient) x (lability class concentration) = flux Time Labile = (45 μ m/s) (250 μ g C/L) = 0.97 g C/m²/d Fig. 8. Diel oxygen curves for the 5 stations used to calculate metabolism. Ecosystem Semi-labile = (2.4 μ m/s) (804 μ gC/L) = 0.19 g C/m²/d metabolism generated from O_2 uptake in the dark, corrected for algal respiration to approximate heterotrophic respiration (H_r), and DOC uptake (1.16 g C/m²/d) can support 85% of the stream converted to units of C. H_r rates ranged from metabolism with the semi-labile constituents supporting 0.05 to 4.8 gO₂/m²/d and averaged 3.2 \pm 1.6 15% of the metabolism. $gO_2/m^2/d$ (mean, s.d.)

Fig. 4. Lability profiles of DOC from White Clay Creek stream water (open bars) and the ¹³C-DOC tracer (solid bars) from fresh (A) and soil-aged (B) tree tissues generated in stream water-fed bioreactors as a function of residence time or EBCT. Labile DOC (≤ 1.5 min. EBCT) accounted for 85% of the fresh tracer and 22% of the soil-aged tracer while the semi-labile class (≥3 min. EBCT) accounted for 15% of the fresh tracer and 78% of the soil-aged tracer. Bioreactor lability profiling was calibrated with glucose and arabinose. Glucose was completely removed in the bioreactors after EBCT of 0.5 min. while arabinose was completely removed after EBCT of 6 min. In whole stream releases of glucose and arabinose, the uptake length of arabinose was ~ 3 times longer than that of glucose $(2.9 \pm 0.7, n = 59; mean, s.d.)$.

The stream water BDOC data from the bioreactors provide concentrations of the labile and semi-labile BDOC classes. These concentrations are used along with mass transfer coefficients derived from the whole stream additions to estimate fluxes of DOC uptake in the stream ecosystem.

Conclusions

Soil-aging of the ¹³C tree tissue tracer increased our ability to measure the uptake of semi-labile DOC classes in whole-stream releases. The bioreactors are laboratory tools that facilitate measurements of DOC uptake without the confounding issues of algal excretion, photolysis, inputs from groundwater, and leaching of benthic organic matter, permitting measurements that can not be made in situ. Increasing bioreactor residence time, EBCT, becomes a surrogate for DOC biological lability. The ¹³C-DOC tracer, derived from tulip poplar trees, is a natural product present in White Clay Creek. However, our goal was not to assess the importance of tulip poplar molecules to stream respiration, but rather the role of the natural, complex mixture of DOC molecules in the stream. The validity of our DOC lability profiling scheme rests on a primary assumption regarding the fidelity of the bioreactors as models of stream processes, i.e., that organic molecules of similar lability at the bioreactor scale will have similar lability in the stream. We assert that the ordering of DOC lability is preserved across systems with different spatial and temporal scales. This method can be used with molecular-level analyses to begin to provide lability information for individual molecules within the >10,000 molecule DOC pool in freshwaters.

