## Fractionation of carbon isotopes in Methanosarcina barkeri

<u>A.S. Bradley</u><sup>1</sup>, H. GROVER<sup>2</sup>, K. LONDRY<sup>2</sup>, AND R.E. SUMMONS<sup>1</sup>

<sup>1</sup> Department of Earth, Atmospheric, and Planetary Sciences, Massachusetts Institute of Technology; bradleya@mit.edu

<sup>2</sup> Department of Microbiology, University of Manitoba

The <sup>13</sup>C content of the organic constituents and products of an organism are controlled by the <sup>13</sup>C content of the growth susbstrate, the growth rate or degree of substrate utilization, the relative flux of carbon through various parts of the biochemical network, and the isotopic fractionation imposed by the enzymes of that network [1].

Methane is an important part of the Earth's carbon cycle, not least because of its role as a greenhouse gas. Despite this, there is little quantitative understanding of the various methane sources to the atmosphere and hydrosphere. These sources may include abiotic CO<sub>2</sub> reduction, thermogenic cracking of hydrocarbons, microbial production by Euryarchaeota, and direct CH<sub>4</sub> emissions by terrestrial plants [2]. Methane sources are often attributed on little more than  $\delta^{13}$ C values. However, the multitude of controls on  $^{13}$ C fractionation suggests that this approach is flawed.

We sought to examine the variability of the  $^{13}$ C content of the products of methanogens, including biomass, methane, and lipids (archaeol and *sn*-2 hydroxyarchaeol), as a function of growth substrate and rate. We chose *Methanosarcina barkeri* because of its ability to utilise various substrates and grew pure cultures on CO<sub>2</sub>/H<sub>2</sub>, acetate, trimethylamine, and methanol.

Growth without substrate limitation yielded a range of  $^{13}$ C discrimination factors, with growth on methanol yielding methane, biomass and lipids most depleted in  $^{13}$ C relative to substrate (77.6‰, 30.5‰, and 50.0‰ respectively) and growth on acetate yielding products least depleted (35‰, 4.6‰, and 5.7‰). Substrate-limited growth afforded smaller depletions in  $^{13}$ C on all substrates.

The compilation of these results allows us to better understand fractionation factors imposed by the various enzymes utilized in growth on different substrates. It also suggests that biological fraction of carbon in methane production can not be generalized to the common inference that biological methane has  $\delta^{13}C < -50\%$ . Carbon isotope fractionations imposed by methanogenic archaea can vary as strongly as those imposed by photoautotrophs, and under substrate-limiting conditions biological methane may be surprisingly enriched in <sup>13</sup>C.

## References

[1] Hayes, J.M. (2001) *Reviews in Mineralogy & Geochemistry* **42**, 225-227.

[2] Keppler, F., Hamilton, J.T., Brass, M., and Rockmann, T. (2006) *Nature* **439**, 187-191.