# Report number: DOE/97ER62479

Title: Cycling of DOC and DON by Novel Heterotrophic and Photoheterotrophic Bacteria in the Ocean

### Author and lead PI:

David Kirchman College of Marine and Earth Studies University of Delaware Lewes, DE 19958 302-645-4375 Kirchman@udel.edu

### Grant number: DE-FG02-97ER62479

DOE office: Office of Biological and Environmental Research

**STI Product Title:** Cycling of DOC and DON by Novel Heterotrophic and Photoheterotrophic Bacteria in the Ocean: Final Report

### Abstract:

The flux of dissolved organic matter (DOM) through aquatic bacterial communities is a major process in carbon cycling in the oceans and other aquatic systems. Our work addressed the general hypothesis that the phylogenetic make-up of bacterial communities and the abundances of key types of bacteria are important factors influencing the processing of DOM in aquatic ecosystems. Since most bacteria are not easily cultivated, the phylogenetic diversity of these microbes has to be assessed using culture-independent approaches. Even if the relevant bacteria were cultivated, their activity in the lab would likely differ from that under environmental conditions.

This project found variation in DOM uptake by the major bacterial groups found in coastal waters. In brief, the data suggest substantial differences among groups in the use of high and molecular weight DOM components. It also made key discoveries about the role of light in affecting this uptake especially by cyanobacteria. In the North Atlantic Ocean, for example, over half of the light-stimulated uptake was by the coccoid cyanobacterium, *Prochlorococcus*, with the remaining uptake due to *Synechococcus* and other photoheterotrophic bacteria. The project also examined in detail the degradation of one organic matter component, chitin, which is often said to be the second most abundant compound in the biosphere.

The findings of this project contribute to our understanding of DOM fluxes and microbial dynamics supported by those fluxes. It is possible that these findings will lead to improvements in models of the carbon cycle that have compartments for dissolved organic carbon (DOC), the largest pool of organic carbon in the oceans.

# **Publications Supported by DOE**

- Cottrell, M. T., and D. L. Kirchman. 2000a. Community composition of marine bacterioplankton determined by 16S rDNA clone libraries and fluorescence in situ hybridization. Applied and Environmental Microbiology 66: 5116-5122.
- Cottrell, M. T., and D. L. Kirchman. 2000b. Natural assemblages of marine proteobacteria and members of the *Cytophaga-Flavobacter* cluster consuming low- and high- molecularweight dissolved organic matter. Applied and Environmental Microbiology 66: 1692-1697.
- Cottrell, M. T., and D. L. Kirchman. 2003. Contribution of major bacterial groups to bacterial biomass production (thymidine and leucine incorporation) in the Delaware Estuary. Limnol. Oceanogr. 48: 168-178.
- Cottrell, M. T., and D. L. Kirchman. 2004. Single-cell analysis of bacterial growth, cell size, and community structure in the Delaware estuary. Aquat. Microb. Ecol. 34: 139-149.
- Cottrell, M. T., J. A. Moore, and D. L. Kirchman. 1999. Chitinases from uncultured marine microorganisms. Applied and Environmental Microbiology 65: 2553-2557.
- Cottrell, M. T., L. A. Waidner, L. Yu, and D. L. Kirchman. 2005a. Bacterial diversity of metagenomic and PCR libraries from the Delaware River. Environ. Microbiol. 7: 1883-1895.
- Cottrell, M. T., D. N. Wood, L. Y. Yu, and D. L. Kirchman. 2000. Selected chitinase genes in cultured and uncultured marine bacteria in the alpha- and gamma-subclasses of the proteobacteria. Applied and Environmental Microbiology 66: 1195-1201.
- Cottrell, M. T., L. Yu, and D. L. Kirchman. 2005b. Sequence and expression analyses of *Cytophaga*-like hydrolases in a western Arctic metagenomic library and the Sargasso Sea. Appl. Environ. Microbiol. 71: 8506-8513.
- Elifantz, H., R. R. Malmstrom, M. T. Cottrell, and D. L. Kirchman. 2005. Assimilation of polysaccharides and glucose by major bacterial groups in the Delaware Estuary. Appl. Environ. Microbiol. 71: 7799-7805.
- Elifantz, H., L. A. Waidner, M. T. Cottrell, and D. L. Kirchman. 2008. Diversity and Abundance of Glycosyl Hydrolase Family 5 in the North Atlantic Ocean. FEMS Microbiol. Ecol. 63: 316-327.
- Kirchman, D. L., and J. White. 1999. Hydrolysis and mineralization of chitin in the Delaware Estuary. Aquat. Microb. Ecol. 18: 187-196.
- Michelou, V. K., M. T. Cottrell, and D. L. Kirchman. 2007. Light-stimulated bacterial production and amino acid assimilation by cyanobacteria and other microbes in the North Atlantic Ocean. Appl. Environ. Microbiol. 73: 5539-5546.
- Svitil, A. L., S. M. N. Chadhain, J. A. Moore, and D. L. Kirchman. 1997. Chitin degradation proteins produced by the marine bacterium Vibrio harveyi growing on different forms of chitin. Applied and Environmental Microbiology 63: 408-413.

#### **Postdoctorate and Research Fellows**

Amy Svitil Matthew Cottrell

# Graduate Students Fully or Partially Supported by this DOE project

- 1. Jessica Moore, M.S. 1999
- 2. Hila Elifantz, Ph.D. 2007
- 3. Vanessa Michelou, Ph.D., projected graduation in 2009
- 4. Lisa Waidner, Ph.D., 2007

Two other students, Rex Malmstrom (Ph.D., 2005) and Tiffany Straza (Ph.D. expected in 2009), were not supported by this project, but benefited from it and helped out in many aspects.

#### **Undergraduate Students**

This research grant support summer intern projects by undergraduate students from Lincoln University, the nation's oldest historical Black college. One or two students were supported each summer during the project.