

Potential impact of the interactions between *Synechococcus* and heterotrophic bacteria on oceanic carbon flow

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Abstract: *Synechococcus* accounts for a considerable proportion of marine primary production and has complex interactions with heterotrophic bacteria at sub-micrometer scales. However, the impact of the *Synechococcus*-bacteria interactions on marine carbon cycle is still unclear. Here, through laboratory experiments, we studied the potential role of the interactions between bacterial communities and two ecotypes of *Synechococcus* (i.e. PCC7002 and CCMP1334) in driving carbon flow. *Synechococcus* released abundant dissolved organic matters (DOM) during their growth. A large fraction of the DOM was rapidly utilized by heterotrophic bacteria and transformed into bacterial biomass. While partly since the DOM released from two *Synechococcus* had different chemical composition, which can be partially reflected by their differences in the fluorescent DOM (FDOM) components, the two *Synechococcus* reshaped the co-cultivated bacterial community into distinct community structure. Among the FDOM released by these two *Synechococcus* ecotypes, a portion of humic-like component was highly resistant to bacterial degradation and could accumulate in the co-culture system, suggesting that *Synechococcus* may directly produce recalcitrant DOM and contribute to the long-term sequestration of carbon in a dissolved form in the ocean. Meanwhile, the *Synechococcus*-bacteria interactions increased aggregate formation and particle sinking, suggesting that their interactions in the ocean may enhance the export of *Synechococcus*-derived organic carbon to deep sea through sinking process of biological pump.

Key words: Marine *Synechococcus*, bacteria, carbon export, carbon sequestration