

Heterotrophic sources for dissolved organic nitrogen in the oligotrophic ocean indicated by nitrogen isotopic analysis of individual amino acids

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Dissolved organic nitrogen (DON) plays the roles in N cycling and ecosystem in the upper ocean. This study explores the use of nitrogen isotopic analysis of individual amino acids ($\delta^{15}\text{N}_{\text{AA}}$) of coupled DON and particulate organic nitrogen (PON) samples as a new approach to examine relative sources, transformation processes, and potential coupling of the main detrital organic nitrogen form in the ocean water column. We measured $\delta^{15}\text{N}_{\text{AA}}$ distributions in high-molecular-weight DON (HMW DON) and suspended PON in the North Pacific Subtropical Gyre (NPSG) from surface to mesopelagic depths. A new analytical approach using HPLC purification of amino acids achieved far greater $\delta^{15}\text{N}_{\text{AA}}$ measurement precision for DON than earlier work, allowing us to resolve previously obscured differences in $\delta^{15}\text{N}_{\text{AA}}$ signatures, both with depth and between ON pools. The $\delta^{15}\text{N}_{\text{AA}}$ values of both surface and mesopelagic HMW DON suggest mainly heterotrophic sources, with the mesopelagic HMW DON bearing signatures of far more degraded material. These results contrast with a previous proposal that DON $\delta^{15}\text{N}_{\text{AA}}$ patterns are essentially “pre-formed” in the surface ocean, undergoing little further change with depth. Together with the results of amino acid enantiomers (D/L), these results suggest that heterotrophic bacteria are the main sources for DON at both surface and mesopelagic depths in the oligotrophic ocean.