



## Transportation and Transformation of Dissolved Organic Matter from Overlying to Bottom Waters of Cold Seeps in the South China Sea: Insights at the Molecular Level

**Shi Tang**<sup>1</sup>, Zhenwei Yan<sup>4</sup>, Yuanbi Yi<sup>4</sup>, Yuan Shen<sup>5</sup>, Wei Xie<sup>1,2</sup>, Ding He<sup>4,6</sup>, and Penghui Li<sup>1,2,3</sup>

<sup>1</sup>School of Marine Sciences, Sun Yat-sen University, Zhuhai, China

<sup>2</sup>Southern Marine Science and Engineering Guangdong Laboratory, Zhuhai, China

<sup>3</sup>Guangdong Provincial Key Laboratory of Marine Resources and Coastal Engineering, Zhuhai, China

<sup>4</sup>Department of Ocean Science and Center for Ocean Research in Hong Kong and Macau, The Hong Kong University of Science and Technology, Hong Kong, China

<sup>5</sup>State Key Laboratory of Marine Environmental Science & College of Ocean and Earth Sciences, Xiamen University, Xiamen, China

<sup>6</sup>State Key Laboratory of Marine Pollution, City University of Hong Kong, Hong Kong, Hong Kong SAR, China

Cold seeps are critical hotspots in marine ecosystems, where the biogeochemical processes of dissolved organic matter (DOM) significantly impact regional carbon reservoirs and the global ocean carbon cycle. To clarify the impact of cold seep activity on the production, transportation and transformation of DOM, we employed Fourier transform ion cyclotron resonance mass spectrometry (FT-ICR MS) to analyze DOM from the water column and sediment overlying water collected from cold seep and non-cold seep regions in the northern South China Sea. Our results showed that the overlying water in cold seeps contained a greater diversity of unique molecules, with a larger proportion of sulfur-containing compounds compared to the non-cold seep area. Approximately half of these unique molecules, characterized by lower H/C ratios, higher molecular weights, and a predominance of highly unsaturated compounds (82.3%), were transferred to the corresponding bottom water during the bubbling process. In contrast, molecules with higher H/C ratios, lower molecular weights, and a larger proportion of aliphatics compounds (40.8%) were lost. Additionally, the bottom water of the active cold seep exhibited the formation of some labile molecules ( $H/C > 1.5$ ) with lower aromaticity ( $AI_{mod} < 0.25$ ) and the decomposition of nitrogen-containing carboxyl-rich alicyclic molecules (CRAMs) with higher aromaticity, driven by the positive priming effect. These findings highlight the profound influence of cold seep activity on DOM properties and dynamics, providing deeper insights into the complex biogeochemical processes in cold seep ecosystems and their critical implications for marine carbon cycling.