

Active methane cycling studied by molecular biomarkers and stable isotope in the sediment of Sakhalin continental slope

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Active methane (CH₄) releases from gas seepages bearing gas hydrate (GH) to water column have been widely observed on the Sakhalin continental slope. However biogeochemical processes about methane-related archaea communities in this region is poorly understood. Hence, we analyzed the distribution of the intact polar lipid (IPL) and core lipid (CL)-glycerol dibiphytanyl glycerol tetraethers (GDGTs) in order to assess its utility as a proxy for *in situ* CH₄ production/consumption in geological systems associated with shallower GH accumulation. In shallow sediments, the most depleted-carbon isotopic values of CH₄ and CO₂ ($\delta^{13}\text{C}_{\text{CH}_4}$: -46.3‰ and $\delta^{13}\text{C}_{\text{CO}_2}$: -26.6‰) indicate that thermogenic CH₄ is mainly oxidized by methanotrophic activities. $\delta^{13}\text{C}$ values of biphytanes ($\delta^{13}\text{C}_{\text{BP1}}$ as low as -69.3‰) derived from IPL-GDGTs, in particular BP-1 as probable methane-related biomarker, are distinctively lighter than those of CH₄, suggesting that methanotrophic archaea is potentially indicative for living biomass performs *in situ* anaerobic oxidation of methane (AOM) in this depth range. Previous 16S rRNA surveys of the microbial populations in this vent deposit have shown predominant ANME-1 among phylogenetically *Euryarchaeota* and appear to be controlled by geochemical and geological processes that drive biogeochemical CH₄ cycle. Overall, IPL results are similar with gene-based survey. We conclude that upward CH₄ sources from thermogenic origin can be continuously reduced by methanotrophic activities in surface seafloor. This study provides the first biogeochemical data set of active CH₄ vent deposits at the Sakhalin continental slope, which help to better understand modern and past CH₄ cycle in this setting.