Mercury methylation and demethylation in paddy soil with and without rice plants

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Methylmercury (MeHg) is a neurotoxin that can accumulate in rice via uptake from the paddy soil into the plant. Mercury methylation and demethylation are microbial processes that control the concentration of MeHg ([MeHg]) in paddy soil, although their relative influence may differ between vegetated and non-vegetated soils. This study is the first to assess potential rates of methylation and demethylation (Kmeth and Kdemeth) simultaneously in a manner that explicitly compares how they affect [MeHg] in the vegetated and non-vegetated paddy soil compartments. Flooded rice paddy soil microcosms (2x10x20") with and without rice plants were amended with enriched Hg stable isotopes (200Hg and Me201Hg) to assess Kmeth and Kdemeth. Overall, the results of this study suggest that MeHg concentrations are differentially controlled by MeHg production and degradation processes, depending on whether plants are present. In non-vegetated soils, Kmeth was positively correlated with both ambient [MeHg] (p = 0.02, r = 0.90) and the percent of total Hg present as MeHg (%MeHg; p = 0.023, r = 0.87), while K_{demeth} had no relationship (p = 0.77-79), indicating that MeHg formation drove MeHg dynamics. In planted soils, by contrast, Kmeth did not correlate to [MeHg] (p = 0.79). Rather, K_{demeth} was negatively correlated with both [MeHg] (p = 0.04, r = -0.82) and %MeHg (p = 0.036, r = -0.84), suggesting that degradation of MeHg was a more important control on the [MeHg] of vegetated soils.