## Anomalous uptake of CH<sub>4</sub> in volcanic soils indicative of a novel sink mechanism for atmospheric CH<sub>4</sub>

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There is a great diversity of natural and anthropogenic sources of atmospheric methane (CH<sub>4</sub>) but only three known sinks: oxidation by hydroxyl radicals in the troposphere, stratospheric loss, and oxidation by soil bacteria. To date, no physical processes have been reported that sequester atmospheric CH<sub>4</sub>, however, sorption of CH<sub>4</sub> to mineral surfaces in marine sediment is common [1] and CH<sub>4</sub> adsorption onto selected nanoporous zeolites has been modelled [2].

We conducted a <sup>13</sup>CH<sub>4</sub> stable isotope switching (SIS) [3] study on Hawaiian soils to investigate the effects of soil weathering and development on atmospheric CH<sub>4</sub> uptake. Results indicated an anomalous sink mechanism for atmospheric CH<sub>4</sub> that appears to be most likely linked with long term abiotic adsorption of atmospheric CH<sub>4</sub> under Earth surface conditions (temperature and pressure). Capture and sequestration of CH<sub>4</sub> was reversible, and adsorbed CH<sub>4</sub> bio-available; demonstrated by subsequent CH<sub>4</sub> oxidation by high affinity methanotrophic bacteria in younger soils (<150 ka).

We suggest the composition of the colloidal fraction (<2  $\mu$ m) of a soil impacts CH<sub>4</sub> sink capacity beyond the direct influence that physical structure has on soil gas diffusion. The type, activity, and surface area of soil minerals appears to exert a direct control on CH<sub>4</sub> uptake, storage and supply, prior to CH<sub>4</sub> access by soil methanotrophs. This additonal uptake mechanism increases the potential microbial CH<sub>4</sub> sink and explains elevated CH<sub>4</sub> uptake capacities reported from minearologically similar andisols in Tenerife [4].

 [1] Ertefai et al. (2010) Geochim. Cosmochim. Acta. 74, 6033–6048, [2] Kim et al. (2013) Nat. Commun. 4, 1694, [3] Maxfield et al. (2012) Rapid Commun. Mass. 26, 997-1004, [4] Maxfield et al. (2008) Environ, Microbiol. Rep. 1, 450-456.