

Biodiversity of methanogenic Archaea in permafrost affected soils of the Lena Delta, Siberia

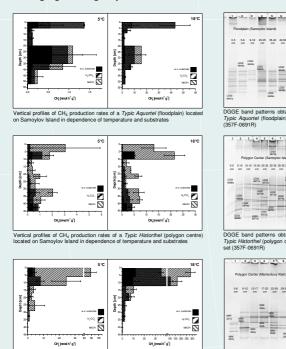
Introduction

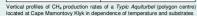
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Hydromorphic arctic tundra soils are an important source of the greenhouse gas methane (CH₄). In these environments carbon is accumulated due to reduced decomposition of organic matter. Most of the climate models predict a global warming for the next century which will lead in deeper and longer thaw processes in the active layer of arctic permafrost soils. Consequently a higher emission of methane and carbon dioxide could be expected. The release of CH₄ is a result of strictly anaerobic methanogenic archaea that use simple compounds (e.g. H₂, CO₂, acetate) formed by bacteria during the anaerobic degradation of organic material. There were only a few investigations about the methanogenic community existing in cold terrestrial habitats (e.g. Høj et al., 2005). Here we investigated the methanogenic community solution tundra soils located in the Laptev Sea coast region (Siberia).

Results

Independent from the chosen temperature or the added substrate the methane production rate reached its maximum in nearly all soil profiles in the upper soil layers. With rising temperature and/or the adding of substrate the methane production rate increase. In all soils we found sequences of *Methanomicrobiales* and *Methanosarcina* but no members of *Methanobacteriales*. In two of the soils we also found sequences belonging to the group of *Methanosaetaceae*.

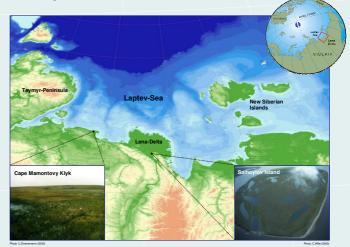


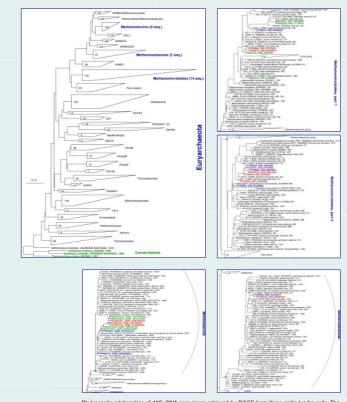


Conclusions

In all three investigated permafrost soils methanogens could be detected by activity tests and molecular methods. Sequences of *Methanosaetaceae* indicating the presence of methanogens that use acetate as the only carbon and energy source. Most of the described *Methanomicrobiales* use mainly H_2/CO_2 or formate for methanogenesis and growth while the genus *Methanosarcina* can use a wide variety of substrates. With increasing depth the decrease of DNA bands in DGGE

Investigation Area





Phylogenetic relationships of 155 rDNA sequences retrieved by DGGE from three arctic tundra solis. The sequences obtained in this study are coloured. FP – Floodplat asmoylov Island, PC - Polyon Centre Samoylov Island, MAK – Polygon Centre Cape Mamontovy Klyk. The scale bar represents 0.1 changes per nucleotide.

patterns could be explained by the decrease of temperature and/or the availibility of substrate. Those bands could represent psychrotolerant or psychrophilic methanogenic archaea that are well adapted to these low temperature conditions near the permafrost table. This is supported by the results of the activity tests that show a distinct CH_4 production at a temperature of 5 °C.

References

Hej et al. (2005) Archaeal communities in High Arctic wetlands at Spitsbergen, Norway (78°N) as characterized by 16S rRNA gene fingerprinting. FEMS Microbiology Ecology 53, 89-101 Alfred-Wegener-Institute for Polar and Marine Research Forschungsstelle Potsdam Telegrafenberg A 43 14473 Potsdam, Germany