



Distinct microbial communities associated with buried soils in the Siberian tundra

Antje Gittel (1,2), Jiří Bárta (3), Iva Kohoutová (3), Robert Mikutta (4), Sarah Owens (5), Jack Gilbert (5), Jörg Schneckner (6,2), Birgit Wild (6,2), Bjarte Hannisdal (7), Joeran Maerz (8), Nikolay Lashchinskiy (9), Petr Čapek (3), Hana Šantrůčková (3), Norman Gentsch (4), Olga Shibistova (4), Georg Guggenberger (4), Andreas Richter (2,6), Vigdis Torsvik (1), Christa Schleper (2,10), Tim Urich (2,10)

(1) Centre for Geobiology, Department of Biology, University of Bergen, Bergen, Norway, (2) Austrian Polar Research Institute, Vienna, Austria, (3) Department of Ecosystems Biology, University of South Bohemia, České Budějovice, Czech Republic, (4) Institut für Bodenkunde, Leibniz Universität Hannover, Hannover, Germany, (5) Institute of Genomics and Systems Biology, Argonne National Laboratory, Argonne, IL, USA, (6) Division of Terrestrial Ecosystem Research, Department of Microbiology and Ecosystem Science, University of Vienna, Vienna, Austria, (7) Centre for Geobiology, Department of Earth Science, University of Bergen, Bergen, Norway, (8) Division of Ecosystem Modelling, Institute of Coastal Research, Helmholtz Zentrum Geesthacht, Geesthacht, Germany, (9) Central Siberian Botanical Garden, Siberian Branch of Russian Academy of Sciences, Novosibirsk, Russia, (10) Division of Archaea Biology and Ecogenomics, Department of Ecogenomics and Systems Biology, University of Vienna, Vienna, Austria

Cryoturbation, the burial of topsoil material into deeper soil horizons by repeated freeze–thaw events, is an important storage mechanism for soil organic matter (SOM) in permafrost-affected soils. Besides abiotic conditions, microbial community structure and the accessibility of SOM to the decomposer community are hypothesized to control SOM decomposition and thus have a crucial role in SOM accumulation in buried soils. We surveyed the microbial community structure in cryoturbated soils from nine soil profiles in the northeastern Siberian tundra using high-throughput sequencing and quantification of bacterial, archaeal and fungal marker genes. We found that bacterial abundances in buried topsoils were as high as in unburied topsoils. In contrast, fungal abundances decreased with depth and were significantly lower in buried than in unburied topsoils resulting in remarkably low fungal to bacterial ratios in buried topsoils. Fungal community profiling revealed an associated decrease in presumably ectomycorrhizal (ECM) fungi. The abiotic conditions (low to subzero temperatures, anoxia) and the reduced abundance of fungi likely provide a niche for bacterial, facultative anaerobic decomposers of SOM such as members of the Actinobacteria, which were found in significantly higher relative abundances in buried than in unburied topsoils. Our study expands the knowledge on the microbial community structure in soils of Northern latitude permafrost regions, and attributes the delayed decomposition of SOM in buried soils to specific microbial taxa, and particularly to a decrease in abundance and activity of ECM fungi, and to the extent to which bacterial decomposers are able to act as their functional substitutes.