



Supplement of

Metagenomic analyses of the late Pleistocene permafrost – additional tools for reconstruction of environmental conditions

E. Rivkina et al.

Correspondence to: E. Rivkina (elizaveta.rivkina@gmail.com)

The copyright of individual parts of the supplement might differ from the CC-BY 3.0 licence.

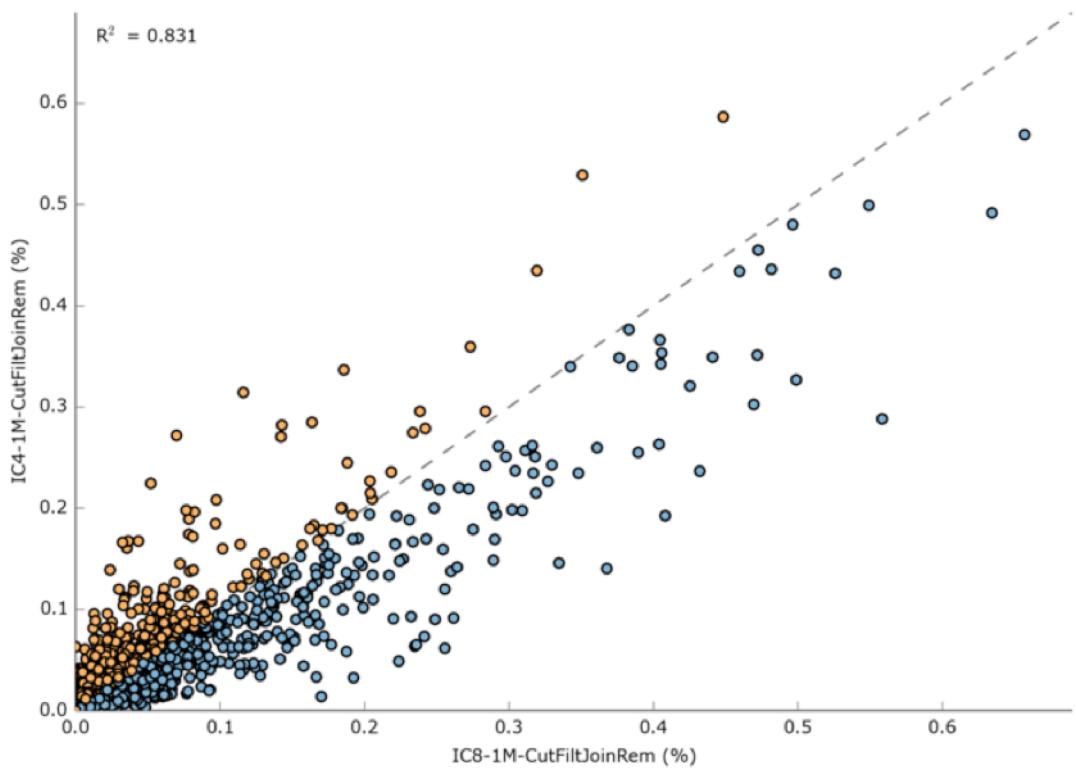


Figure S1. Profile scatter plot indicating the similarity of the metagenomes based on the SEED functional annotation.