

**Table S3:** The detection of archaeal sequences in environmental samples at resolution down to the family level. The classification was conducted on quality filtered and chimera free sequence libraries for each sample. Green boxes indicate sequence classes that were detected.

Classification	ANBR	FAS	GC	OP-YNP
Crenarchaeota				
Thermoprotei				
Caldisphaerales				
Caldisphaeraceae				
Desulfurococcales				
Desulfurococcaceae				
Sulfolobales				
Sulfolobaceae				
Thermoproteales				
Thermoproteaceae				
unclassified				
env_sample				
Euryarchaeota				
Archaeoglobi				
Archaeoglobales				
Archaeoglobaceae				
Marine_Group_II				
env_sample				
Marine_Group_III				
env_sample				
Methanobacteria				
Methanobacteriales				
Methanobacteriaceae				
Methanomicrobia				
Methanomicrobiales				
Genera_incertae_sedis				
Methanospirillaceae				
Methanosarcinales				
Methanosaetaceae				
Methanosarcinaceae				
Thermoplasmata				
Thermoplasmatales				
Picrophilaceae				
Nanoarchaeota				
env_sample				
Thaumarchaeota				
marine_archaeal_group_1				
Nitrosopumilales				
Nitrosopumilaceae				
env_sample				

env\_sample  
unclassified
