

Table S1. PCR primers and conditions for screening of archaeal single amplified genomes (SAGs).

Primer	Sequence (5' → 3')	Annealing temp (°C)	Gene	Metabolic process	Reference
SSU rRNA PCR primers					
parch_519F	CAGCMGCCGCGGTAA	50	SSU rRNA	NA	Ovreas <i>et al.</i> , 1997
Arch_915R	GTGCTCCCCCGCCAATTCCT				DeLong, 1992
Arch_345F	CCTAYGGGGYGCASCAG	57	SSU rRNA	NA	Gantner <i>et al.</i> , 2010
Arch_1000R	GGCCATGCACYWCYTCTC				
Metabolic gene PCR primers					
Arch-amoAF	STAATGGTCTGGCTTAGACG	50	<i>amoA</i>	Ammonia oxidation	Francis <i>et al.</i> , 2005
Arch-amoAR	GCGGCCATCCATCTGTATGT				
NirKMF	CCWGGWCCAACHYTDAGA	50	<i>nirK</i>	Nitrite reduction	This study
NirkMR	ATTTCWCCDACAAYRTG				
nifHF	GGHAARGGHGGHATHGGNAARTC	55	<i>nifH</i>	Nitrogen fixation	Mehta <i>et al.</i> , 2003
nifHR	GGCATNGCRAANCCVCCRCANAC				

References:

- Ovreas L, Forney L, Daae F, Torsvik V (1997) Distribution of bacterioplankton in meromictic Lake Saelenvannet, as determined by denaturing gradient gel electrophoresis of PCR-amplified gene fragments coding for 16S rRNA. *Appl Environ Microbiol* 63: 3367-3373.
- DeLong EF (1992) Archaea in coastal marine environments. *Proc Natl Acad Sci USA* 89: 5685-5689.
- Gantner S, Andersson AF, Alonso-Sáez L, Bertilsson S (2010) Novel primers for 16S rRNA-based archaeal community analyses in environmental samples. *J Microbiol Methods* 84: 12-18.
- Francis CA, Roberts KJ, Beman JM, Santoro AE, Oakley BB (2005) Ubiquity and diversity of ammonia-oxidizing archaea in water columns and sediments of the ocean. *Proc Natl Acad Sci USA* 102: 14683-14688.
- Mehta MP, Butterfield DA, Baross JA (2003) Phylogenetic diversity of nitrogenase (*nifH*) genes in deep-sea and hydrothermal vent environments of the Juan de Fuca Ridge. *Appl Environ Microbiol* 69: 960-970.