

Supplementary Table S2 Primers and PCR conditions of target genes for qPCR

Gene	Primer set	Primer sequence (5'-3')	Thermal profile	References
<i>nirK</i>	F1aCu	ATCATGGTSC TGCCGCG	98 °C for 2 min; 30 cycles of 98 °C for 30 s, 58°C for 30 s, and 72 °C for 30 s	[12]
	R3Cu	GCCTCGATCAGRTTGTGGTGGTT		
<i>nirS</i>	Cd3aF	GTSAACGTSAAGGARACSGG	98 °C for 2 min; 30 cycles of 98 °C for 30 s, 55°C for 30 s, and 72 °C for 30 s	[13]
	R3cdR	GASTTCGGRTGSGTCTTGA		
<i>hzsB</i>	<i>hzsB</i> 396F	ARGGHTGGGGHAGYTGGAAG	3 min at 95 °C, 40 cycles consisting of 30 s at 95 °C, 30 s at 59°C and 30 s at 72 °C	[14]
	<i>hzsB</i> 742R	GTYCCHACRTCATGVGTCTG		

References

- [12] Hallin S, Lindgren P. 1999. PCR detection of genes encoding nitrite reductase in denitrifying bacteria. *Applied and Environmental Microbiology* 65: 1652-1657 <https://doi.org/10.1128/aem.65.4.1652-1657.1999>
- [13] Kandeler E, Deiglmayr K, Tscherko D, Bru D, Philippot L. 2006. Abundance of *narG*, *nirS*, *nirK*, and *nosZ* genes of denitrifying bacteria during primary successions of a glacier foreland. *Applied and Environmental Microbiology* 72: 5957-5962 <https://doi.org/10.1128/aem.00439-06>
- [14] Wang S, Zhu G, Peng Y, Jetten M, Yin C. 2012. Anammox bacterial abundance, activity, and contribution in riparian sediments of the Pearl River Estuary. *Environmental Science & Technology* 46: 8834-8842 <https://doi.org/10.1021/es3017446>