

Supplementary data

Table S1 Characteristics of domestic wastewater

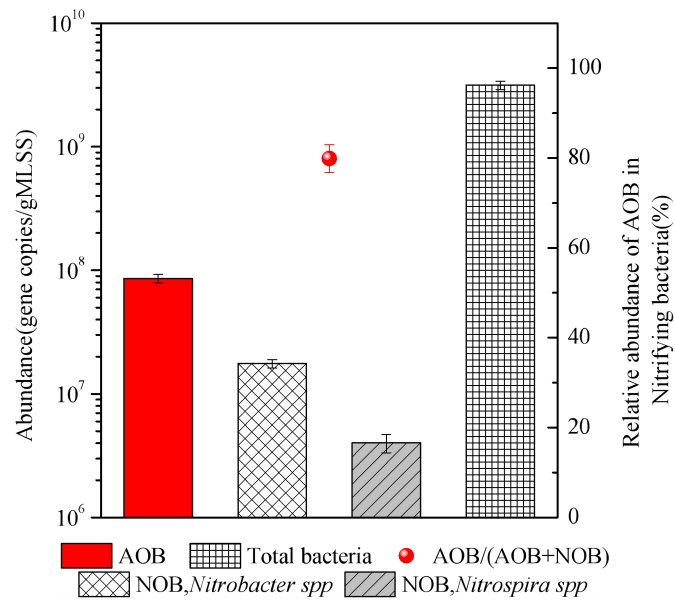
Characteristics	Abbr.	Unit	Range	Average
pH	pH	–	7.3–7.7	–
Chemical oxygen demand	COD	mg/L	167–223	195
Ammonium nitrogen	NH ₄ ⁺ -N	mg/L	54.8–75.6	65.2
Nitrite nitrogen	NO ₂ ⁻ -N	mg/L	0.08–0.79	0.32
Nitrate nitrogen	NO ₃ ⁻ -N	mg/L	0–0.87	0.21
Total nitrogen	TN	mg/L	55.4–88.3	67.1

Table S2 Temperature coefficient (θ) from the literature and this study

MLVSS (g/L)	T (°C)	θ	Ea (KJ/mol)	Conditions	Culture	Source
–	10–25	1.12	–	–	–	Henze et al. (2000)
0.15–0.2	10–20	1.098	–	Domestic wastewater	Activated sludge	Sözen et al. (1996)
4	20–35	1.045	–	Leather-tanning wastewater	Activated sludge	Gorgun et al. (2007)
0.43	4–33	1.028	–	Digester	Activated	Shammas (1986)
1.2	4–25	1.061	–	supernatant	sludge	
3.2	4–25	1.129	–	–	–	–
1.1		1.122 ± 0.010	74.9 ± 6.6	–	–	–
1.8		1.111 ± 0.006	71.8 ± 3.9	–	–	–
3.8	10–30	1.082 ± 0.008	58.4 ± 5.4	Domestic wastewater	Partial nitrification sludge	This study
5.3		1.084 ± 0.005	54.5 ± 2.3			
6.8		1.072 ± 0.004	51.3 ± 3.5			
8.7		1.078 ± 0.005	49.4 ± 5.8			
1.5–2	5–20	1.172	111.5	Domestic sewage	Partial nitrification sludge	Guo et al. (2010)
	20–35	1.062	42.0			
1.6–1.8	10–20	–	119.7	Synthetic wastewater	Partial nitrification sludge	Wu et al. (2012)
	20–30	–	54.3			
0.9–1.1	12–34	–	24.6	Synthetic wastewater	Activated sludge	Weon et al. (2004)
	4–12	–	253			
–	10–20	–	163	Synthetic wastewater	Immobilized cells	Benyahia and Polomarkaki (2005)
–	20–35	–	56			

Table S3 The primers targeting AOB, NOB (*Nitrobacter spp* and *Nitrospira spp*), and total bacteria

Specificity	Primer	Sequence (5'-3')	Reference
<i>amoA</i> (AOB)	<i>amoA</i> -1f	GGGGTTTCTACTGGTGGT	Wang et al.
	<i>amoA</i> -2r	CCCCTCKGSAAGCCTTCTTC	(2011)
16S rRNA gene for <i>Nitrospira spp.</i>	341f	CCTACGGGAGGCAGCAG	Geets et al.
	NTSPAr	CGTTATCCTGGGCAGTCCTT	(2007)
16S rRNA gene for <i>Nitrobacter spp.</i>	FGPS872f	CTAAAACTCAAAGGAATTGA	Ahn et al.
	FGPS1269r	TTTTTTGAGATTTGCTAG	(2008)
16S rRNA gene for the total bacteria	Amx820r	AAAACCCCTCTACTTAGTGCCC	
	341f	CCTACGGGAGGCAGCAG	Koike et al.
	534r	TTACCGGGCTGCTGGCAC	(2007)

**Fig. S1** Abundance of total bacteria, AOB and NOB (*Nitrobacter spp* and *Nitrospira spp*) in the pilot-scale SBR activated sludge. Error bars indicate standard deviation, $n = 3$

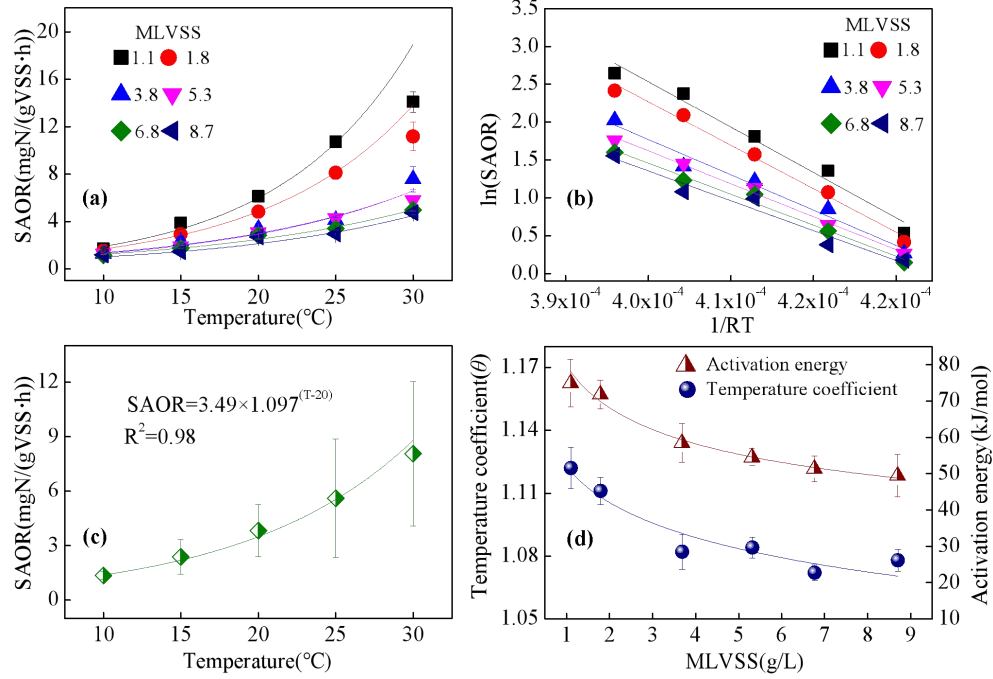


Fig. S2 Effect of MLVSS on the temperature coefficient and activation energy. (a, represents the relationships between SAOR and temperature under different MLVSS; b, represents the relationships between ln(SAOR) and 1/RT under different MLVSS; c, represents the relationships between mean SAOR of different temperatures at 10 °C, 15 °C, 20 °C, 25 °C, 30 °C and temperature; d, represents the relationships between MLVSS and temperature coefficient/activation energy) (Error bars indicate standard deviations of measurements)

Detailed analysis method

Quantification of nitrifiers

The sludge samples were taken from the pilot-scale SBR parent reactor. Using FastDNA SPIN kits for soil (Q BIOgene Inc, CA, USA), DNA of sludge samples were extracted. DNA concentrations were estimated using Nanodrop 1000 Spectrophotometer (Nanodrop, DE, USA). Using the SYBR-Green real-time PCR, three genes targeting AOB, NOB (*Nitrobacter spp* and *Nitrospira spp*) and total bacteria were quantified (Table S3). The measurement was according to the previous report Liu et al. (2017). The results of efficiency and correlation coefficient above 95% and 0.98 were employed.

References

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