

1 **Supplementary material**

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3 Effect of elevated nitrate and sulfate concentrations on selenate

4 removal by mesophilic anaerobic granular sludge bed reactors

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19 **Method for FISH imaging**

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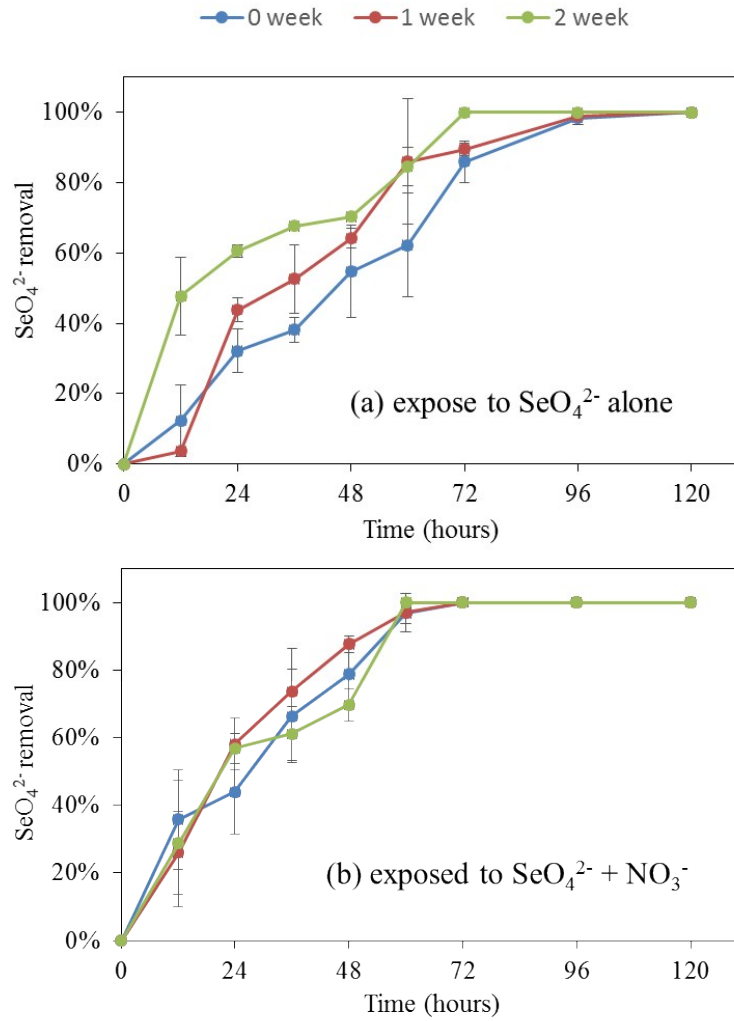
21 The microbial communities present in the sludge samples were qualitatively analyzed using  
22 fluorescence in situ hybridization (FISH). Vigorous biomass mixing within the reactor was  
23 done first by increasing the liquid upflow velocity to ensure a good representation of biomass  
24 sampling. All microbial cells were visualized by DAPI staining with specific oligonucleotide  
25 probes ARCH915 <sup>1</sup> and EUB I-III <sup>2</sup> for archaeal and bacterial cells, respectively. Two  
26 subgroups of the proteobacteria (beta-proteobacteria and gamma-proteobacteria) were included  
27 using BET42a and GAMMA42a probes. <sup>3</sup> Sulfate-reducing bacteria (SRB) were visualized  
28 using the *Desulfobulbaceae* DBB60 <sup>1</sup> and *Desulfovibrionales* SRB385 <sup>4</sup> probes. The inoculum  
29 anaerobic granular sludge was used as the control.

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31 **Table S1** Batch test results with standard deviation after 5 days incubation under different molar ratios of  $\text{NO}_3^-$  and  $\text{SO}_4^{2-}$  to  $\text{SeO}_4^{2-}$ .

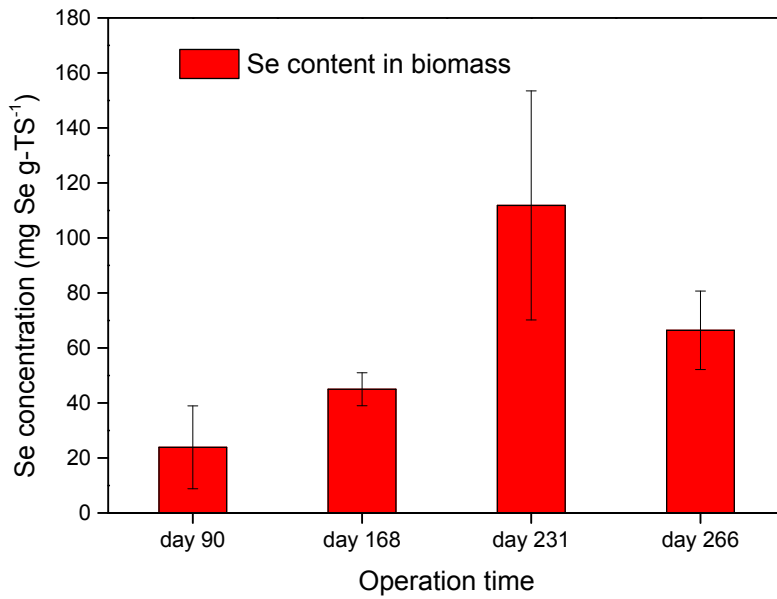
Condition/molar ratio	Lactate		* $\text{NO}_3^-$		$\text{SO}_4^{2-}$		TDS	$\text{SeO}_4^{2-}$		
	Initial conc. (mM)	% Removal	Initial conc. (mM)	% Removal	Initial conc. (mM)	% Removal	Produced (mM)	Initial conc. ( $\mu\text{M}$ )	% Removal	
Sterilized biomass	21±1	0±0	2±1	0±0	3±1	0±0	0.0±0.0	179±0	0±0	
No lactate added	n/a***		2±0	0±0	3±0	0±0	0.0±0.0	163±1	0±0	
$\text{NO}_3^-$ only	12±1	100±0	4±0	100±0	n/a	n/a	n/a	n/a	n/a	
$\text{SO}_4^{2-}$ only	14±1	100±0	n/a	n/a	25±1	15±1	4.0±0.0	n/a	n/a	
$\text{NO}_3^- + \text{SO}_4^{2-}$	14±0	100±0	4±1	100±0	24±0	6±2	3.0±0.0	n/a	n/a	
$\text{SeO}_4^{2-}$ only	25±4	83±0	n/a	n/a	n/a	n/a	n/a	108±0	67±2	
** $\text{NO}_3^- / \text{SeO}_4^{2-}$	1	13±0	100±0	0.5±0.0	100±0	n/a	n/a	n/a	410±0	62±12
	20	17±2	85±3	2±0	100±0	n/a	n/a	n/a	96±0	100±0
	40	17±2	85±3	4±0	100±0	n/a	n/a	n/a	141±0	91±12
	70	19±2	90±1	7±0	100±0	n/a	n/a	n/a	117±0	81±10
	100	18±)	92±1	10±0	100±0	n/a	n/a	n/a	98±0	59±4
$\text{SO}_4^{2-} / \text{SeO}_4^{2-}$	1	13±0	91±2	n/a	n/a	0.5±0.0	63±9	0.3±0.2	410±0	71±4
	50	24±4	83±2	n/a	n/a	4±0	39±10	0.5±0.2	236±0	85±0
	100	24±5	92±0	n/a	n/a	9±1	28±1	1.0±0.2	237±0	79±1
	150	24±5	86±1	n/a	n/a	14±1	23±1	1.0±0.2	194±0	40±0
	200	24±3	92±1	n/a	n/a	20±1	25±7	0.9±0.3	235±0	52±0
	300	22±4	89±1	n/a	n/a	28±1	16±1	1.7±0.9	247±0	39±1
$\text{SO}_4^{2-} / \text{SeO}_4^{2-}$ with co-exposure of 4 mM $\text{NO}_3^-$	1	13±0	92±1	0.5±0.0	100±0	0.5±0.0	27±4	0.1± 0.0	500±0	100±0
	50	25±3	86±0	3±1	100±0	4±0	30±2	0.3±0.1	201±0	92±1
	100	24±5	84±1	3±1	100±0	9±1	25±5	0.7±0.3	197±0	91±1
	150	24±4	86±0	3±1	100±0	14±1	25±6	0.6±0.2	217±0	61±2
	200	24±5	92±1	3±1	100±0	19±1	20±4	0.4±0.0	240±)	61±2
	300	24±5	92±1	3±1	100±0	31±1	19±3	0.5±0.1	221±0	32±0

Note: results are all reported as mean ± standard deviation, n = 4 replicates (total); \* $\text{NO}_3^-$  concentration reached 100% removal for all concentration at 48 h; n/a – not applicable



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34 **Fig. S1**  $\text{SeO}_4^{2-}$  reduction using untreated sludge and treated sludge (reused from the previous  
 35 week) (a) expose only to  $\text{SeO}_4^{2-}$  and (b) expose to both  $\text{NO}_3^-$  and  $\text{SeO}_4^{2-}$ . Batch experiments  
 36 were conducted under excess lactate (10mM), 30°C, 200 rpm and pH 7.0.

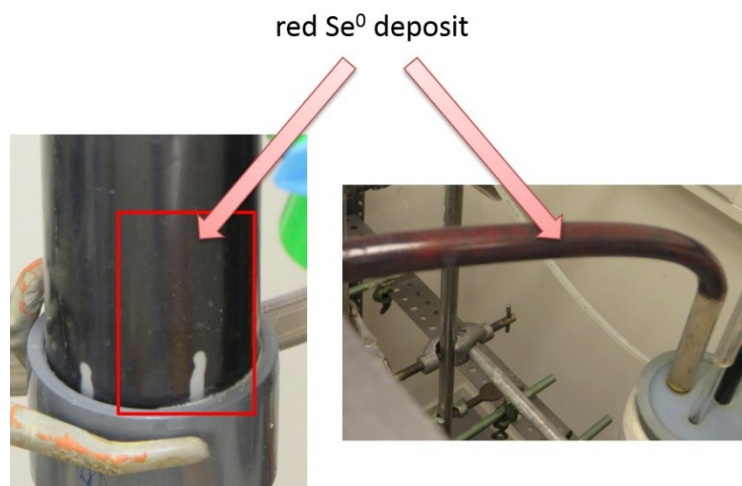


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38 **Fig. S2** Se concentration detected in the biomass at various operation times (n=3 biomass  
 39 sample replicate at each time period).

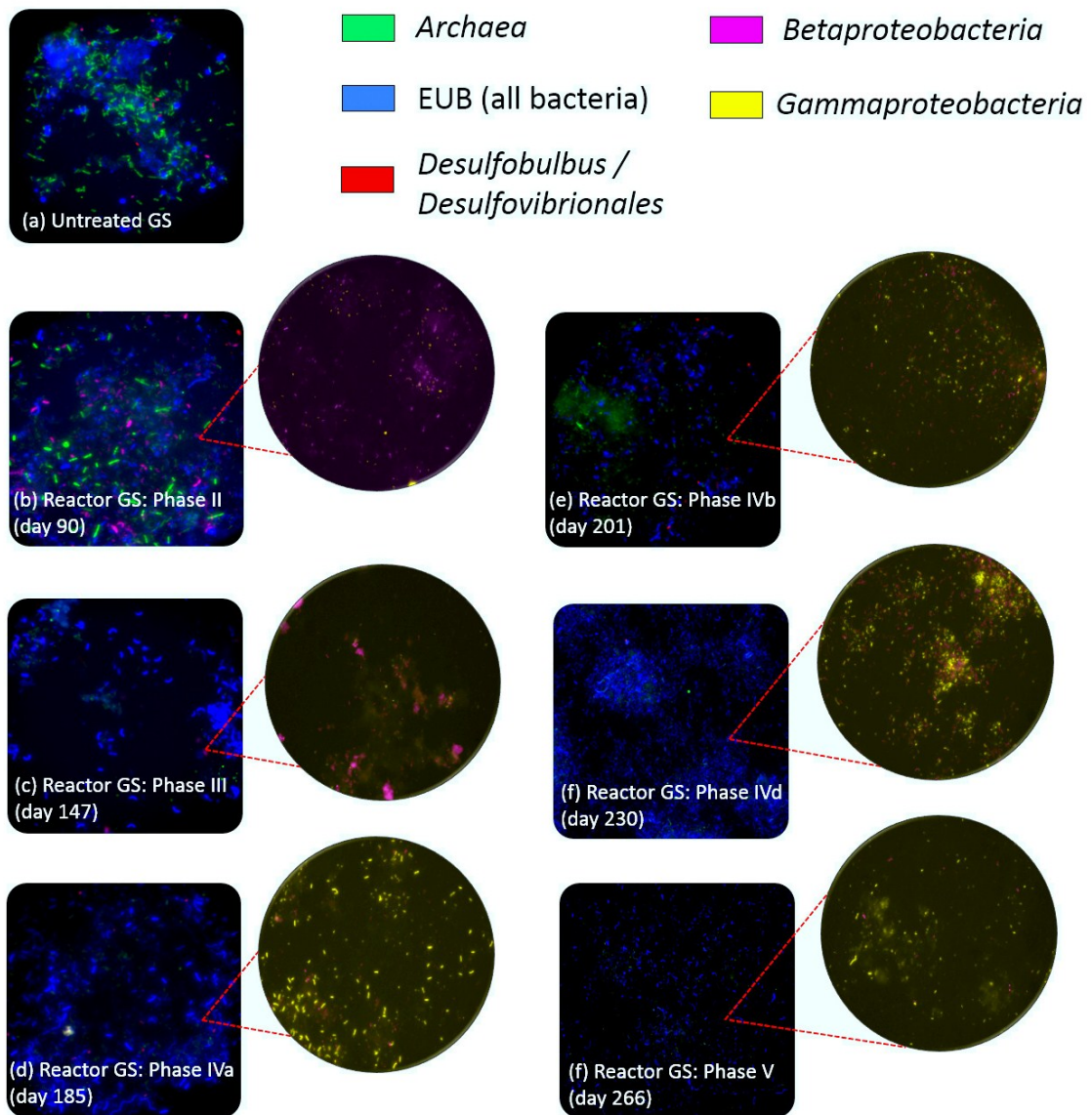
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43 **Fig. S3** Red-color formation in the reactor walls and tubings due to Se<sup>0</sup> production.



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45 **Fig. S4** Microbial community analysis of anaerobic granular sludge sampled from the  
 46 laboratory-scale UASB reactor using *fluorescence in situ hybridization* (FISH) technique.  
 47 FISH images are grouped into EUB-SRB-Archaea (box shape) and BETA-GAMMA (circle  
 48 shape).

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50 **References**

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