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## **Electronic Supplementary Information**

for

Inhibition and recovery of anaerobic granular sludge performances in response to shortterm polystyrene nanoparticles exposure

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This electronic supplementary information contains 4 tables and 4 figures.

#### **1.** Synthesis of polystyrene nanoparticles (PS-NPs)

Functionalized PS-NPs (PS-SO<sub>3</sub>H and PS-NH<sub>2</sub>) were synthesized in our laboratory through the miniemulsion polymerization method. The styrene monomer was washed three times by a base solution (10% NaOH) to remove its inhibitor. The anionic PS-SO<sub>3</sub>H was synthesized with sodium dodecyl sulfate (SDS) as emulsifiers and potassium persulphate (KPS) as initiator.<sup>1, 2</sup> The cationic PS-NH<sub>2</sub> was synthesized through nitrogen-protected polymerization for 5 h at 80 °C with the stirring rate fixed at 400 rpm with dodecylamine hydrochloride and 2, 2'-azobisisobutyronitrile (AIBN). After the synthesis, the latex was transferred to dialysis bag (1 kDa) for removing styrene monomer, emulsifier, initiator and other impurities in the reaction system. The diameter and morphologies of PS-NPs were characterized by a scanning electron microscope (Hitachi S-5000, Hitachi, Japan). Size (Z-average) and  $\zeta$ -potential (mV) were determined using dynamic light scattering (Zetasizer Nano ZS, Malvern, UK). The number and size/concentration of particles were measured by a NanoSight NS500 (NanoSight, Amesbury, UK).

### 2. Modified Gompertz fitting

The modified Gompertz model was used to simulate methane production. Equation 1 shows modified Gompertz equation.<sup>3</sup> And the model simulate results are shown in Table S3.

$$Y(t) = Aexp\left(-\exp\left(\frac{\mu e}{A}(\lambda - t) + 1\right)\right) \qquad Equation \ 1$$

Where:

Y—Cumulative of specific methane production (mL)

A—Methane production potential (mL)

 $\mu$ —Maximum methane production rate (mL/h)

 $\lambda$ —Lag phase period (h)

t—Cumulative time for methane production (h)

e—Mathematical constant

(2.718282)

#### **3. EPS extraction**

The EPS matrix was extracted using cation exchange resin method to elucidate the possible roles of EPS wrapping the AGS in response to PS-NPs.<sup>4</sup> Specifically, the sludge was centrifuged at 6000 r/min for 15 min at 4 °C to remove any EPS from the bulk water. Then, the sludge pellets were resuspended to their original volume in 1% sodium chloride solution containing cation exchange resin (CER) (60 g/g MLSS). The suspension was stirred at 500 r/min for 12 h at 4 °C. The extracted EPS were collected by centrifugation of the CER and sludge mixture at 10,000 r/min for 10 min to remove the CER. Finally, the supernatants were filtered through 0.22 µm acetate cellulose membranes and lyophilized to obtain crude EPS.

#### 4. FTIR analysis

The extracted EPS and PS-NPs were dissolved with 0.2 M phosphate buffer solution (PBS, pH 7.4) in 50 mL centrifuge tubes. Then, the mixed solution was put into an oscillator to mix and balanced without stirring for 6 h at 37 °C before spectral analysis. The infrared spectra of EPS with or without PS-NPs were measured in potassium bromide pellets using an FTIR spectrometer (Aratar, Thermo NicoLet, USA) at wavenumbers from 4000 to 400 cm<sup>-1</sup>. Each spectrum was collected with a resolution of 2 cm<sup>-1</sup>, and the ordinate was expressed as absorbance. To obtain detailed information regarding protein secondary structures, the amide I region (1700–1600 cm<sup>-1</sup>) of the FTIR spectrum was further analyzed. The amide I region was deconvoluted to divided overlapping peaks by increasing the spectral resolution. Then, the spectrum was further broken up into component peaks as protein secondary structures through nine-point Savitzky-Golay derivative function analysis and smoothing.<sup>5</sup> A Lorentzian line shape was quantitated for the amide I region prior to curve fitting the original spectra using Peakfit 4.12 software.

Components	Elements	Concentration (g/L)		
Constant elements	glucose	2.00		
	NH <sub>4</sub> Cl	0.20		
	KH <sub>2</sub> PO <sub>4</sub>	0.05		
Trace elements Stock solution	$CaCl_2 \cdot 2H_2O$	16.70		
(Take 13.5 mL to 1 L when using)	MgCl <sub>2</sub> ·6H <sub>2</sub> O	120.00		
	KCl	86.70		
	$MnCl_2 \cdot 4H_2O$	1.33		
	CoCl <sub>2</sub> ·6H <sub>2</sub> O	2.00		
	H <sub>3</sub> BO <sub>3</sub>	0.38		
	$CuCl_2 \cdot 2H_2O$	0.18		
	$Na_2MoO_4$ ·2H <sub>2</sub> O	0.17		
	ZnCl <sub>2</sub>	0.14		
	FeCl <sub>2</sub> ·4H <sub>2</sub> O	24.67		
Buffer solution	NaHCO <sub>3</sub>	2.00		

Table S1 The compositions of synthetic wastewater

Table S2 Primers used for qPCR amplifications in this study.

	Primer								
Name	Sequence of the primer	Target gene	Expected	Annealing	Reference				
			amplicon	(°C)					
K90	GCGGTGGAGCATGTGGTTTA	16S rRNA	181bp	55	6				
K94	GATAAGGGTTGCGCTCGTTG								
mcrA-F	TGTAAAACGACGGCCAGTGGTGGTGTMGGATTC	mcrA	464bp	51	7				
	ACACARTAYGCWACAGC								
<i>mcrA</i> -R	CAGGAAACAGCTATGACCTTCATTGCRTAGTTW								
	GGRTAGTT								
ACAS-F	TAATCCGCCAAAAGAGTTGG	ACAS	138bp	56	8				
ACAS-R	TCTTCTGGACTGGCTGGTCT								

Sample	First cycle				Second cycle			
	A (mL)	$\lambda$ (h)	$\mu(mL/h)$	R <sup>2</sup>	A (mL)	$\lambda$ (h)	$\mu(mL/h)$	<b>R</b> <sup>2</sup>
blank	51.29	9.11	0.39	0.99	9.75	2.33	0.21	0.98
60 µg/mL PS-SO <sub>3</sub> H	52.10	8.88	0.39	0.98	9.40	6.51	0.28	0.99
80 µg/mL PS-SO <sub>3</sub> H	50.88	1.48	0.44	0.98	10.94	9.57	0.34	0.99
100 μg/mL PS-SO <sub>3</sub> H	42.81	-1.32	0.30	0.97	11.39	8.53	0.27	0.99
$5 \ \mu g/mL \ PS-NH_2$	49.84	4.41	0.38	0.99	11.41	11.20	0.27	0.99
$10 \ \mu g/mL \ PS-NH_2$	45.87	6.55	0.39	0.98	8.45	8.73	0.23	0.99
$20 \ \mu g/mL \ PS-NH_2$	40.17	5.37	0.41	0.99	10.26	6.12	0.25	0.99

# Table S3 Modified Gompertz parameters.

		First cycle		Second cycle			
tax_name	Dlaula	100 µg/mL	20 μg/mL	Dlaula	100 µg/mL	$20 \ \mu g/mL$	
	Blank	PS-SO <sub>3</sub> H	PS-NH <sub>2</sub>	Blank	PS-SO <sub>3</sub> H	PS-NH <sub>2</sub>	
Euryarchaeota	99.61	99.81	99.52	99.60	99.43	99.48	
Pacearchaeota	0.22	0.05	0.39	0.09	0.05	0.09	
Crenarchaeota	0.10	0.06	0.05	0.17	0.45	0.34	
Woesearchaeota	0.07	0.08	0.02	0.14	0.06	0.09	
Other	0.0	0.0	0.02	0.0	0.01	0.0	

Table S4. The relative abundances of Archaea phyla in the first and second cycles.



Fig. S1 The dynamic light scattering of the synthetic (A) PS-SO<sub>3</sub>H and (B) PS-NH<sub>2</sub>.









Fig. S3 Curves of other VFAs over time for the first (black) and second (red) cycle: propionic acid (A: PS-SO<sub>3</sub>H and B: PS-NH<sub>2</sub>); n-butyric acid (C: PS-SO<sub>3</sub>H and D: PS-NH<sub>2</sub>); isobutyric acid (E: PS-SO<sub>3</sub>H and F: PS-NH<sub>2</sub>); isovaleric acid (G: PS-SO<sub>3</sub>H and H: PS-NH<sub>2</sub>).



Fig. S4 The relative abundances of (A) bacterial and (B) archaeal phylum of A1: blank, A2: 100  $\mu$ g/mL PS-SO<sub>3</sub>H and A3: 20  $\mu$ g/mL PS-NH<sub>2</sub> after 1<sup>st</sup> cycle; B1: blank, B1: 100  $\mu$ g/mL PS-SO<sub>3</sub>H and B3: 20  $\mu$ g/mL PS-NH<sub>2</sub> after 2<sup>nd</sup> cycle.

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