

1 **Reduction of soluble microbial products in activated sludge system with pure**
 2 **oxygen aeration under toxic stress condition of phenol**

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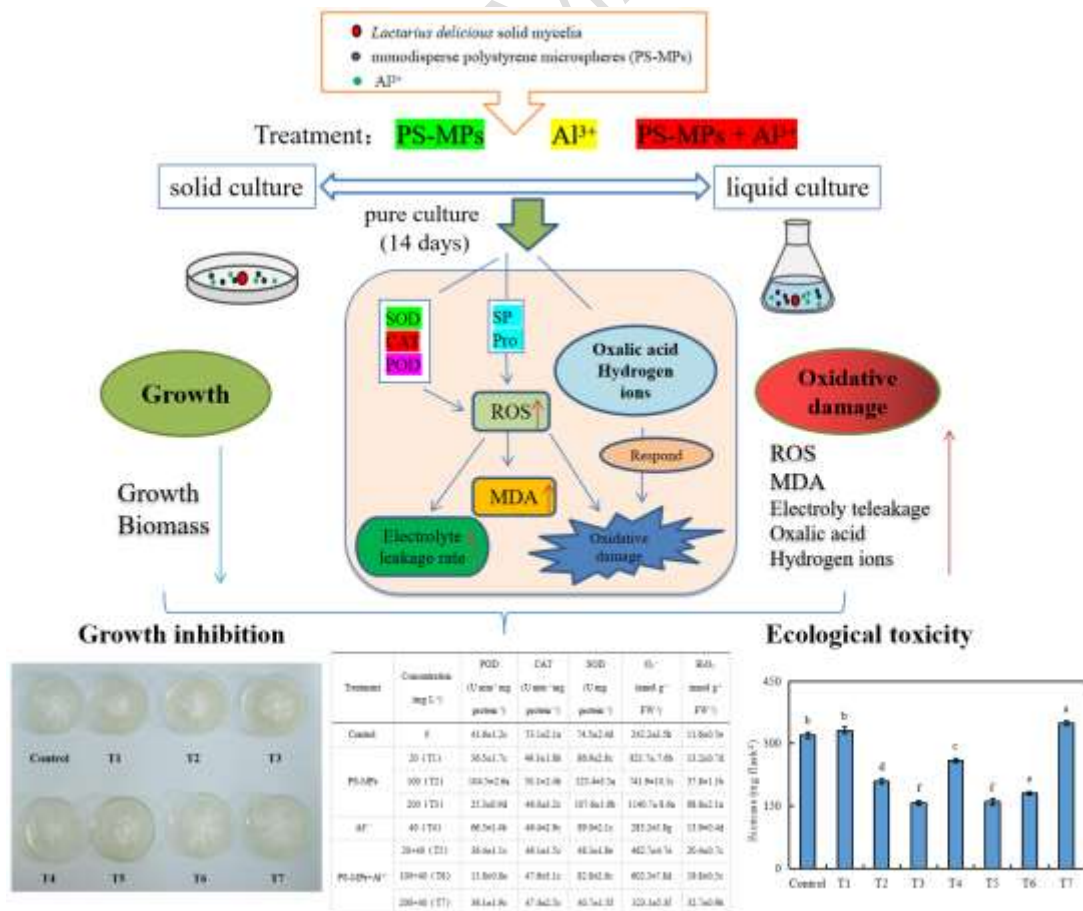
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11 **GRAPHICAL ABSTRACT**



13 **Abstract**

14 This study aimed to explore the effect of pure oxygen aeration on the generation of
15 soluble microbial products (SMP) when activated sludge was under toxic stress
16 condition of phenol. Two parallel sequencing batch reactors, one with air and the other
17 with pure oxygen aeration, were used to investigate the relationships among effluent
18 chemical oxygen demand (COD), N-acyl-homoserine lactones (AHLs), microbial
19 community, extracellular polymeric substances (EPS), and SMP. The results showed
20 that pure oxygen aeration caused rapid recovery of the activated sludge and low effluent
21 COD concentration in steady condition when the two reactors were fed with phenol. In
22 contrast, these results were not observed with sodium acetate as substrate. A strong
23 correlation was found among C4-HSL, loosely bound EPS (LB-EPS), and SMP. The
24 genera *Zoogloea*, *Bacteroides*, and *Flavobacterium* were also related to effluent COD,
25 SMP, and LB-EPS production under toxic stress condition of phenol. Pure oxygen
26 aeration reduced AHL production and decreased the relative abundance of specific
27 genera (*Zoogloea* and *Flavobacterium*). These changes caused the reduction of SMP
28 generation in the pure oxygen-aerated activated sludge systems. These results provide
29 insights into the effect of pure oxygen aeration on SMP generation of activated sludge
30 with toxic substance existence.

31 **Keywords**

32 extracellular polymeric substances, N-acyl-homoserine lactones, pure oxygen aeration,
33 quorum sensing, soluble microbial products

34 **INTRODUCTION**

35 Activated sludge processes are widely applied not only in municipal wastewater but
36 also in industrial wastewater treatment. In these processes, oxygen plays an essential
37 role in contaminant biodegradation by microbes. However, some characteristics of
38 industrial wastewater, such as high concentration of pollutants, high salinity, and oil
39 content, generally have an adverse effect on oxygen transfer (Zhuang *et al.* 2016; Hu *et*
40 *al.* 2019; Wang *et al.* 2021). Moreover, industrial wastewater also probably contains
41 toxic and refractory pollutants. Some researchers promoted oxygen transfer and
42 improved the removal of toxic and refractory compounds through pure oxygen-aerated
43 activated sludge processes (Zhuang *et al.* 2016).

44 Pure oxygen aeration is not always indispensable. The pure oxygen-aerated
45 activated sludge processes exhibit several advantages: improved oxygen transfer, fast
46 organic removal rate, high enzyme activity, and less electricity consumption during
47 aeration (Skouteris *et al.* 2020). Although oxygen production was considered as an
48 additional energy-consuming process, the oxygen source can be replaced by oxygen
49 waste gas obtained from industrial manufacturing processes (Wang *et al.* 2021). Thus,
50 pure oxygen-aerated activated sludge systems can be advisable to treat industrial
51 wastewater.

52 Soluble microbial products (SMP) are mainly derived from soluble organic
53 substances arising from substrate degradation, metabolism, and microbial decay (Soh
54 *et al.* 2020). SMP is generally the main component of secondary effluent organic matter
55 (Yu *et al.* 2015). The contribution of SMP is also non-negligible in the effluent of
56 biological treatment processes for industrial wastewater (Yu *et al.* 2015). Moreover,

57 SMP is still considered as the potential precursor of disinfection by-products and the
58 foremost reason for membrane fouling (Yu *et al.* 2015).

59 Pure oxygen aeration affected the production of SMP and extracellular polymeric
60 substances (EPS) (Skouteris *et al.* 2020). The content of EPS was less under pure
61 oxygen aeration because of its decomposition (Zhang *et al.* 2019). The production of
62 SMP under pure oxygen aeration was affected by the food-to-microorganism (F/M)
63 ratio (Zhang *et al.* 2019). There was no significant difference in SMP contents between
64 air and pure oxygen aeration under low F/M ratio conditions. In contrast, SMP with
65 pure oxygen aeration increased in the conditions with high F/M ratios. These batch
66 experiments were operated with readily biodegradable substances as substrate.

67 The effect of aeration type on EPS and SMP production was complex when the
68 activated sludge was under stress conditions. However, limited research has focused on
69 it (Hu *et al.* 2019). Pure oxygen aeration caused more EPS production than air aeration
70 under stress conditions of salinity (Hu *et al.* 2019). The increase in EPS was related to
71 the self-protection of microorganisms against stress conditions. However, the
72 production of SMP under pure oxygen aeration depended on the extent of stress
73 conditions (Hu *et al.* 2019). Pure oxygen aeration promoted higher content of SMP
74 production than air aeration at low stress conditions (salinity), whereas the opposite was
75 observed under high stress conditions. The researchers believed that the high content
76 of SMP was related to EPS release to bulk solution and a high rate of substrate
77 utilization under pure oxygen aeration under stress conditions of salinity. However, the
78 stress effect of industrial wastewater on the activated sludge frequently originated from

79 toxic and refractory compounds. Thus, it is more important to investigate the effect of
80 pure oxygen aeration on SMP production when the activated sludge was under stress
81 conditions of the toxic substances.

82 The generation of EPS and SMP is related to the microbial community, which can
83 be affected by the type of aeration (Skouteris *et al.* 2020). Moreover, EPS excretion
84 also depends on quorum sensing (QS), and stress conditions provoke signal molecule
85 excretion by microbes (Shi *et al.* 2017). Thus, microbial community structure and signal
86 molecule variation may reveal the effect of pure oxygen aeration on EPS and SMP
87 production under stress conditions. The objective of this study was to investigate the
88 effect of oxygen aeration on EPS and SMP production under stress condition of phenol.
89 Two parallel SBRs with different aeration types (air and pure oxygen) were operated
90 under shock loading of phenol. The relationships among the type of aeration, signal
91 molecule, microbial community, EPS, and SMP were investigated during the non-
92 steady and steady states of the two SBR systems.

93 **MATERIALS AND METHODS**

94 **Reactor set up, operation, experimental design, and sampling**

95 Two identical sequencing batch reactors, A-SBR and O-SBR (with a working
96 volume of 4.5 L), were used and operated parallel using air and pure oxygen aeration,
97 respectively (Fig. S1). An oxygen generator supplied pure oxygen in O-SBR, while the
98 air was aerated to the A-SBR system by a pump. The flow rates of aeration in A- and
99 O-SBRs were kept at 1 L/min during aeration. The SBR systems run at room
100 temperature and were operated as follows: 10 min of influent filling, 240 min of

101 aeration, 30 min of settling, 10 min of effluent withdrawal, and 70 min of resting.

102 The characteristics of the synthetic wastewater were as follows: chemical oxygen
103 demand (COD) of 500 mg/L, NH_4Cl of 95 mg/L, and KH_2PO_4 of 22.4 mg/L. Then, Ca^{2+} ,
104 Mg^{2+} , and trace elements were also added to the synthetic wastewater. The sludge was
105 obtained from the secondary sedimentation tank at Xianyang Road Wastewater
106 Treatment Plant in Tianjin, China. The details of the experimental design and system
107 operation parameters are shown in Table S1.

108 The water samples of influent and effluent from both reactors were collected from
109 the two SBRs and then passed through 0.45 μm membrane filters for water quality
110 analysis. The parameters of COD, phenol, acetate, dissolved oxygen (DO), total
111 phosphorus, ammonia nitrogen, and total nitrogen were determined by the standard
112 methods (SEPA, 2002). The acetate concentration was detected by a gas chromatograph
113 with a flame ionization detector. The phenol concentration was measured using 4-
114 Aminoantipyrine and spectrophotometry. The sludge samples (approximately 200 mg
115 in dry weight) were obtained from the two reactors with replication. These replications
116 were mixed into single samples and frozen at $-20\text{ }^\circ\text{C}$ for DNA extraction.

117 **Extraction and analysis of EPS and SMP**

118 A modified heat extraction method was used to separate EPS from activated sludge (Li
119 & Yang, 2007). The total EPS included tightly bound EPS (TB-EPS) and loosely bound
120 EPS (LB-EPS). Then, 50 mL of the sample with sludge and water mixture was
121 centrifuged and dewatered at 4,000 g for 5 min. The residual sludge was resuspended
122 and diluted with a heated NaCl solution (0.05 %, 70 $^\circ\text{C}$) to the original volume (50 mL).

123 The sludge mixture could reach a temperature of approximately 50 °C. The mixture was
124 shaken by a vortex mixer (VM-370, INTLLAB) for 1 min and centrifuged at 4,000 g
125 for 10 min. The supernatant was collected and passed through a 0.45 µm membrane
126 filter. The collection was considered as the LB-EPS of the biomass. The NaCl solution
127 was added to the residual solid again (50 mL). The supernatant containing TB-EPS was
128 extracted from the resuspended mixture at 4,000 g centrifuged (15 min) after a water
129 bath at 60 °C for 30 min.

130 The SMP was extracted from the effluents of the two SBR systems. The effluent
131 samples were passed through a 0.45 µm membrane filter, and the SMP was involved in
132 the effluent samples.

133 The compositions and concentration of the EPS and SMP extractions were
134 determined by humic-like substances (HS), polysaccharides (PS), and proteins (PN).
135 The PN and HS were analyzed through the modified Lowry method, and the PS was
136 determined with the phenol-sulphuric acid method.

137 **Sampling, extraction, and analysis of AHLs**

138 N-acyl-homoserine lactones (AHLs) were extracted from the supernatant, and 400 mL
139 of supernatant was harvested. The method of AHL extraction was introduced by Tang
140 *et al.* (2015). The supernatant was passed through a cellulose filter (0.22 µm) and
141 extracted with an equal volume of ethyl acetate in triple. The solvent was collected and
142 mixed in a conical flask. The collected solvent was dried with anhydrous sodium sulfate
143 and then evaporated at 40 °C via a rotary evaporator (Yarong, China). The extract was
144 resolute by 2 mL methanol and stored at -20 °C for further analysis.

145 The AHLs in the samples were identified and determined via high-performance
146 liquid chromatography-mass spectrometry (Agilent, USA) with an XDB-C18 column
147 (5 μm , 4.6 mm \times 180 mm, Agilent, USA). In the high-performance liquid
148 chromatography condition, the mobile phase consisted of solvent A (ultrapure water
149 with 0.1% of formic acid in v/v and 2 mmol/L ammonium acetate) and solvent B
150 (methanol with 0.1% of formic acid in v/v and 2 mmol/L ammonium acetate) for
151 gradient elution. The column temperature was kept at 40 $^{\circ}\text{C}$, and the flow rate was set
152 at 0.2 mL/min.

153 In the mass spectrometry condition, electric spray ion scanning and multi-reaction
154 ion monitoring mode were used. The voltage was set at 4 kV, and the heated capillary
155 was maintained at 350 $^{\circ}\text{C}$.

156 In this study, N-Butanoyl-L-homoserine lactone (C4-HSL), N-hexanoyl-L-
157 homoserine lactone (C6-HSL), N-octanoyl-L-homoserine lactone (C8-HSL), and N-
158 Decanoyl-L-homoserine lactone (C10-HSL) were continuously found and determined
159 in the two reactors. The details of the standard curves are shown in Supplementary
160 Materials.

161 **High-throughput sequencing analysis**

162 High-throughput sequencing analysis was implemented by Novogene (Beijing, China).
163 The processes of DNA extraction were followed according to our previous study (Wang
164 *et al.* 2021).

165 First, DNA from the sludge samples was extracted via CTAB method. In this
166 process, 1% agarose gel was used to monitor the DNA concentration and purity. Then,

167 the DNA of the samples was diluted to 1 ng/ μ L. The V4 region of the 16S rDNA gene
168 with the forward primer 515F and reverse primer 806R was amplified. Quantification
169 of amplicons was detected through electrophoresis with 1X loading buffer and 2%
170 agarose gel. Then, the amplicons were mixed in equidensity ratios and purified with a
171 Qiagen Gel Extraction Kit (Qiagen, Germany). The sequencing was processed through
172 the Illumina NovaSeq platform. Data split, sequence assembly, and filtration were also
173 completed via FLASH (V1.2.7, <http://ccb.jhu.edu/software/FLASH/>) and QIIME
174 (V1.9.1, http://qiime.org/scripts/split_libraries_fastq.html). Canonical correspondence
175 analysis (CCA) was conducted from the online platform (Novogene) and regenerated
176 through Origin 2021.

177 **Statistical analyses**

178 SPSS 22 was used in statistical analyses. The differences in effluent COD, SMP and
179 EPS contents between the two SBR systems were obtained via paired t-test analysis.
180 The relationships among AHL contents and EPS and SMP components were obtained
181 via Pearson correlation analysis. The negative coefficients (R) of the Pearson
182 correlation (-1-0) indicated a negative correlation. The positive R values (0-1) show a
183 positive correlation. R = 0 represents noncorrelation. The coefficient matrixes were
184 shown by heatmap via Origin 2021.

185 **RESULTS AND DISCUSSION**

186 **Organic removal performance**

187 The COD removal performances of A- and O-SBR are depicted in Fig. 1. When the
188 substrate was altered from sodium acetate to phenol, the effluent COD of both reactors

189 increased and gradually decreased. Phenol cannot be degraded by non-incubation
190 activated sludge in the initial period. The toxic effect of phenol also caused biomass
191 decay and cell lysis, which built up SMP (Wu *et al.* 2016). The residual phenol and
192 SMP caused the increase of effluent COD.

193 The changes of effluent COD from A- and O-SBRs showed a significant difference
194 with phenol as substrate. Pure oxygen aeration accelerated the recovery of COD
195 removal performance compared with air aeration in the non-steady state of the phenol
196 shock loading stage. This condition indicates that pure oxygen aeration alleviated the
197 toxic effect of phenol on the activated sludge due to the high enzyme activity during
198 pure oxygen aeration (Calderon *et al.* 2012).

199 In the steady state (Fig. 1b), the effluent COD concentration of A- and O-SBRs
200 showed no significant difference when the two reactors were fed with sodium acetate.
201 In contrast, the effluent COD of O-SBR was lower than that of A-SBR with phenol as
202 substrate via t-test analysis ($p < 0.001$). This finding implies that pure oxygen aeration
203 can polish the effluent quality with toxic carbon source feeding rather than a readily
204 biodegradable carbon source. Both sodium acetate and phenol were in low
205 concentration because of their removal by incubation activated sludge (Table S2)
206 (Zhuang *et al.* 2016). The residual COD mainly consisted of SMP in the two reactors.
207 The relatively low COD concentration in O-SBR may be related to reducing SMP by
208 pure oxygen aeration.

209 **Content variations of EPS and SMP**

210 The changes of EPS and SMP contents were investigated to elucidate the effect of pure

211 oxygen aeration on the SMP production of activated sludge with toxic substrate feeding
212 (Fig. 2). As shown in Fig. 2a, the type of aeration seemed not to affect the SMP
213 production when the systems were fed with sodium acetate. This result coincides with
214 the findings of Zhang *et al.* (2019b). The reason is that the substrate plays a more critical
215 role in the SMP generation of activated sludge compared with other factors such as
216 oxygen supply (Xu *et al.* 2011). Sodium acetate, a readily biodegradable substrate, has
217 no adverse influence on the SMP generation of the activated sludge. Hence, the SMP
218 contents had no difference between air and pure oxygen aeration in this study.

219 When the substrate was altered to phenol, the SMP contents initially increased
220 significantly and gradually decreased in the two reactors. The activated sludge in O-
221 SBR released less SMP than that in A-SBR (Fig. 2a). The increase of SMP was related
222 to cell lysis and EPS release into the bulk solution with toxic substrate existence
223 (Aquino & Stuckey, 2004). This result also represents that pure oxygen aeration
224 reduced the SMP production of activated sludge under stress conditions, which was
225 different from previous findings (Hu *et al.* 2019). Hu *et al.* (2019) found that pure
226 oxygen aeration increased the SMP generation of the activated sludge under stress
227 conditions of low salinity due to the high organic removal rate. The reason is that the
228 high content of SMP release was related to biomass-associated product generation of
229 improved microbial metabolism and utilization-associated products from rapid organic
230 degradation (Hu *et al.* 2019). In this study, a high organic removal rate was also
231 obtained (Table 1). The reduced SMP production by oxygen-aerated sludge under toxic
232 stress condition of phenol may be related to rapid substrate removal rate and toxicity

233 alleviation.

234 One of the SMP production sources originates from EPS hydrolysis (Ni *et al.* 2010)
235 The changes of EPS content are shown in Figs. 2b, 2c, and 2d. When the substrate was
236 altered to phenol, the total and LB-EPS contents increased in both the reactors and
237 gradually decreased. However, the TB-EPS content increased insignificantly in the two
238 reactors with phenol feeding. The contents of the total, LB-, and TB-EPS in the
239 presence of phenol were higher than those in the absence of phenol. The increase of
240 EPS excretion is beneficial in protecting microbes from toxic substrates (Zhang *et al.*
241 2021b). In this study, the changing pattern of LB-EPS was similar to that of SMP in the
242 two reactors. It indicates that LB-EPS had the potential to be released into the effluent
243 as SMP due to their loose structure.

244 The type of aeration caused a difference in EPS content between O- and A-SBR.
245 The TB-EPS content in O-SBR were higher than those in A-SBR (Fig. 2c). These
246 results showed that pure oxygen aeration caused more TB-EPS excretion of activated
247 sludge than air aeration. Some researchers found that pure oxygen aeration made less
248 EPS generation than air aeration due to EPS consumption as substrate at the end of the
249 aeration stage (Zhuang *et al.* 2016; Zhang *et al.* 2019b). However, the opposite
250 conclusion was found under stress conditions (Hu *et al.* 2019). In this study, high
251 content of TB-EPS was excreted to protect microbes and contain extracellular enzymes
252 with pure oxygen aeration.

253 The variations of EPS components are shown in Fig. 3. The PS and PN contents of
254 LB-EPS in O-SBR were lower than those in A-SBR except the samples in Day 20 (Figs.

255 3a and 3b), while the PS and PN contents of TB-EPS in O-SBR were higher than those
256 in A-SBR (Figs. 3c and 3d). The low PS and PN contents of LB-EPS in O-SBR may
257 be related to toxicity alleviation of toxic substrate removal (Shi *et al.* 2021). The high
258 PS and PN contents of TB-EPS in O-SBR may contain more abundant extracellular
259 redox-active components to promote pollutant reduction than those in A-SBR, such as
260 reducing saccharides and various extracellular enzymes (Yu 2020; Zhou *et al.* 2020).
261 Substrate bioavailability and its removal rate may be promoted by the high content of
262 TB-EPS in O-SBR. The higher efficiency of toxic substrate removal may cause less
263 SMP and LB-EPS generation in O-SBR. Meanwhile, pure oxygen aeration influenced
264 the signal molecule of QS and microbial community structure, which also played an
265 essential role in EPS generation.

266 **Variations of AHLs**

267 The QS plays a vital role in regulating EPS secretion, which further influences SMP
268 production. The variations of AHLs (C4-, C6-, C8-, and C10-HSL) were investigated
269 and shown in Fig. 4. In A-SBR, when the substrate was altered from sodium acetate to
270 phenol, the concentration of C4-HSL in A-SBR increased initially with phenol as
271 substrate and then decreased gradually. The content of C4-HSL in A-SBR was
272 correlated with SMP and LB-EPS (Fig. 5a). The concentrations of C6-, C8-, and C10-
273 HSL in A-SBR increased until day 20 and then decreased gradually.

274 In the activated sludge systems with air aeration, various studies found that AHLs
275 played an essential role in EPS and SMP generation (Zhang *et al.* 2019e; Zhang *et al.*
276 2019d; Xiong *et al.* 2020). However, the effect of AHLs on SMP and EPS generation

277 cannot draw a uniform conclusion. Xiong *et al.* (2020) believed that C6-HSL and
278 3OC6-HSL controlled EPS and SMP production. Some researchers also found that the
279 C8- and 3OHC8-HSL might be necessary for EPS secretion (Zhang *et al.* 2019d; Zhang
280 *et al.* 2019e). In this study, C4-HSL played a critical role in SMP and LB-EPS
281 production. This result is different from previous findings because the experiment was
282 under the stress condition of toxic compound in the present study. In contrast, the
283 former experiments were based on no stress conditions. Ding *et al.* (2016) found that
284 C4-HSL was related to LB-EPS production under oligotrophic nitrogen supply
285 conditions, which is similar to the result of our study. Thus, C4-HSL may be related to
286 LB-EPS and SMP production in stress conditions with air aeration. Meanwhile, Zhang
287 *et al.* (2019c) found that the concentrations of C6- and C8-HSL increased to maintain
288 bacterial activity after substrate shock loading. In our study, C6-, C8-, and C10-HSL
289 increased after phenol shock loading, which could be related to the self-defense
290 behavior of microbes and maintained bacterial activity under stress conditions (Zhang
291 *et al.* 2019c).

292 However, the four AHLs contents in O-SBR were lower than those in A-SBR
293 under shock loading of phenol (Fig. 4). This result indicates that pure oxygen aeration
294 or DO significantly affected AHL generation. Recent research found that facultative
295 anaerobe can generate AHLs only in low DO conditions ($DO < 2$ mg/L) (Huang *et al.*
296 2021). In this study, the production of AHLs by the facultative bacteria may be inhibited
297 during pure oxygen aeration. Thus, the concentrations of AHLs in O-SBR were lower
298 than those in A-SBR. The lower C4-HSL concentrations with pure oxygen aeration

299 caused less SMP and LB-EPS generation. The content of C4-HSL also showed no
300 correlation with LB-EPS and SMP production (Fig. 5b). The lower concentrations of
301 C6-, C8-, and C10-HSL in O-SBR showed non-significant effects on organic removal
302 performance.

303 **Changes in microbial community**

304 The microbial diversity and richness indices are shown in Table 2. The estimators of
305 Chao 1 and ACE indicate the richness of the microbial community. The results of Chao
306 1 and ACE illustrate that the O-SBR had a lower richness of microbial community than
307 A-SBR with sodium acetate as substrate. In contrast, the richness of microbial
308 community in O-SBR was higher than that in A-SBR when the substrate was altered
309 with phenol. The estimators of Shannon and Simpson indicate the diversity of the
310 microbial community. The values of Shannon and Simpson showed that O-SBR had a
311 higher diversity of microbial community than A-SBR with sodium acetate and phenol
312 as substrate. These results demonstrate that the diversity and richness of the microbial
313 community with pure oxygen aeration were higher than those with air aeration with
314 phenol as substrate. The differences in diversity and richness between A- and O-SBR
315 were related to the type of aeration (Zhuang *et al.* 2016). The high diversity and richness
316 of the microbial community with pure oxygen aeration was beneficial to the rapid
317 recovery under toxic stress condition of phenol.

318 The notable changes in microbial communities at the phylum level are considered
319 in Fig. 6a. The predominant phyla in all samples contained *Proteobacteria*,
320 *Bacteroidetes*, and *Firmicutes*. *Proteobacteria* was always dominant in both reactors,

321 with sodium acetate and phenol as substrates. However, the relative abundance of
322 *Proteobacteria* decreased gradually when the substrate was altered from sodium acetate
323 to phenol in both two reactors. On the contrary, the phyla of *Bacteroidetes* and
324 *Firmicutes* increased in both reactors with phenol as substrate. *Proteobacteria*,
325 *Bacteroidetes*, and *Firmicutes* were usually the dominant phyla in activated sludge
326 samples from municipal and industrial wastewater treatment plants (Sebastian *et al.*
327 2020). The relative abundance of *Bacteroidetes* and *Firmicutes* indicates that these
328 phyla contained the functional species related to phenol degradation.

329 The variations of the top 20 genera in the samples are shown in Fig. 6b. The
330 dominant genera in the two reactors contained *Zoogloea*, *Bacteroides*, *Flavobacterium*,
331 *Competibacter*, *Rhizobium*, and *Luteococcus*. The changes in the microbial community
332 in both the SBRs were different when the substrate was altered from sodium acetate to
333 phenol. In the A-SBR system, the genera of *Zoogloea*, *Bacteroides*, and
334 *Flavobacterium* increased significantly with phenol as substrate. These genera were
335 related to EPS generation against toxic conditions (Zhang *et al.* 2019a; Zhang *et al.*
336 2021a). In O-SBR, the genera *Bacteroides*, *Competibacter*, *Rhizobium*, and
337 *Luteococcus* were found to increase and dominate with phenol as substrate. *Rhizobium*
338 and *Luteococcus* were related to phenol degradation (Tong *et al.* 2018; He *et al.* 2021;
339 Zhang *et al.* 2021a;). The genera *Competibacter* and *Luteococcus* were also considered
340 as EPS producers (Wei *et al.* 2020; He *et al.* 2021). However, *Zoogloea* and
341 *Flavobacterium* changed insignificantly in O-SBR. The specific dominant genera in O-
342 SBR, such as *Rhizobium*, preferred high DO concentration (Yadav *et al.* 2020). Thus,

343 the dominant and functional microbes difference between A- and O-SBR was due to
344 the type of aeration and DO concentration difference.

345 The relationships among system performances, EPS, SMP, AHLs, and microbial
346 community are further illustrated via CCA. In Fig. 6c, the acute angles among effluent
347 COD, SMP, LB-EPS, and AHLs indicated a positive correlation. *Zoogloea*,
348 *Bacteroides*, and *Flavobacterium* were also related to effluent COD, AHLs, SMP, and
349 LB-EPS content. It should also be noted that TB-EPS production was correlated with
350 *Luteococcus*. In previous studies, *Zoogloea* and *Flavobacterium* correlated with EPS
351 generation under toxic conditions (Zhang *et al.* 2019a; Zhang *et al.* 2021a).
352 *Flavobacterium* was also proved to be the AHL producer (Zhang *et al.* 2018). In this
353 study, AHL content and microbial community composition affected LB-EPS and SMP
354 production. Compared with A-SBR, the lower relative abundance of *Zoogloea* and
355 *Flavobacterium* in O-SBR caused less LB-EPS and SMP production (Fig. 6b).
356 Furthermore, the high content of TB-EPS in O-SBR was related to the high relative
357 abundance of *Luteococcus*.

358 According to the results above, pure oxygen aeration influenced AHL excretion
359 and microbial community structure. The reduced AHL contents and low abundance of
360 functional bacteria for LB-EPS generation in the reactor with pure oxygen aeration
361 tended to excrete less LB-EPS and SMP. This result caused lower effluent COD with
362 oxygen aeration than with air aeration. These results imply that high dissolved oxygen
363 concentration could reduce the SMP production of activated sludge under stress
364 conditions of the toxic substrates from industrial wastewater.

365 **CONCLUSIONS**

366 In this study, the effect of pure oxygen aeration on activated sludge with phenol as a
367 toxic substrate was investigated via two parallel SBRs using pure oxygen and air
368 aeration. Compared with air aeration, pure oxygen aeration promoted the recovery of
369 activated sludge under shock loading of phenol. Pure oxygen aeration also caused the
370 reduction of AHL excretion and decreased the abundance of functional bacteria for LB-
371 EPS production under stress condition of toxic substrate. This reduced LB-EPS and
372 SMP production of activated sludge in the system with pure oxygen aeration and finally
373 decreased the effluent COD. These results may provide insights into the effect of pure
374 oxygen aeration on SMP generation under stress conditions of toxic compounds.

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378 **Author Contributions**

379 All the co-authors read and approved the final manuscript. Z.H.S. carried out the
380 experiments and data. D. W. conceptualized and supervised the project, acquired
381 funding, provided resources for the experiments, wrote the original draft and
382 contributed to the final revision of the manuscript. C.S.Q. and S.P.W. co-supervised the
383 student and reviewed the written manuscript.

384 **DATA AVAILABILITY STATEMENT**

385 All relevant data are included in the paper or its Supplementary Information.

386 **CONFLICT OF INTEREST**

387 The authors declare that they have no conflict of interest.

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491

492

Table and Figure captions

493

494 **Table 1** COD changes in batch assays of different aeration types with phenol as
495 substrate

496 **Table 2** Bacterial richness and diversity estimators of both reactors with different
497 substrates

498 **Fig. 1** Organic removal performance of two SBR systems. (a) COD removal
499 performance and (b) Effluent COD comparison of the two SBR systems in steady state
500 via t-test analysis

501 **Fig. 2** Content variations of (a) SMPs, (b) total EPS, (c) TB-, and (d) LB-EPS. *: $p <$
502 0.05 ; **: $p < 0.01$; ***: $p < 0.001$

503 **Fig. 3** Content variations of PN in (a) LB-EPS, (b) PS in LB-EPS, (c) PN in TB-EPS,
504 and (d) PS in TB-EPS. *: $p < 0.05$; **: $p < 0.01$; ***: $p < 0.001$

505 **Fig. 4** Changes of (a) C4-HSL, (b) C6-HSL, (c) C8-HSL, and (d) C10-HSL
506 concentration in the reactors

507 **Fig. 5** Pearson correlation among AHLs, SMP, and EPS components in (a) A-SBR and
508 (b) O-SBR (*: $p < 0.05$, **: $p < 0.01$)

509 **Fig. 6** Relative abundance changes of microbial communities at (a) phylum and (b)
510 genus level in the samples, and (c) CCA on correlation of genus with various factors

511

512 **Table 1.** COD changes in batch assays of different aeration types with phenol as
513 substrate.

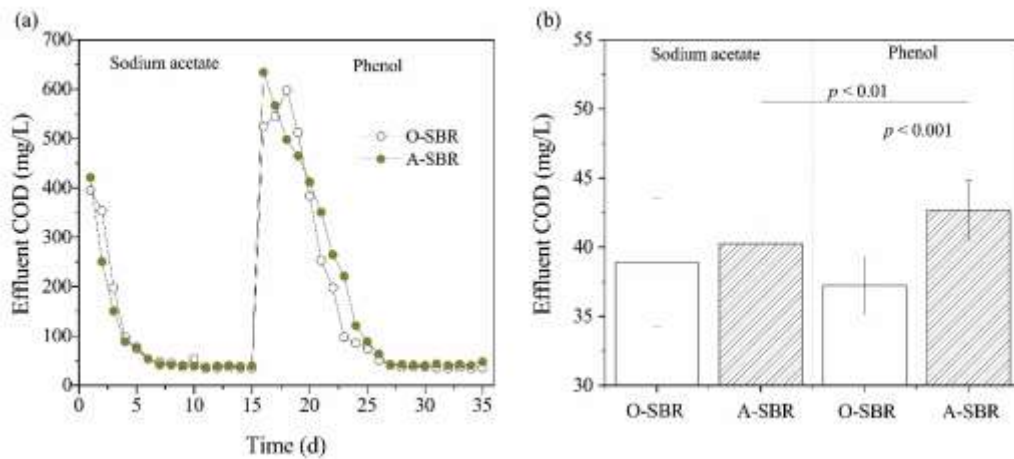
Reactor	Effluent COD (mg/L)					
	0 h	0.5 h	1 h	2 h	3 h	4 h
A-SBR	240±12	153±8	67±5	46±7	43±3	45±7
O-SBR	235±14	48±5	40±3	41±2	35±5	36±9

514

515 **Table 2.** Bacterial richness and diversity estimators of both reactors with different
 516 substrates.

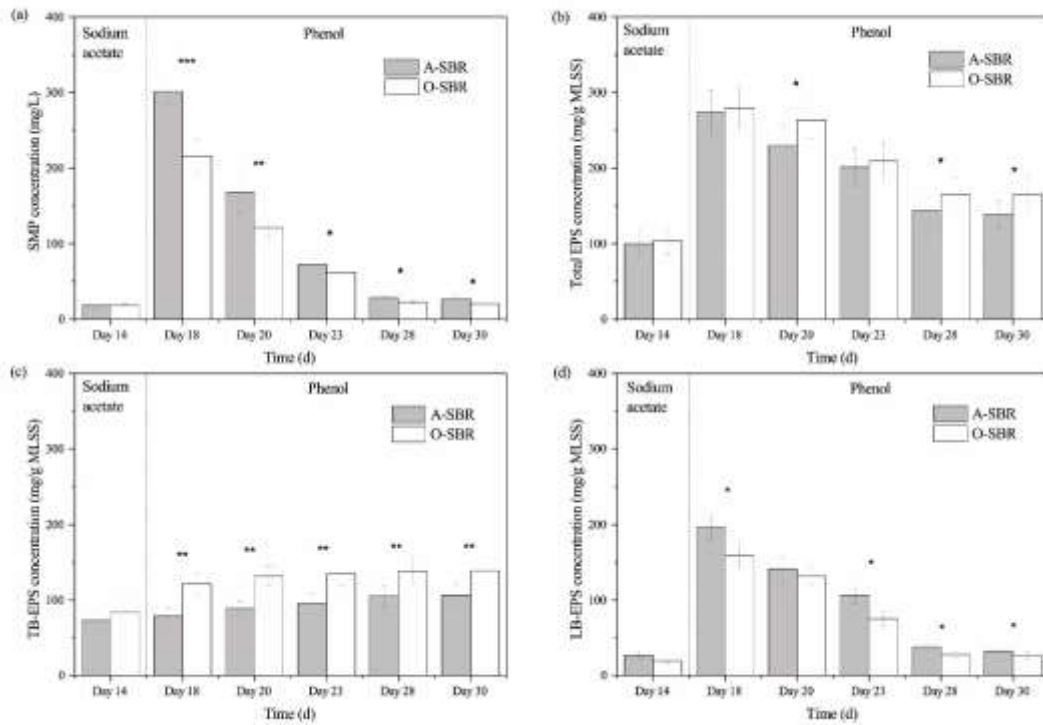
Reactor	Substrate	OTUs	Shannon	Simpson	Chao1	ACE	Goods coverage (%)
A-SBR	Sodium acetate	1407	7.141	0.955	1497.004	1545.389	99.5
	Phenol	848	6.674	0.968	914.275	929.731	99.7
O-SBR	Sodium acetate	1229	7.686	0.987	1332.147	1359.238	99.5
	Phenol	1209	7.491	0.9845	1330.921	1341.0245	99.4

517



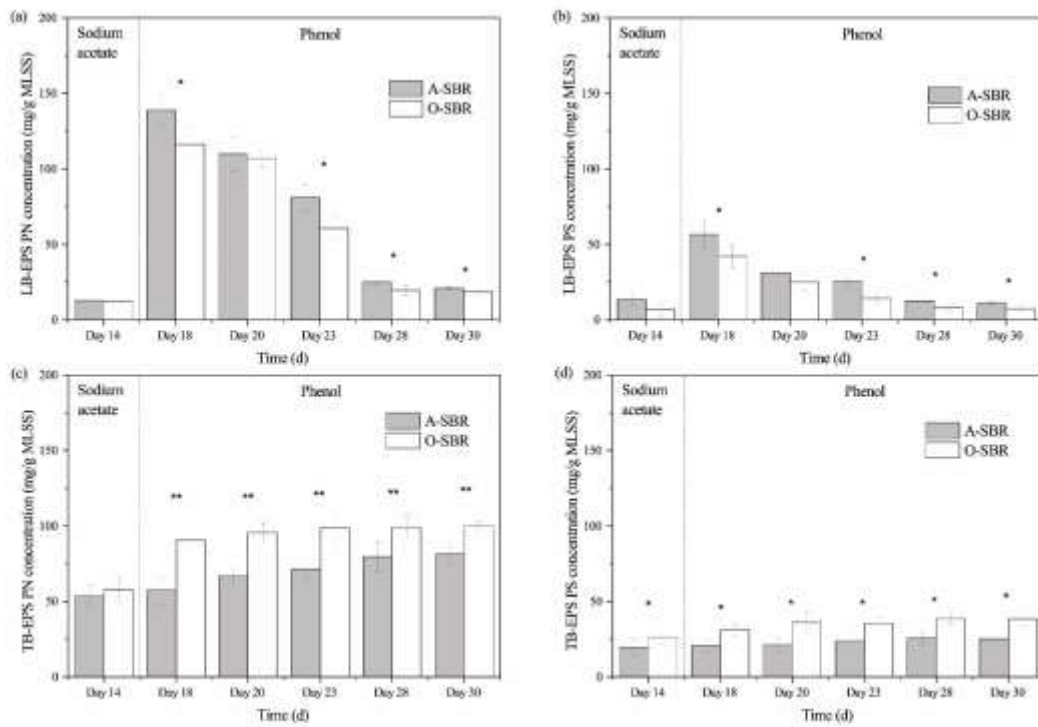
518

519 Figure 1.



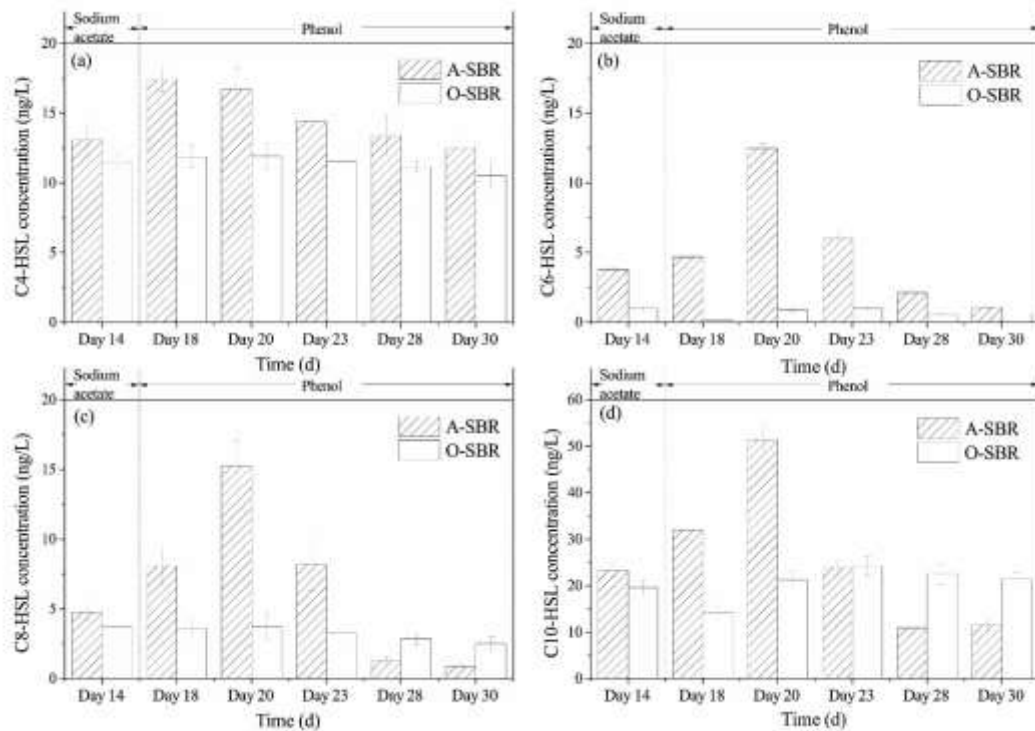
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521 Figure 2.



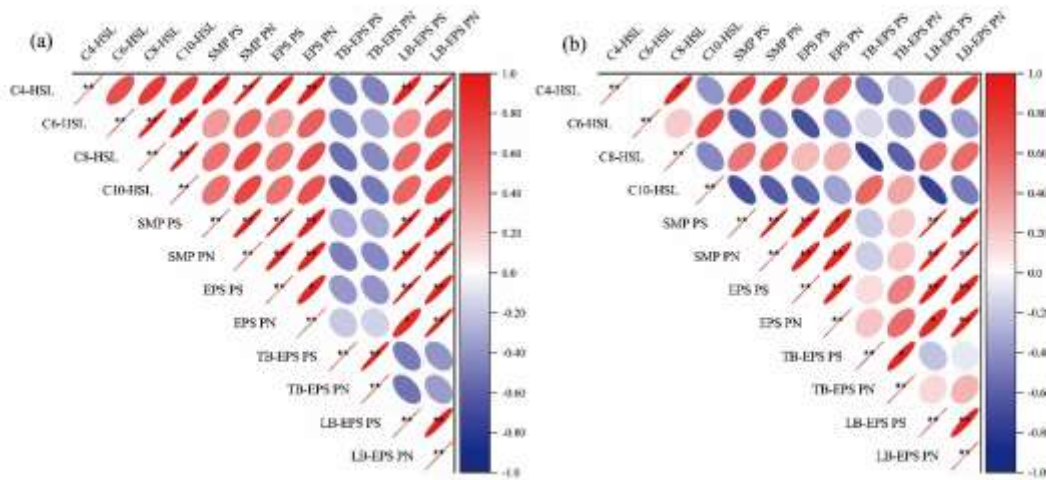
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523 Figure 3.



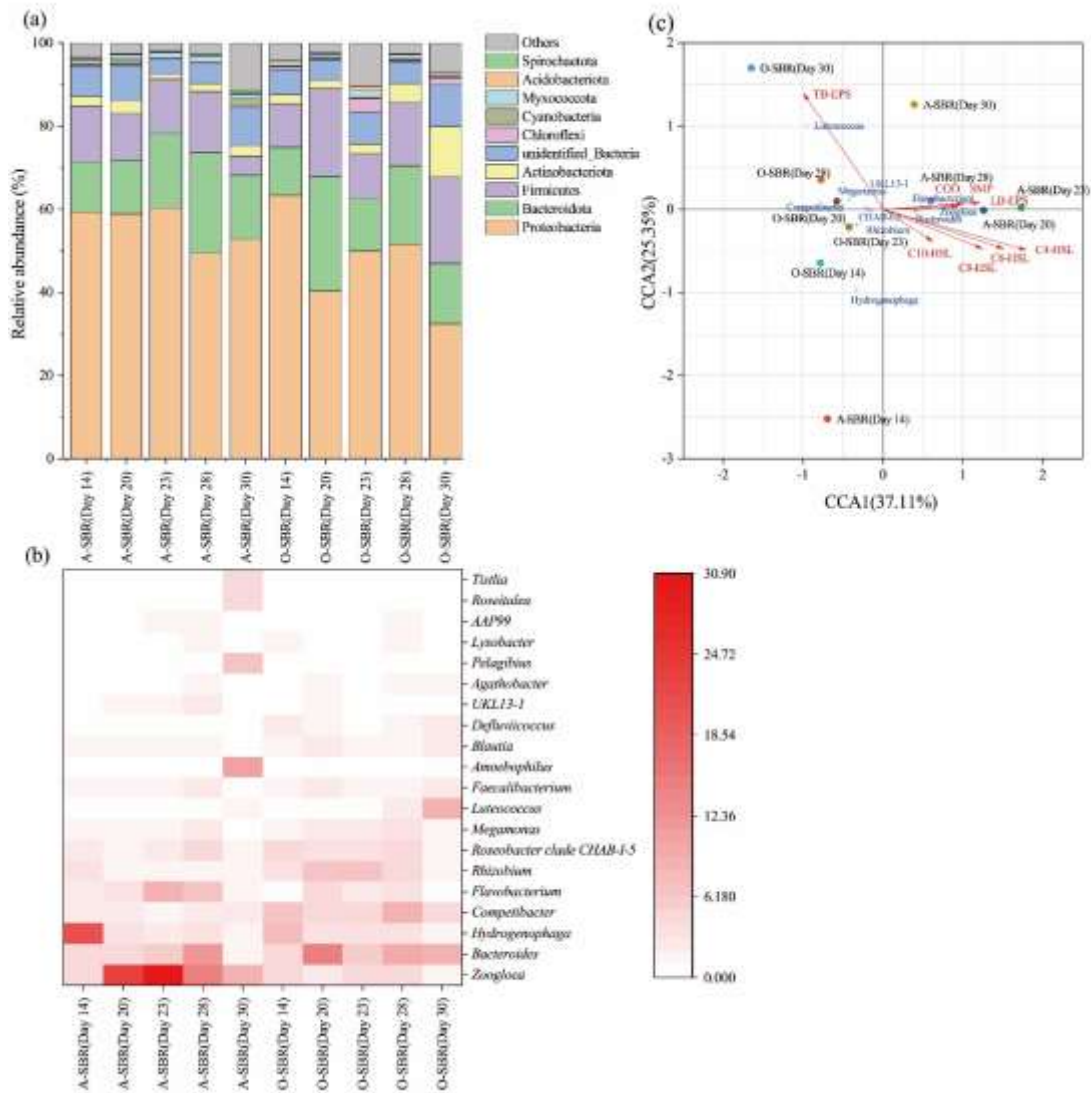
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525 Figure 4.



526

527 Figure 5.



528

529 Figure 6.

ACCEPTED MANUSCRIPT

531 **Supplementary Materials**

532

533 **Reduction of soluble microbial products in activated sludge system**
534 **with pure oxygen aeration under toxic stress condition of phenol**

535

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543

544 **EXPERIMENT:**

545 **Determination of AHLs**

546 In this study, the AHLs standards (C4-HSL, C6-HSL, C8-HSL and C10-HSL) were purchased from
547 Sigma Aldrich. AHLs standards were configured into standard solutions of AHLs with concentration
548 gradients of 2, 4, 6, 8, 10, 20 and 40 µg/L. The standard curves were generated with the peak area
549 corresponding to quantitative ion of m/z 102 via HPLC-MS.

550

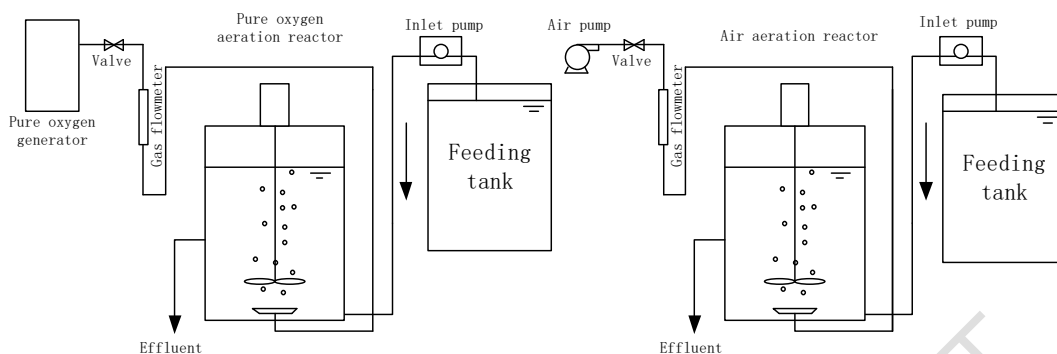
AHL	Standard curve	Recovery (%)
C4-HSL	Y = 163.29X+576.93 R ² = 0.9952	86.43
C6-HSL	Y = 221.76X+224.82 R ² = 0.9935	89.85
C8-HSL	Y =279.07X-37.78 R ² = 0.9943	95.77
C10-HSL	Y =250.29X+7.16 R ² = 0.9911	83.64

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552 **FIGURE:**



553

554

Figure S1. Schematic diagram of the SBR systems with pure oxygen and air aeration.

555

556 **TABLE:**

557 **Table S1.** Details of system operation parameters and sampling.

Reactor	Carbon source	Operation time	State	Sampling time	COD _{inf} (mg/L)	HRT (h)	MLSS (g/L)	SRT (d)
A-SBR	Sodium acetate	Day 1–15	Sludge cultivation	—	500	4	4.1 ± 0.5	10-20
		Day 7–15	Steady state	Day 14	500	4	4.0 ± 0.2	
	Phenol	Day 16–26	Shock loading	Day 18, 20 and 23	500	4	4.0 ± 0.6	
		Day 27–33	Steady state	Day 28 and 30	500	4	3.9 ± 0.1	
O-SBR	Sodium acetate	Day 1–6	Sludge cultivation	—	500	4	4.0 ± 0.3	10-20
		Day 7–15	Steady state	Day 14	500	4	4.1 ± 0.1	
	Phenol	Day 16–26	Shock loading	Day 18, 20 and 23	500	4	3.9 ± 0.4	
		Day 27–33	Steady state	Day 28 and 30	500	4	4.0 ± 0.1	

558

559 **Table S2.** Substrate removal performance of the two SBRs in steady state

Reactor	Carbon source	Sampling time	Concentration (mg/L)
A-SBR	Sodium acetate	Day 10–15	5.2 ± 2.3
	Phenol	Day 30–33	3.2 ± 0.3
O-SBR	Sodium acetate	Day 10–15	4.7 ± 1.5
	Phenol	Day 30–33	3.7 ± 0.6

560

561