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4 1 **Insights into quorum quenching mechanisms to control membrane biofouling under**
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6 2 **changing organic loading rates**
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8
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4 **19 Abstract**

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7 20 A quorum quenching (QQ) consortium comprised of both acyl homoserine lactones
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9 21 (AHLs)- and autoinducer-2 (AI-2)-degrading bacteria, either immobilized in polymer-coated
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11 22 alginate beads or in liquid suspension, was examined for fouling control in lab-scale MBRs
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14 23 under both steady and changing organic loading rates (OLRs). Under steady conditions the QQ
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16 24 consortium retarded biofouling by a factor of 3. However, a continuous increase in OLR vastly
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18 25 reduced the effectiveness of QQ bacteria; the biofouling was retarded only by factors of 1.4-1.8.
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20 26 A significant increase in extracellular polymeric substance (EPS), especially loosely-bound EPS
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22 27 in mixed liquor together with an increase in polysaccharide content up to 4 times in EPS resulted
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24 28 from the increase in OLR, was attributed to the impaired QQ efficacy. In control MBRs, cake
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26 29 layer resistance was the major factor (>60%) contributing to the increased trans-membrane
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28 30 pressure, as compared with pore blockage resistance and intrinsic membrane resistance. In
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30 31 contrast, the pore blockage resistance became dominant in QQ MBRs (>40%).
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36 **32 Key words**

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38 33 Membrane bioreactor (MBR), Biofouling, Quorum sensing, Quorum quenching, organic
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40 34 loading rate (OLR)
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43 **35 1. Introduction**

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45 36 Membrane bioreactor (MBR) technology is becoming a widely recognized and accepted
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47 37 wastewater treatment technology owing to its distinctive advantages, including high treatment
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49 38 efficiency, low sludge production and a small footprint compared to conventional activated
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51 39 sludge (CAS) (Hong et al., 2002; Lin et al., 2014). However, membrane fouling remains a major
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53 40 impediment that limits the widespread application of MBR due to high energy demand and
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55 41 operational costs associated with fouling control (Drews, 2010; Yu et al., 2010). Complex
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4 42 interactions between membrane and mixed liquor components like suspended solids, colloids,
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6 43 biopolymers, and solutes introduced from raw wastewater or produced during biomass growth or
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9 44 decay play a significant role in membrane fouling (Hu et al., 2016). It is known that cake layer
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12 45 formation on the membrane surface accounts for over 80% of the total filtration resistance in
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14 46 most MBR studies (Miura et al., 2007; Wang et al., 2007).

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16 47 In past few years great interest has been focused on mitigating membrane biofouling via
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19 48 quorum quenching (QQ) where bacterial communication is disrupted to reduce cake layer
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22 49 formation on the membrane (Oh et al., 2012). To achieve this, bacteria which produce QQ
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24 50 enzymes were isolated from sludge and applied in MBRs, either in the form of a microbial
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26 51 culture (Cheong et al., 2014; Jahangir et al., 2012), or immobilized in alginate beads (Kim et al.,
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29 52 2013). Successful application of QQ bacteria in terms of biofouling control in MBR systems in
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31 53 lab (Kim et al., 2013, 2015; Maqbool et al., 2015; Oh et al., 2012) as well as pilot scale (Lee et
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33 54 al., 2016) has been reported. These studies have proved the effectiveness of QQ mechanisms for
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36 55 biofouling control under steady conditions; however, the efficacy of QQ under transient
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38 56 conditions has not been studied yet. An increase in organic loading rate (OLR), a typical
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41 57 transient condition, may trigger increased EPS production, causing EPS bulking and increased
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43 58 biofouling in MBRs (Abdollahzadeh Sharghi and Bonakdarpour, 2013; Shariati et al., 2013).
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46 59 This poses a challenge to the application of QQ for biofouling control under transient conditions,
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48 60 as QQ bacteria are believed to disrupt QS molecules, thus reducing EPS production and
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51 61 biofouling (Kim et al., 2013, 2015). Therefore, it is important to comprehensively investigate the
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53 62 factors which may contribute to membrane biofouling, and examine their changes in the presence
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56 63 of QQ bacteria under transient loads.

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4 64 This study aimed to investigate the impact of OLR on QQ-based biofouling control. To
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7 65 achieve this objective, a consortium of QQ strains comprised of AHLs- and AI-2-degrading
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9 66 bacteria, both in immobilized and suspension forms, was applied to MBRs. The influence of QQ
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11 67 bacteria on EPS production, sludge floc size, and sludge filterability in term of capillary suction
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14 68 time (CST) under both steady and transient organic loads was examined to gain a better
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16 69 understanding of the underlying relationship between QQ and biofouling. The concentrations of
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19 70 AHLs and AI-2 were also monitored to confirm the QQ effects.

21 71 **2. Material and Methods**

22 72 2.1. Chemicals

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24 73 The AHLs including N-butyryl-, N-hexanoyl-, N-heptanoyl-, N-octanoyl-, N-decanoyl-, N-
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26 74 dodecanoyl-, and N-tetradecanoyl-DL-homoserine lactone (C4-, C6-, C7-, C8-, C10-, C12- and
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29 75 C14-HSL); N-(3-oxohexanoyl)-, N-(3-oxooctanoyl)-, N-(3-oxododecanoyl)-, and N-(3-
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31 76 oxotetradecanoyl)-L-homoserine lactone (3OC6-, 3OC8-, 3OC12-, 3OC14-HSL) were obtained
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34 77 from Sigma-Aldrich (Singapore). The precursor of AI-2, 4,5-dihydroxy-2,3-pentanedione (DPD)
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36 78 was purchased from Omm Scientific, Inc (Dallas, TX, USA).

37 79 2.2. Experimental setup

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39 80 Experiment was conducted under varying (Phase I) and fixed OLR (Phase II) in duplicate.
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41 81 (Table 1) In Phase I four MBRs, each with a working volume of 450 mL; (i) control MBR (C-
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43 82 MBR), (ii) control MBR with empty beads (Cb-MBR), (iii) MBR with addition of QQ bacteria
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46 83 entrapped in polymer-coated alginate beads (QQb-MBR), and (iv) MBR with addition of QQ
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48 84 bacteria in broth (QQs-MBR) were set up and operated simultaneously. Under fixed OLR (Phase
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51 85 II), two more QQb-MBR were added in addition to the four MBRs; when C-MBR, Cb-MBR and
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54 86 QQs-MBR fouled, one of the QQb-MBR was stopped, and the level of EPS and other fouling
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87 parameters in both MBRs were analyzed to gain better understanding of the QQ mechanism
 88 along with the operation cycle. In Phase I, the OLR was gradually increased from 4 to ~ 7 mg
 89 COD/ g VSS • hr via wasting sludge from the MBRs; while in Phase II, the OLR was maintained
 90 constantly via supplementing the seed sludge.

91 Hollow fiber membranes with a pore size of 0.1 μm and surface area of 0.0045 m^2 were
 92 used in the present study. Sludge (collected from Ulu Pandan Water Reclamation Plant,
 93 Singapore) was acclimatized to a synthetic wastewater, comprising glucose (500 mg/L), NH_4Cl
 94 (190 mg/L), KH_2PO_4 (55.6 mg/L), CaCl_2 (5.5 mg/L), $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ (5.7 mg/L), FeCl_3 (1.5
 95 mg/L), MnCl_2 (1mg/L) and NaHCO_3 to maintain pH in the range of 7.0-7.5, for one month
 96 before being used as the seed sludge. During the study, the solids retention time (SRT) and
 97 hydraulic retention time (HRT) were maintained at 30 d and 12 h, respectively, at an operational
 98 flux of 8.5 $\text{L}/\text{m}^2/\text{h}$ (LMH) unless otherwise mentioned. A constant flux of 8.5 LMH was
 99 maintained to keep fouling at a reasonable rate.

100 **Table 1: Experimental setup and operating conditions**

	Phase I	Phase II
Experimental conditions	Varying Organic loading rate	Fixed organic loading rate
MBRs setup	i. Control MBR (C-MBR) ii. Control MBR with vacant beads (Cb-MBR) iii. MBR with QQ beads (QQb-MBR) iv. MBR with QQ sludge (QQs-MBR)	
Reactor volume (mL)	450	450
Membrane type	Hollow fiber, PVDF	Hollow fiber, PVDF
Flux (LMH)	8.5	8.5
HRT (h)	12	12
SRT (d)	30	90

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4 101 2.3. Bacterial immobilization

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6 102 A QQ consortium comprising of AHLs degrading strains *Enterobacter cloaca* (QQ13),
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9 103 *Microbacterium sp.* (QQ1), *Pseudomonas sp.* (QQ3), which were isolated from activated sludge
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11 104 in the authors' lab (the 16S rRNA gene accession number: KR058854, KR058848 and
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13 105 KR058846, respectively), *Rhodococcus sp.* RBH4 (Kim et al., 2015) and an AI-2 degrading *E.*
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15 106 *coli* strain Δ *lsrRAluxS* (Thompson et al., 2015) was selected for the study. Polymeric beads,
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17 107 entrapped with a single QQ strain, were prepared according to the method described by Kim et
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19 108 al. (2015) with some modifications. Briefly, fresh QQ bacterial culture, grown in Luria Bertanni
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21 109 (LB) broth with an optical density of 1.00 measured at 400 nm, was centrifuged at 4000 rpm for
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23 110 30 min and re-suspended in a phosphate buffer solution (PBS). For QQs-MBR, strain
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25 111 suspensions were directly added into the sludge, keeping sludge to volume ratio at 2%. Whereas
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27 112 for QQ beads, 5 mL of bacterial suspension was mixed with sterile sodium alginate (2% w/v)
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29 113 and the final suspension dropped into CaCl₂ solution (4% w/v) through a nozzle at a rate of 1
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31 114 mL/min using a peristaltic pump. For polymeric coating, pellets of polysulfone were dissolved in
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33 115 N-methyl-2-pyrrolidone (8% w/v) at 60 °C; finally, alginate beads were dipped in polymeric
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35 116 solution for 15 s and stored in deionized water at 4 °C. The QQ strains were entrapped separately
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37 117 in polymeric beads to avoid non-homogenous mixing, and added in equal proportions to MBRs
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39 118 keeping the overall ratio of beads to original MBR working volume at 2%.
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48 119 2.4. Membrane resistance analysis

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50 120 Membrane resistances were evaluated in terms of their fouling potential using a simple
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52 121 resistance in series model (Choo and Lee, 1996) based upon Darcy's Law:

$$R = \frac{\Delta P}{J\mu}$$

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4 122 where R is hydraulic resistance (1/m), ΔP is the TMP rise (Pa), J is the operational flux
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7 123 ($\text{m}^3/\text{m}^2/\text{s}$), and μ is the permeate dynamic viscosity (Pa.s).

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$$R$$

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11 125 where, R_t is the total hydraulic resistance, comprised of cake layer resistance (R_c), pore blockage
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14 126 resistance (R_p) and intrinsic membrane resistance (R_m). R_m was measured by filtering clean water
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17 127 through new membrane, while R_t was determined at the end of an experiment by measuring flux
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19 128 and TMP when deionized (DI) water was filtered through the membrane. The cake layer was
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22 129 then removed via sonication and the membrane submerged in deionized (DI) water to measure
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24 130 R_m+R_p . Finally, R_c was calculated by subtracting $R_m + R_p$ from R_t .

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26 131 2.5. Quantification of signal molecules

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29 132 AHLs were extracted from the biocake and activated sludge (50 mL) of the MBR as per
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31 133 the method described by Waheed et al. (2016). Samples were concentrated 100 times and
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34 134 analyzed using an Agilent 6460 (USA) QQQ triple quadrupole mass spectrometer (MS)
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36 135 equipped with an AJS electro-spray interface. The instrument settings were optimized as follows:
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39 136 interface voltage, 3.5 kV; drying gas temperature, 325 °C; drying gas (N_2) flow rate, 8 L/min;
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41 137 sheath gas temperature, 300°C; sheath gas (N_2) flow rate, 11 L/min. An Agilent 1290 Infinity II
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43 138 HPLC was used for chromatographic separation, which was achieved on a 100x2.1 mm Kinetex
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46 139 2.6 μm Bi-phenyl column (Phenomenex, USA) using 0.1% formic acid (A) and methanol with
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49 140 0.1% formic acid (B) as the mobile phase at a gradient of 5 to 85% B within 12 min. The limit of
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51 141 quantification (LOQ) for this method was 0.01 $\mu\text{g}/\text{L}$.

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53 142 One milliliter of sample was used to analyze the concentration of AI-2 in the sludge
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56 143 supernatant via the bioluminescence method using the reporter strain *Vibrio harveyi* BB170
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58 144 (Taga and Xavier, 2011). Briefly, the fresh culture of BB170 ($\text{OD}_{600} = \sim 0.3$ to 0.5) was diluted

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4 145 2000 times in a freshly-prepared AB medium. Duplicate mixture containing 10 % (v/v) sample
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6 146 along with 90 % (v/v) diluted BB170 culture was added to a microtitre plate and incubated at 30
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9 147 °C with rotary mixing at 150 rpm. The same amount of Milli-Q water was used as for negative
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11 148 controls. The luminescence produced in each sample was measured using a microtitre plate
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14 149 reader (Infinite M200 Pro, Tecan, Switzerland) at a wavelength of 490 nm. Measurements were
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16 150 recorded every 30 min from 5 to 8 h, until the negative control samples showed the lowest
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19 151 amount of luminescence. A precursor of AI-2, 4,5-dihydroxy-2,3-pentanedione (DPD) (Omm
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21 152 Scientific, Dallas, TX, USA), was used as the calibration standard.

23 2.6. Sludge characterization & water quality analysis

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26 154 The heat extraction method (Li and Yang, 2007) was used for extraction of extracellular
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28 155 polymeric substances (EPS). Briefly, 15 mL of sludge was collected from the MBRs, centrifuged
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31 156 at 4000 rpm and 4 °C for 30 min (this condition was also used for the following centrifugation
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33 157 steps unless otherwise stated), and supernatant stored as soluble EPS. The sludge pellet was re-
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36 158 suspended in 0.05 % (w/v) NaCl and homogenized; then the sample was centrifuged (4000 rpm
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38 159 for 15 min at 4 °C) and supernatant stored as loosely bound (LB) EPS. The sludge pellet was re-
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41 160 suspended in same NaCl solution and incubated at 60°C for an hour; after centrifugation the
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43 161 supernatant was stored as tightly bound (TB) EPS. For protein (PN) analysis, the Lowry method
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45 162 (Lowry et al., 1951) with the folin-ciocalteu phenolic reagent was used, and absorbance at 750
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48 163 nm was measured via a spectrophotometer (UV-2600, Shimadzu). For the quantification of
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51 164 polysaccharides (PS), the absorbance at 490 nm was measured using the phenol sulfuric acid
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53 165 method (DuBois et al., 1956).

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55 166 The dissolved oxygen (DO) and specific oxygen uptake rate (SOUR) were monitored using
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58 167 a multi meter (Fisher Scientific, XL600). The chemical oxygen demand (COD), sludge volume

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4 168 index (SVI), mixed liquor suspended solids (MLSS) and volatile suspended solids (VSS) were
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6 169 analyzed as per the Standard Methods (APHA, 2012). Sludge dewaterability was evaluated in
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9 170 terms of CST using a CST apparatus (Ofite 294-50, OFI Testing Equipment Inc., US). The
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11 171 sludge particle size distribution was monitored using a particle size analyzer (SALD-3101,
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14 172 Shimadzu).

16 173 **3. Results and discussion**

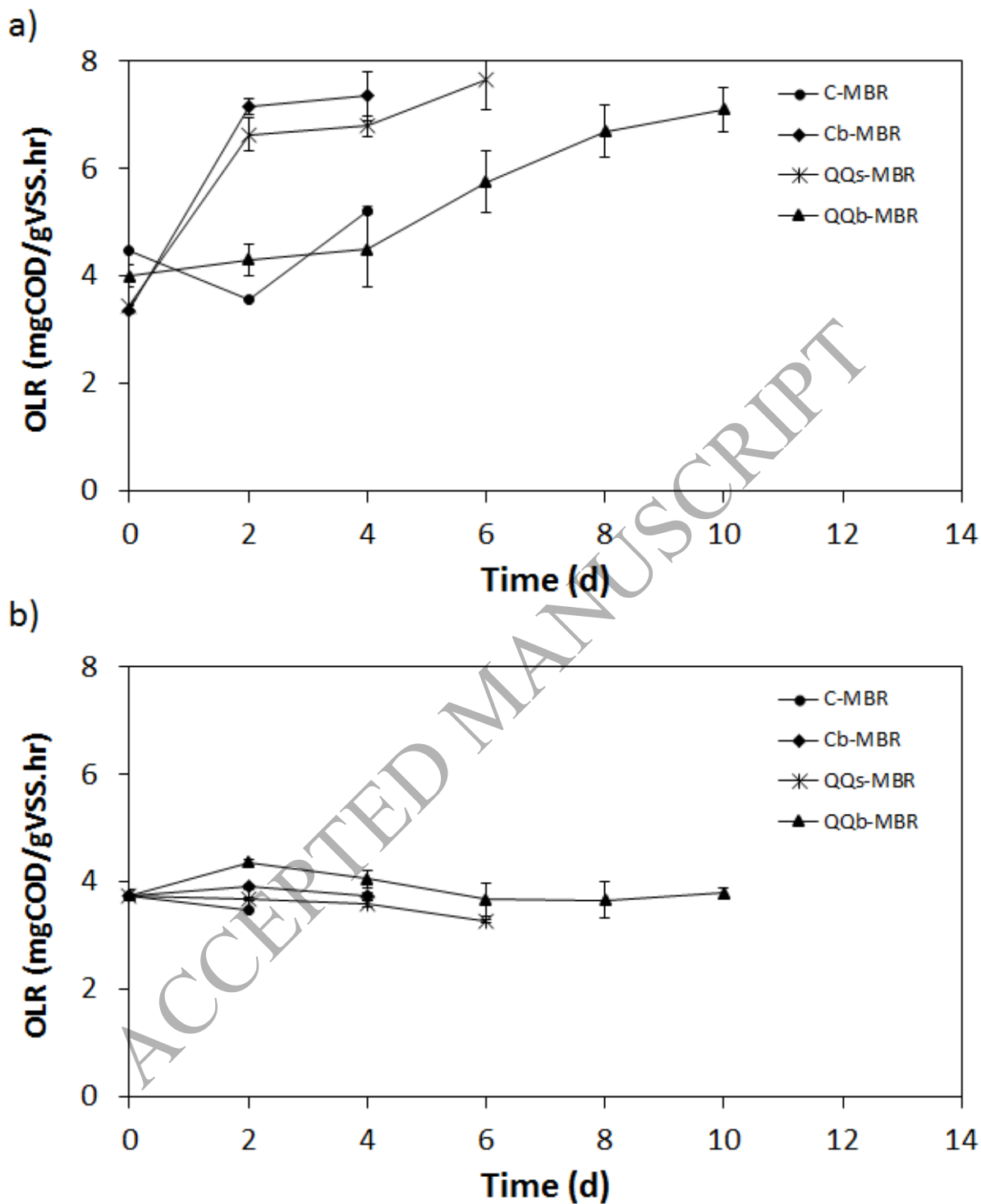
18 19 174 3.1. MBR operation and trans-membrane pressure (TMP) trends

21 175 The experiment was conducted in two separate phases, as described in Table 1. The DO
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23 176 levels in all reactors throughout the phases were maintained within 5.5 to 6.5 mg/L. The COD
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26 177 removal efficiency was more than 90 % in all MBRs (Table 2), and similar COD removal in the
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28 178 QQ MBRs indicate that QQ beads or sludge did not affect the performance of MBRs. A gradual
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31 179 decrease in mixed liquor volatile suspended solids (MLVSS) concentration was carried out
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33 180 purposely in Phase I via wasting sludge and controlling a SRT of 30 d, thus creating a continuous
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36 181 increase in the OLR. While in Phase II, all MBRs were supplemented with the seed sludge at
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38 182 regular intervals to maintain a constant MLVSS and OLR (Figure 1). To evaluate MBR
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41 183 performance, TMP was monitored for both experiments as the extent of membrane fouling is
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43 184 directly linked to TMP increase. Figure 2 shows the TMP profiles of MBRs of both Phases
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46 185 which were terminated when the TMP reached ~30 kPa. In Phase I (Figure 2a), no significant
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48 186 difference was observed between both control MBRs, as the overall fouling rates ($\Delta\text{TMP}/\Delta t$) for
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51 187 C-MBR and Cb-MBR were found to be 5.81 and 5.12 kPa/d, respectively. However, addition of
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53 188 QQ sludge or QQ beads in MBRs reduced the fouling rate by 29% and 49%, respectively, as
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56 189 compared to C-MBR under transient conditions. While in steady state (Phase II in Figure 2b),
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58 190 addition of QQ consortium, either in culture broth or in immobilized beads, significantly reduced
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4 191 the overall fouling rate by more than 66 % as compared to C-MBR, with the QQ beads slightly
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6 192 better than the QQ sludge. The vacant beads in Cb-MBR also reduced the fouling rate by 35%. It
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9 193 was found that OLR caused mainly external fouling due to accumulation of colloidal and
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11 194 dissolved organic molecules (Trussell et al., 2006), thereby, MBRs at low OLR during Phase II
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14 195 operated longer than MBR at high or changing OLR.
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16 196 **Table 2 Sludge characterization and performance efficiency of MBRs (n=3)**

	Phase I				Phase II			
	C- MBR	Cb- MBR	QQs- MBR	QQb- MBR	C- MBR	Cb- MBR	QQs- MBR	QQb- MBR
SOUR (mg/hr/g)	32.4	40.7	35.8	79.9	70.2	77.5	122.3	160.3
CST (s)	15.0	8.9	18.5	12.0	19.8	16.0	12.5	13.5
PSA mean (μm)	50.6	87.5	125.3	48.6	69.0	38.5	60.6	35.0
COD removal (%)	97.5	96.9	98.0	85.2	90.5	90.6	91.5	92.0



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198 **Figure 1 The organic loading rate (OLR) of (a) Phase I and (b) Phase II**

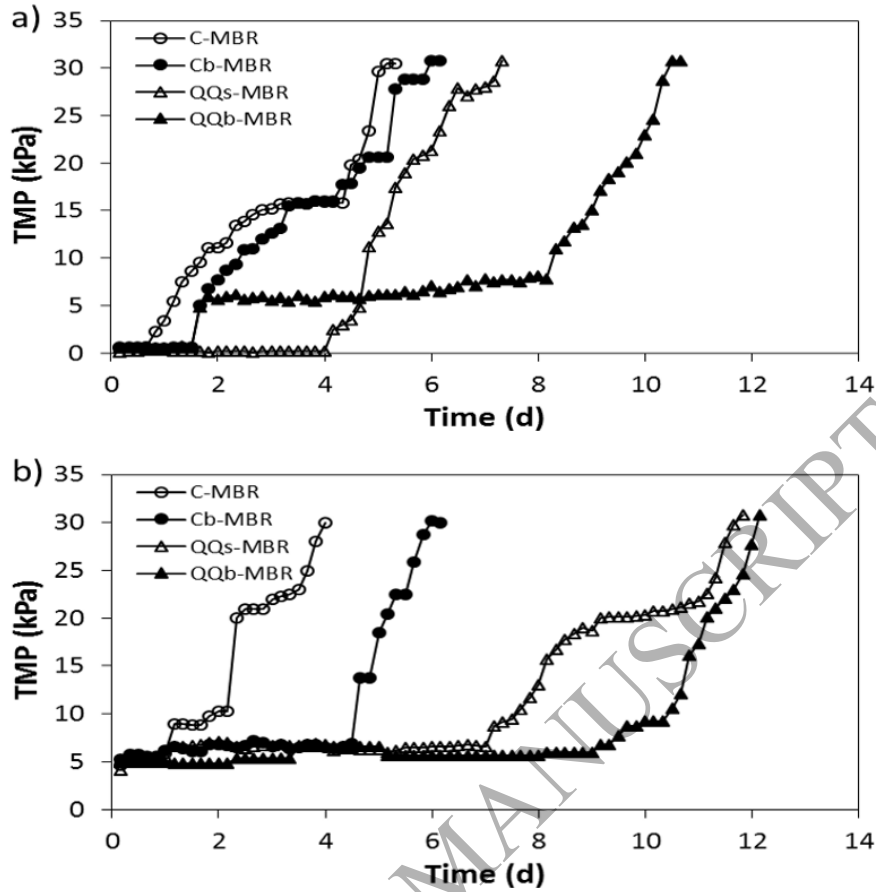


Figure 2 TMP profiles of (a) Phase I and (b) Phase II

The similar TMP profiles of QQs-MBR and QQb-MBR in Phase II suggest that the QQ consortium was effective in fouling control when the MBR was in steady state, irrespective of whether it is in broth or in immobilized beads. However, under transient conditions, robust fouling control was achieved only when the QQ consortium was in immobilized beads. QQ strains may grow well in immobilized form since beads provide protective environment under harsh conditions Whereas, activity/growth of QQ strains was suppressed when they were added directly to sludge, which could be due to the presence of predators or QQ enzyme suppressors released by other competitors. Previous studies (Gao et al., 2013; Zhang et al., 2006) demonstrated three dominant stages in overall TMP profiles, including an initial rise due to pore

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210 blockage (stage 1), a slow rise due to accumulation of EPS on the membrane surface (stage 2),
211 and a sharp jump resulted from biofilm formation (stage 3). In this study, a slower rise in TMP
212 during stage 2 in the QQ MBRs compared to control MBRs was observed for both Phases,
213 indicating that QQ may have played a role in retarding biofouling by postponing cake layer
214 formation.

215 3.2. Membrane resistance and foulant characterization

216 Membrane resistances from all six reactors of Phase II were compared to further elucidate
217 the mechanism of QQ (Figure 3). A clear decrease in the fraction of pore blockage resistance
218 (R_p/R_t) with time was observed in QQb-MBR, from >70% on Day 4, to ~ 45% on Day 6, and
219 further to ~ 40% on Day 12 when it was fouled. The fraction of cake layer resistance (R_c/R_t)
220 remained very low (<7%) on Day 4 and 6, but increased to ~30% on Day 10. This clearly
221 indicates the strong inhibition of cake layer formation by QQ in the early stage. Compared to
222 other MBRs, the fraction of pore blockage resistance (R_p/R_t) of both QQ MBRs (>40%) were
223 the greatest, when they were fouled. This suggests that more molecules or small particles had
224 blocked the membrane pores in QQ MBRs, as the QQ had postponed the cake layer formation.
225 Hence in QQ MBRs, the pore blockage may be the leading factor that had led to the final
226 fouling.

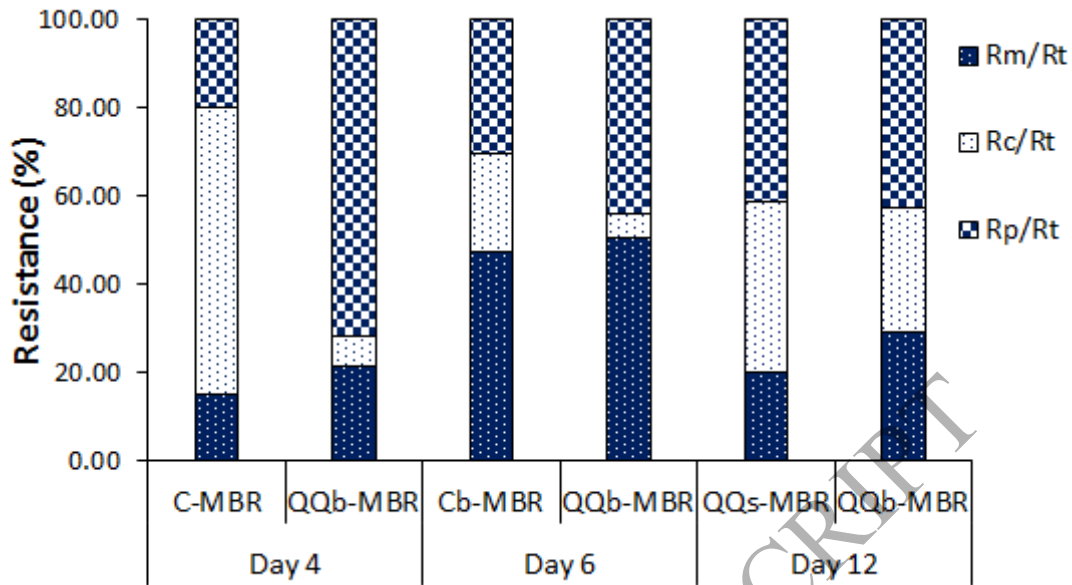


Figure 3 Membrane resistance analysis under steady conditions

The significantly higher portion of cake layer resistance (R_c/R_t) in C-MBR as compared to Cb-MBR indicates that beads scouring had greatly reduced the cake layer formation. This was also observed when QQs-MBR was compared to QQb-MBR.

Fouled membranes from all reactors were analyzed via Fourier transform infrared (FTIR) spectroscopy (AIM-8800, Shimadzu) to detect any foulant composition changes posed by quenching mechanism. QQ strains either immobilized or added as a suspension did not modify the membrane foulants, as the organic or functional groups in all membranes were found to be same, as shown in Figure S2 in the Supplementary Information. It can be inferred that quenching activity delayed biofilm formation, but kept the foulant composition intact.

3.3. Effect of QQ on QS signal molecules

Biofilm formation (biofouling) could be mitigated substantially by disrupting signal molecules such as AHLs, oligopeptides or AI-2 using enzymes released by QQ bacteria (Shrout

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and Nerenberg, 2012). In the present study, the role of QQ bacteria was confirmed by monitoring the concentrations of two signal molecules, AHLs and AI-2 using LC-MS and *V. harveyi* bioluminescence assays, respectively.

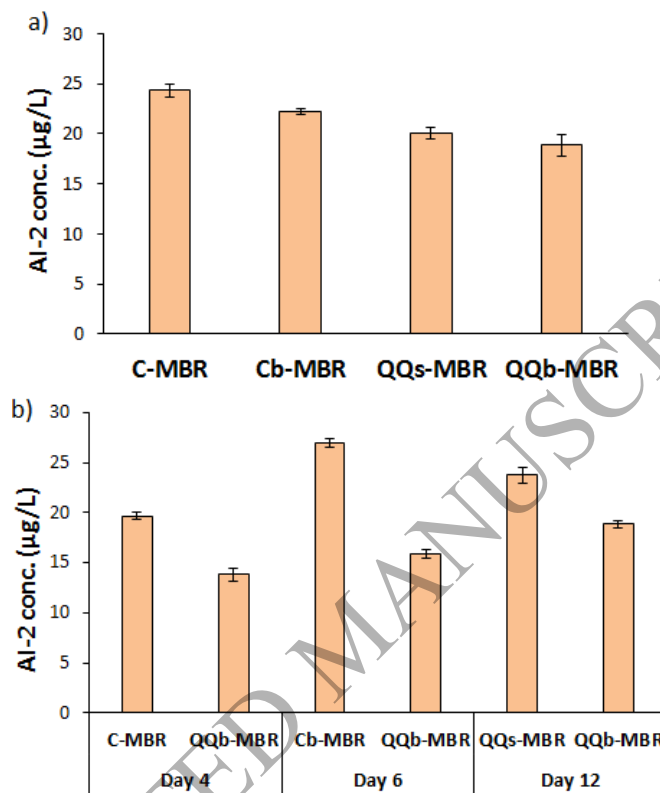
3.3.1. Quenching of AHLs

AHLs in the seed sludge were analyzed to screen the major types of QS signal molecules; C4-HSL, 3OC6-HSL, 3OC8-HSL and 3OC12-HSL were detected as the dominant AHLs present. Table S1 presents the detailed quantitative results of AHLs in the biocake, mixed liquor and effluents from each reactor. Various AHLs including C4-, 3OC6-, C6- and 3OC14-HSL were detected in the mixed liquor of C-MBR and Cb-MBR, whereas they were not detected or lower than the quantification limit in QQs-MBR or QQb-MBR. This indicates that of the QQ consortium had quenched these QS molecules, thus reducing the biofilm formation. High concentrations of C4-HSL and 3OC6-HSL in the biocake sample of C-MBR and Cb-MBR as compared to that in QQ MBRs may signify the role of small chain signal molecules in the formation of matured biofilms.

3.3.2. AI-2

The AI-2 concentration was determined by bioluminescence assay with 4,5-dihydroxy-2,3-pentanedione (DPD), a precursor of AI-2, as the calibration standard. The effectiveness of QQ bacteria in reducing the AI-2 signal molecules is shown in Figure 4. The AI-2 concentrations in QQb-MBR were the lowest when all MBRs were fouled in both Phases. However, the AI-2 in QQs-MBR of Phase II was higher than that of C-MBR, though it was still lower than both C-MBR and Cb-MBR in Phase I. Higher concentrations of AI-2 in QQs-MBR could be due to the less survival rate of the *E. coli* strain *ΔlsrRΔluxS*, since the cells were not protected in the beads. A slow but gradual increase in AI-2 concentrations with time in QQb-MBR was also noticed.

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4 264 These may suggest that the QQ consortium were effective in quenching AI-2, but only at the
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6 265 early stage; with time the quenching effect vanished. Their quenching effect was more robust
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9 266 when they were immobilized in beads.



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268 **Figure 4 Level of autoinducer (AI-2) in the mixed liquor of MBRs, (a) Phase I and (b)**
269 **Phase II**

270 3.4. Effect of QQ on sludge characteristics

271 The mixed liquor was sampled regularly to monitor the effect of QQ on the different
272 sludge characteristics as listed in Table 2. The SOUR was measured as a representative of sludge
273 respirometric activity, and Phase II showed higher activity compared to the Phase I. The SOUR
274 of QQb-MBR was almost 2 times as many as that of control MBRs in both Phases. This shows
275 that QQ phenomenon or disruption in bacterial signaling had a direct impact on metabolic

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4 276 activity of microbial cells; which could be related to starving conditions developed during
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7 277 interruption in bacterial communications, or dominance of strains those require very high OUR
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9 278 (Garcia-Ochoa et al., 2010; Gomez et al., 2006).

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11 279 Mean floc size for the QQb-MBR was found to be 49 and 35 μm for Phases I and II
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13 280 respectively. Overall, the sludge floc sizes in Cb-MBR and QQb-MBR were smaller than C-
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15 281 MBR and QQs-MBR in both Phases, respectively; the presence of beads in the former two
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17 282 MBRs contributed to this reduction in floc size. The dewaterability of sludge determined by CST
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19 283 is also an indication of its permeability in the MBR (Table 2), and a high EPS content will
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21 284 increase sludge viscosity thus reducing its permeability. However, no clear impacts of the QQ
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23 285 consortium on CST could be observed. Reduction in floc size still resulted in the better sludge
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25 286 dewaterability, however, its effects on sludge settling property cannot be ignored.

31 287 3.5. QQ effects on EPS production

33 288 3.5.1. EPS concentrations in the biocake

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36 289 EPS, either in bound or soluble form, is considered to be the major component in cell flocs
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38 290 and biofilms which ultimately leads to membrane fouling in MBRs. In this context, the soluble
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40 291 EPS (S-EPS), loosely bound EPS (LB-EPS) and tightly bound EPS (TB-EPS) extracted from the
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42 292 mixed liquor and sludge cake were analyzed for all phases. The protein and carbohydrate
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44 293 distributions in different EPS fractions extracted from the sludge cake is shown in Figure 5. EPS
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46 294 fractions retained on the membranes in the form of a fouling layer led to a high resistance on the
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48 295 membrane surface, and a reduction in permeate flux (Lin et al., 2014). In the biocake sample,
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50 296 bound EPS was higher than soluble EPS in all phases. Bound EPS facilitates more in floc
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52 297 formation, and an increase in floc size tends to enhance floc deposition on the membrane surface,
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54 298 which ultimately leads to biocake formation (Chu and Li, 2005).

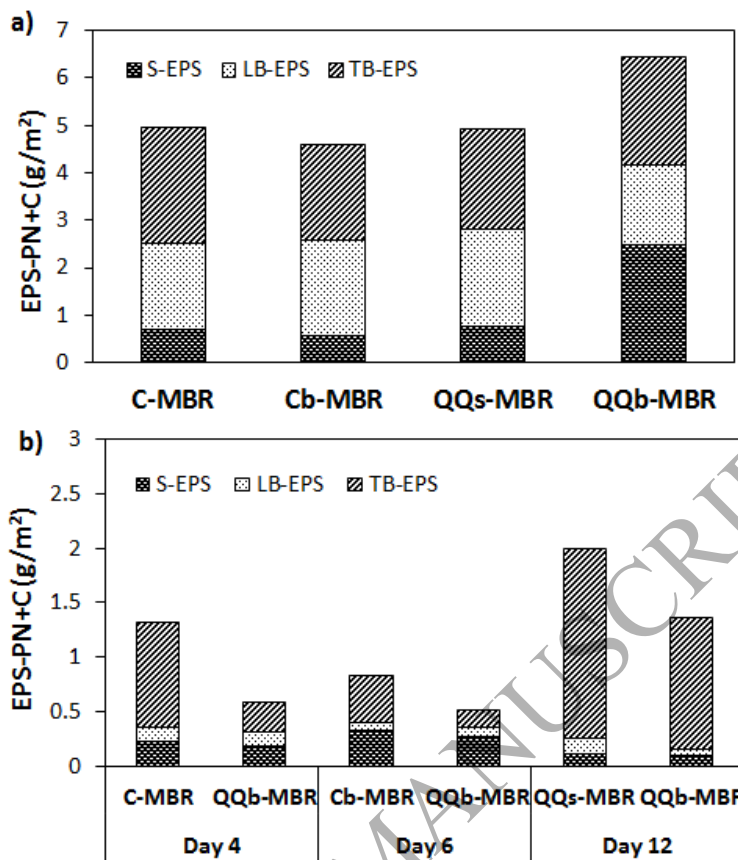


Figure 5 EPS concentration in the biocake samples (a) Phase I and (b) Phase II

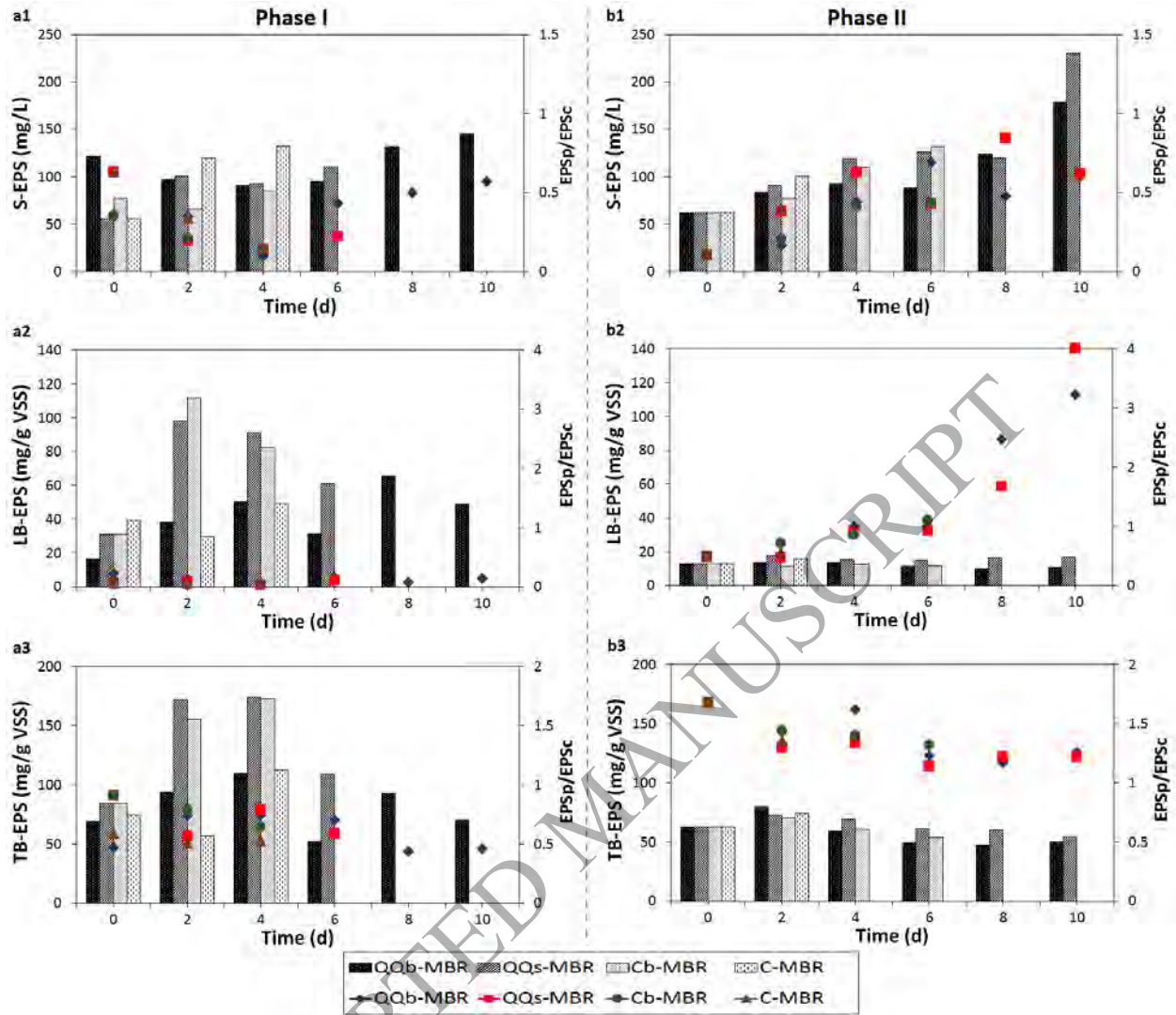
The total biocake EPS in Phase I was more than 3 times higher than Phase II, which indicates that the change in OLR had a dominant effect on the total biocake EPS content. However, the TB-EPS were in the same range ($\sim 0.95\text{-}2.2\text{ g/m}^2$) amongst all MBRs in both phases except a little lower amount was observed in Cb-MBR of Phase II (Figure 5). The high S-EPS and LB-EPS contents in Phases I could have resulted from new or loose sludge flocs being deposited on the surface of the compacted biocake. It also suggests that the scouring effects of the beads were minimal for fouling control, which corresponded well to the TMP profiles in Figure 2.

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309 The total and especially the TB-EPS in QQb-MBR were much lower on Day 4 and 6 as
310 compared to the control reactors of the same day (Figure 5b), but they increased to levels higher
311 than the control reactors when it was fouled. This corresponded well with the membrane
312 resistance analysis.

313 3.5.2. EPS concentrations in the mixed liquor

314 Figure 6 shows the protein-carbohydrate ratio (EPS_p/EPS_C), as well as soluble and bound
315 EPS concentrations at different time intervals in the mixed liquor. When the MBRs were started,
316 the bound-EPS (more dominantly for LB-EPS) contents of all MBRs in Phase II were lower than
317 those of Phase I, which could be the reason of the higher SOUR. The prominent increases (from
318 Day 0 to Day 2) in LB- and TB-EPS in Cb-MBR and QQs-MBR of Phase I corresponded well to
319 the significant decreases in MLSS in the MBRs (Figure S1). But after that, both LB-EPS and
320 TB-EPS in the two MBRs gradually decreased. The LB- and TB-EPS in QQb-MBR also
321 increased since the start of the MBR, but started to decrease from Day 8 and 4, respectively. The
322 initial increase could be partially due to the decrease in MLSS concentrations, which led to an
323 increase in OLR and severe biofouling. Similar results were also reported in sludge granulation
324 (Kim et al., 2013) and MBR operation (Trussell et al., 2006).



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Figure 6 Profiles of extracellular polymeric substances (EPS) in terms of the total concentrations of polysaccharide and protein (histograms) and the ratio of protein to polysaccharide (scattered-dots) in the mixed liquor of (a) Phase I and (b) Phase II. From top to bottom are the soluble EPS (S-EPS), loosely bound EPS (LB-EPS) and tightly bound EPS (TB-EPS), respectively.

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In Phase II, decreases in LB- and TB-EPS in all MBRs were clearly seen from the beginning of the operation. However, no significant differences between the QQ MBRs and the

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333 control MBRs were observed. Instead, the S-EPS gradually increased in all MBRs in both Phases
334 except of a slight decrease in QQb-MBR in Phase I before Day 4. Many studies have directly
335 correlated the S-EPS with biomass concentrations (Drews et al., 2008; Geng and Hall, 2007;
336 Jeong et al., 2007). In our study, the MLSS levels dropped substantially in Phase I; however,
337 continuous rise in S-EPS were still observed. These results suggest that intensive aeration and
338 fluidization of polymeric beads significantly reduced the sludge floc size (Table 2), and the
339 dissociation of sludge flocs had resulted in the decrease in bound EPS and increase in S-EPS and
340 eventually contributed to membrane pore blockage (Meng et al., 2009; Metzger et al., 2007).

341 The ratio of EPS_P (EPS-protein) to EPS_C (EPS-carbohydrate) was also investigated to gain
342 better understanding of QQ mechanism. In Phase I, all MBRs showed a decrease in EPS_P/EPS_C
343 ratio of the S-EPS with an increase in OLR, except for that of QQb-MBR, which slightly
344 increased (Figure 6:a1). In contrast, a clear increase in the EPS_P/EPS_C in the sludge supernatant
345 was observed in Phase II (Figure 6:b1). The gradual increase in EPS_P/EPS_C of the LB-EPS was
346 also observed in all MBRs in Phase II, while it remained unchanged in Phase I. However, for
347 TB-EPS, a slight decrease in EPS_P/EPS_C was observed in both Phases. These results suggest that
348 the increase in OLR had more impacts on LB-EPS and S-EPS; it had maintained the high
349 fraction of polysaccharides in the LB-EPS, which contributed to fouling more than protein-like
350 substances (Rosenberger et al., 2006; Yigit et al., 2008). The EPS_P/EPS_C ratio in the LB-EPS
351 corresponded well to the higher concentrations of polysaccharides than protein in the biocake in
352 Phase I, and the higher protein contents in Phase II. This suggests that LB-EPS were more
353 significantly correlated with membrane fouling compared with TB- and S-EPS, as they
354 contributed to flocculation and sedimentation processes, which was also reported in literatures
355 (Lin et al., 2014; Ramesh et al., 2007; Wang et al., 2009).

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4 356 The biofouling was deferred by factors of 3.04 ± 0.5 and 2.96 ± 0.7 (mean \pm standard
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6 357 deviation, $n=2$) in QQb-MBR and QQs-MBR, respectively, compared to the C-MBR for Phase
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9 358 II. However, these factors decreased to 1.8 ± 0.3 and 1.4 ± 0.1 in Phase I when the OLR kept
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12 359 increasing. A continuous increase in OLR, which resulted in significant EPS production and
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14 360 higher content of polysaccharide in LB-EPS, was the major factor resulting in the impaired
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16 361 efficiency of immobilized QQ strains. The reduction in S-EPS concentration in the mixed liquor
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19 362 in MBRs containing QQ bacteria were reported previously (Lee et al., 2016; Maqbool et al.,
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21 363 2015). However, it was not observed in our study; instead, the QQ consortium retarded the EPS
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24 364 accumulation on the membranes.

26 365 **4. Conclusions**

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29 366 • The QQ consortium in immobilized beads showed robust fouling control capacity under
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31 367 both steady state and transient condition. When added in broth, they remained effective
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33 368 only under steady state.
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36 369 • The QQ consortium significantly reduced the AHL levels in mixed liquor inside MBR
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38 370 and in sludge cake on membrane surface.
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41 371 • LB-EPS of the mixed liquor was the major contributor to membrane fouling, and within
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43 372 the EPS, polysaccharides contributed more to membrane fouling than proteins.
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46 373 • A continuous increase in OLR resulted in higher EPS levels and higher polysaccharide
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48 374 ratios in the LB-EPS, which in turn lead to faster fouling in the MBRs.

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58 378 and Environment & Water Industry Programme Office (EWI), SEO Nanyang Technological
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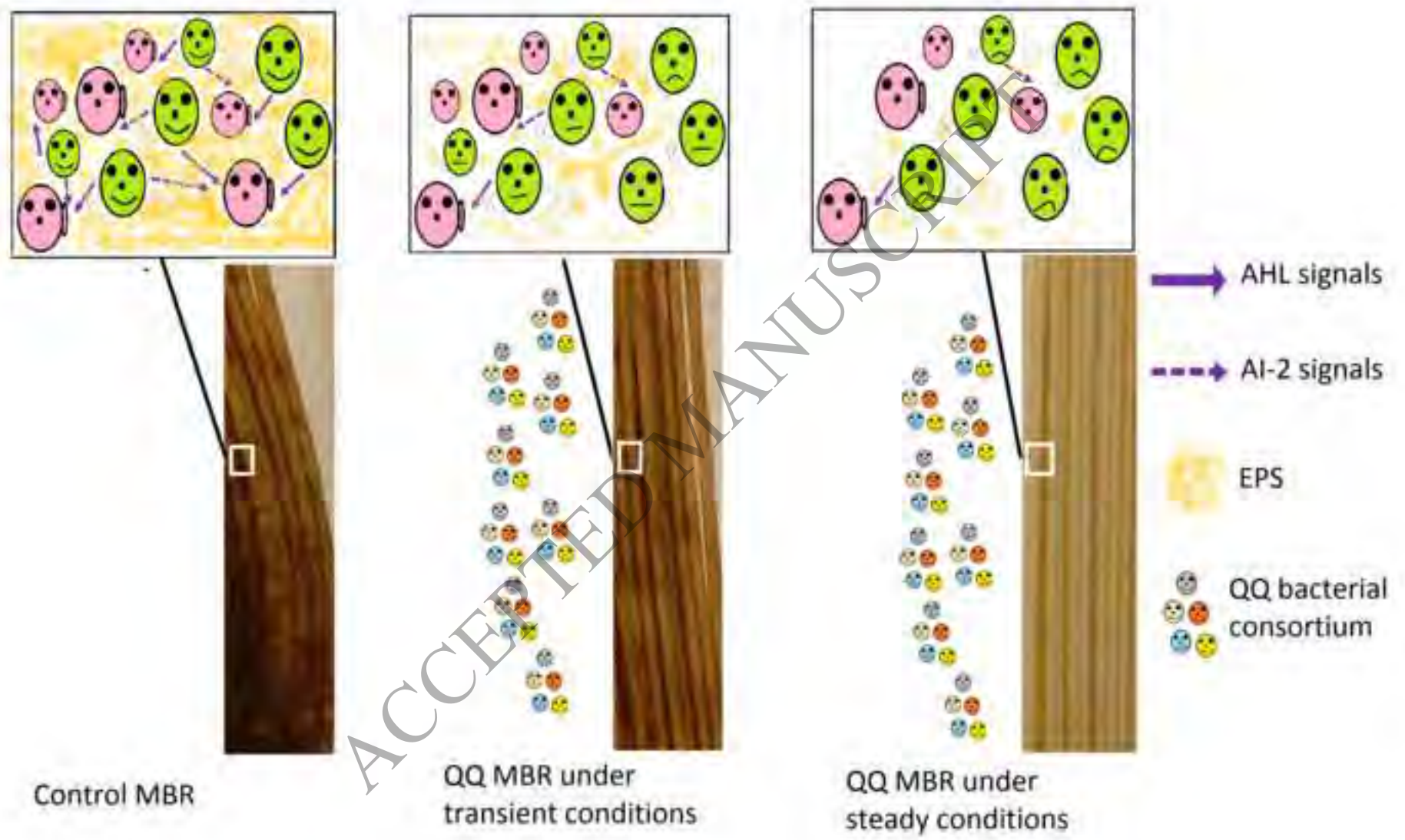
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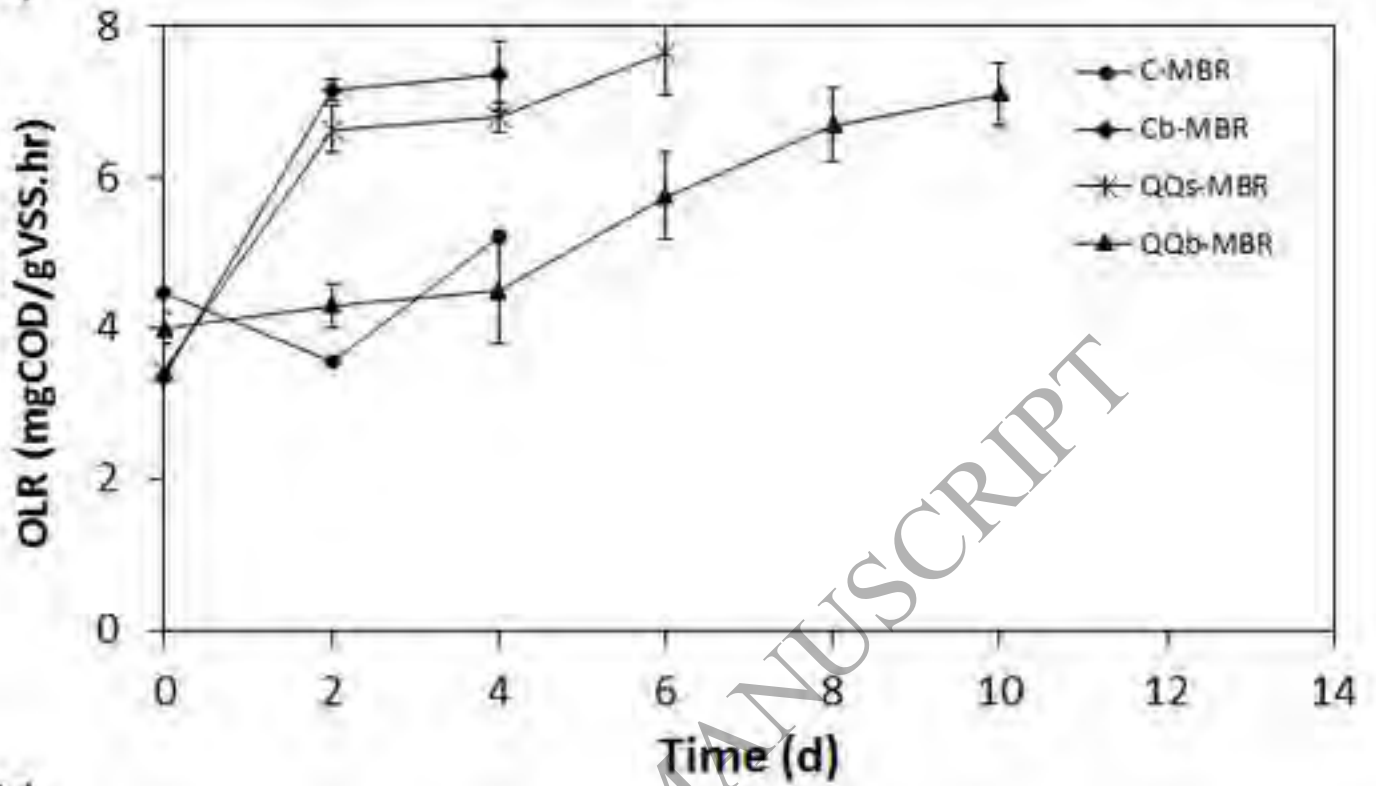
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