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1	Aggregation of low-concentration dirhamnolipid biosurfactant in
2	electrolyte solution
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32 Abstract:

Dynamic light scattering (DLS) and cryo-transmission electron microscopy 33 34 (Cryo-TEM) tests demonstrated aggregate formation for dirhamnolipid biosurfactant (diRL) at concentrations lower than surface-tension-based critical micelle 35 36 concentration (CMC_{st}). Increase of diRL concentration and solution pH results in 37 decrease of the aggregate size at diRL concentration below CMCst, whereas it has no 38 influence at diRL concentration above CMCst. The cryo-TEM micrographs show spherical morphology of aggregates, and logarithm of aggregate size follows Gaussian 39 40 distribution. The aggregates are negatively charged. Zeta potential of the aggregates 41 decreases with increase of diRL concentration to CMCst, and stabilizes at diRL concentrations higher than CMCst. Increase of solution pH causes decrease of zeta 42 43 potential. A transitional state assumption is raised for interpretation of the diRL 44 aggregation behavior. The results demonstrate formation of aggregates at significantly 45 low diRL concentrations, which is of importance for cost-effective application of 46 rhamnolipid biosurfactant.

47 Keywords: dirhamnlipid; aggregation; critical micelle concentration; zeta potential.

48 **1 Introduction**

49 Biosurfactants are surfactants produced by microbes. Due to expanding application of biosurfactants in many fields, e.g. biomedicine and bioremediation, 50 their aggregation behaviour in electrolyte has received much attention in recent years. 51 ¹⁻³ The aggregates have a variety of microstructures, including spherical, globular or 52 cylindrical micelles, ^{1,4-8} spherical or irregular vesicles,^{2,9,10} tubular or irregular 53 bilayers,¹¹ and lamellar sheets.^{3,12,13} Also, the lyotropic liquid crystalline phases with 54 lamellar, hexagonal and cubic aggregate morphologies are observed at high surfactant 55 concentrations¹⁴. The morphology of these aggregates has been demonstrated to be 56 affected by surfactant concentration,^{8,15} pH,^{4,10,16} temperature,¹² counterions,^{1,10,15} and 57 ionic strength.¹⁷ 58

59 Rhamnolipid is the most widely studied biosurfactant and its aggregates exhibit versatile structures at concentrations higher than the critical micelle concentration 60 (CMC). For example, Ishigami et al. investigated the effect of solution pH on 61 rhamnolipid aggregate structure at concentrations of 500-20000 mg/L in phosphate 62 buffered saline solution. The result showed that the aggregates existing in form of 63 bilayers vesicle at pH of 4.3-5.8, bilayer lamella with pH rising to 6.0-6.5, and 64 micelles with further increase of pH to 6.8.¹⁸ Champion et al. determined the 65 rhamnoliopid aggregate morphology at various pH at the concentration of 60 mM by 66 Cryo-TEM. The results show that aggregate phase transitioned in an order of bilayer 67 lamella, large vesicles, small vesicles and micelle with the increase of pH.⁴ In 68

addition, transformation of dirhamnolipid aggregations from large particles into small
 particles with the increase of concentration at fixed pH has also been reported.¹⁹

71 All these prior researches, however, were almost implemented with surfactant concentrations far higher than CMC. However, there are studies showing that 72 rhamnolipid exhibited excellent HOC-solubilizition activity at significantly low 73 concentrations. For example, rhamnolipid can enhance the solubility of hexadecane 74 75 and octadecane by 3~4 orders of magnitude at concentrations lower than CMC determined by surface tension method, and such solubilization efficiency is much 76 CMC.^{16,20} concentrations Hypothetically 77 higher than at above these HOC-solubilization activities of rhamnolipid surfactant may be related to its 78 aggregation behavior at low concentrations, e.g. lower than CMC. Furthermore, signs 79 of aggregate formation at concentrations lower than CMC for multi-component 80 rhamnoliplids were observed using dynamic lighter scattering method.^{7,15} Formation 81 of premicelles for a variety of surfactants also have been reported.²¹⁻²⁴ These 82 observations indicate the probability of sub-CMC aggregate formation for 83 rhamnolipid, which still remains unexplored. 84

In this study, the aggregation behavior of dirhamnolipid in phosphate buffered electrolyte solution with concentrations near surface-tension-based CMC (or CMC_{st}) was investigated. The objective of this study is to examine whether rhamnolipid forms aggregate at concentrations below CMC_{st} , and to explore the effect of solution conditions on aggregate formation at low rhamnolipid concentration range.

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91 **2 Materials and methods**

92 **2.1 Biosurfactant and chemicals**

The dirhamnolipid biosurfactant was produced, extracted, purified, and characterized using the method described by Zhong et al.²⁵ NaOH and HCl (analytical chemistry grade) were purchased from Damao Chemical (Tianjin, China). All other chemicals (NaNO₃, KH₂PO₄, Na₂HPO₄·12H₂O, MgSO₄, FeSO₄·7H₂O) were analytical chemistry grade with purity > 99% and purchased from Sinopharm (Beijing, China).

99 **2.2 Determination of CMC**_{st}

The stock solution of the diRL were prepared in phosphate buffered saline 100 solution (PBSS, NaNO₃ 2g/L, KH₂PO₄ 1.5g/L, Na₂HPO₄ 12H₂O 1.5g/L, MgSO₄ 101 102 0.1g/L, FeSO₄·7H₂O 0.01g/L). PBSS solutions of diRL in a series of concentrations were prepared using dilution method. Surface tension of diRL solutions was measured 103 at 30°C with surface tensiometer (JZ-200A, Chengde, China) using the Du Noüy 104 Ring method. CMCst of diRL was obtained from the relation of surface tension and 105 diRL concentration using the scheme described by Yuan et al.²⁶ Electrical 106 107 conductivity of diRL solutions was measured with DDS-11A Conductivity Meter 108 (Shengci, Shanghai, China).

109 2.3 Hydrodynamic aggregate size and zeta potential.

pH of PBSS solutions of diRL was adjusted to 8.0 with 20% NaOH solution
using a capillary glass pipe. High concentration of NaOH was used to minimize
change of solution volume during pH adjustment. These solutions were then filtered

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113 through a 0.22 µm filter (Millex-HV, Millipore, Billerica, Ma, US) to remove 114 suspended solid particles that may interfere with the measurement. Results of 115 preliminary test showed that the size of rhamnolipid aggregates is far less than 116 $0.22\mu m$ at pH 8.0, so they will not be retained by filtering. The solutions were allowed to stand still for 2 h. Then pH of the solutions was adjusted in sequence to 117 118 7.5, 7.0, 6.5, or 6.0 with 20% hydrochloric acid. For each sample, aggregate size and 119 zeta potential were measured using Zetasizer Nano ZEN3600 (Malvern Instruments, 120 U.K.).

121 The aggregate particle size was determined based on dynamic light scattering (DLS) mechanism using He-Ne laser at wavelength of 623nm and working power of 122 123 4.0 mW. 1 ml of the sample was loaded to the DTS-0012 cell and measured at 124 temperature of 30°C. The scattered light was collected by receptor at angle of 173° 125 from light path. A mean size provided by DTS Nano software (Malvern Instruments, 126 U.K.) was used to represent the aggregate size of the sample. Also, the number-based 127 particle size distribution (number PSD) data generated by the software were used for 128 the statistical analysis of aggregate size.

The zeta potential measurement is based on the mechanism of particle electrophoresis in aqueous solution. 1 mL sample is loaded to DTS 1060 folded capillary cell and the electrophoretic mobility of the aggregate was measured at 30°C under automatic voltage using a laser Doppler velocimetry with M3-PALS technique to avoid electrossmosis. The measured data was converted into corresponding zeta potential by applying the Helmholtz-Smoluchowski equation.²⁷

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DiRL solution or electrolyte solution in the absence of diRL in volume of 4uL 136 was placed on the grid with a holy polymer film using a microsyringe, and then sent 137 138 to a FEI Vitrobot sample plunger system (FEI, Hillsboro, Oregon). Excessive sample was removed by a filter paper. Then the sample grid was rapidly vitrified in liquid 139 140 ethane and transferred to a liquid nitrogen bath. The morphology of diRL aggregate 141 were then viewed and photographed on a Tecnai F20 Transmission Electron 142 Microscope (FEI, Hillsboro, Oregon) at an acceleration voltage of 120 kV. Nano 143 measurer 1.2.5 software (Shanghai, Fudan University) was used to process the micrograph images. The program marked the recognized particles in an image with 144 145 circles. The diameter of every circle was measured by pairing the circle to a screen 146 ruler calibrated by the reference bar in the image and used as the size of the particle. 147 In order to obtain statistically representative sample for aggregate size distribution analysis, the size data were collected on more than 100 or 200 particles from multiple 148 149 images for rhamnolipid concentrations of 25 or 250 µM, respectively.

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151 **3 Results and Discussion**

For all the pHs, surface tension of the solution decrease significantly with the increase of rhamnolipid concentration at low surfactant concentrations, and then further increase of surfactant concentration has no significant effect on surface tension (Fig. 1). Based on the method of Yuan et al.,²⁶ the CMC_{st} values obtained are 62, 78, 82, 83 and 82 μ M for pH of 6.0, 6.5, 7.0, 7.5 and 8.0, respectively. The result showed

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157 that the increase of solution pH resulted in increase of diRL CMC_{st} for pH not higher 158 than 7.0. The electrical conductivity of diRL solution increased with the increase of diRL concentration for all pH conditions, however, the two-line profile with a 159 160 distinguishable slope inflection is not observed for any of the curves (Fig. S1a, SI). 161 The plot of conductivity derivative versus diRL concentration is presented in Fig. 162 S1b, SI. The conductivity derivative shows a gradual decrease at the concentration 163 below CMC_{st}, which is in contrast to an abrupt decrease at CMC general for regular 164 surfactants.

165 Results of DLS-size measurement show that diRL aggregates were detected at diRL concentration both below and above CMCst. The number PSD profiles generated 166 167 by Malvern DTS Nano software show only one peak for all the conditions of measurements (typical profiles are presented in Fig. S2, SI), indicating presence of 168 169 only one type of aggregate. The influence of diRL concentration and solution pH on aggregate size is shown in Fig. 2a. The aggregate size is in a range of 8 to 72 nm. 170 171 When the solution pH is not higher than 7.0, the aggregate size decreased with the 172 increase of diRL concentration up to 100 µM. At diRL concentrations ranging from 10 to 100 μ M (close to CMC_{st}), the aggregate size decreases rapidly with increase of 173 174 pH. When diRL concentration is higher than 100 μ M, both diRL concentration and pH have no observable influence on the aggregate size. The relation between DLS 175 176 diffusion coefficients and diRL concentrations is shown in Fig. S3, SI. The diffusion coefficient increases with increase of diRL concentration when the concentration is 177 178 lower than CMC_{st}. This result is in contrast to DLS diffusion coefficient for regular

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surfactants, for which an abrupt decrease of the coefficient is observed at CMC.²⁸
This result, however, matches with the result of size measurement in that diffusion
coefficient is larger for smaller particles.

Aggregate zeta potential variation with diRL concentration and pH is presented 182 in Fig. 2b. Because rhamnolipid is an anionic surfactant with carboxyl group in the 183 hydrophilic moiety of the molecule, dissociation of the carboxyl groups yields 184 185 negatively-charged aggregate surface and hence negative zeta potential of the 186 aggregates. For all the pHs, the zeta potential decreases significantly the increase of 187 diRL concentration from 25 μ M to 100 μ M. Further increase of concentration has minimal influence. For all the diRL concentrations tested, increase in solution pH 188 causes decrease in aggregate zeta potential. Increase of solution pH results in 189 190 enhanced dissociation of diRL carboxyl group, which in turn increases the aggregate 191 surface charge density and lowers zeta potential (provided a dissociation equilibrium constant of 10^{-5.6} for rhamnolipid carboxyl group at room temperature.¹⁸ the 192 dissociation rate of the rhamnolipid is 71.5, 88.8, 96.2, 98.8, 99.6% at pH of 6.0, 6.5, 193 194 7.0, 7.5, 8.0, repectively).

195 25 μ M (below CMC_{st}) and 250 μ M (above CMC_{st}) of diRL solution at pH of 6.0 196 or 8.0 were examined using cryo-TEM. Typical images of the aggregates are 197 presented in Fig. 3. Aggregates are observed for all the four conditions, which is in 198 contrast to the observation in the absence of diRL for which no aggregates are 199 observed (Fig. S4a, SI). The morphology of the aggregates is spherical with minimal 1200 transparency, indicating micelle-type structure. Other aggregate structures reported in 201

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literatures at relatively high rhamnolipid concentrations, e.g. vesicles, lamella or
microtubes, ^{4,8,9,19} are not observed for any of the conditions tested. This is consistent
with the result of DLS size measurement that only one type of aggregate is observed.
The cryo-TEM result further confirms formation of diRL aggregates at concentrations
below CMC. On the other hand, the DLS size and cryo-TEM results also shows that
the aggregates are not premicelles, which are defined as dimers and low-aggregation
number aggregates of surfactant molecules before micelle formation. ²¹⁻²⁴
All the cryo-TEM images used for aggregate size distribution analysis are shown
in Fig. S4, SI. Gaussian distribution is commonly used to depict natural phenomena

209 in Fig. S4, S associated with real-valued random variables whose distributions are unknown. The 210 211 distributions of aggregate sizes obtained with either DLS or cryo-TEM method appear 212 to deviate from Gaussian distribution (data not shown), however, natural logarithm of 213 the sizes follows Gaussian distribution very well for all the four conditions examined 214 (Fig. 3). Values of the parameters for the fit are presented in Table 1. The mean of cryo-TEM size at diRL concentration of 25 μ M (lower than CMC_{st}) is larger than at 215 216 diRL concentration of 250 μ M (higher than CMC_{st}), for pH of either 6.0 or 8.0. The cryo-TEM size at diRL concentration of 25 μ M is larger for pH 6.0 than for pH 8.0, 217 218 and they are identical at diRL concentration of 250 μ M. These results show that 219 change of the cryo-TEM size is similar to that of DLS size in terms of trend, indicating good consistency. The cryo-TEM sizes obtained at the condition of 25 µM 220 221 diRL and pH 6.0 (24.9 nm) is significantly smaller than the DLS-based size (43.2 nm). 222 The particle size obtained by DLS method is hydrodynamic diameter, which is the

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223 diameter of a sphere that has the same translational diffusion coefficient as the particle. This hydrodynamic size is usually larger than the real particle size.²⁹ Either the DLS 224 225 size or the cryo-TEM size obtained at high diRL concentration (0.5mM) in our study 226 is smaller than that measured at similar concentrations using similar methods in the study of Guo and Hu,⁸ in which formation of large vesicles was observed. The ionic 227 strength of diRL solution in that study is approximately10 mM, which is significantly 228 229 lower than that in our study (55 mM with divalent ions). The hydrophilic head of diRL molecule contains a carboxylic group. At pH higher than 6.0, the majority of 230 231 carboxylic groups are dissociated and negatively charged. Cations in the diRL 232 electrolyte solution can easily bind with the carboxylate groups, resulting in the induction of the solvated groups and disfavours formation of large aggregates.⁹ Such a 233 conversion of large vesicles to small ones was also observed when $Cd^{2\scriptscriptstyle +}$ was 234 introduced in solution of rhamnolipid solution.⁴ In addition, the dirhamnolipid used in 235 the study of Guo and Hu contains higher ratio of long-chain species (Rha₂C₁₀C_{12:1} and 236 $Rha_2C_{10}C_{14:1}$. $Rha_2C_xC_{v(:z)}$ designates the diRL homologue with x and y as the carbon 237 atom number of each aliphatic acid chain in the lipid moiety, and z as the number of 238 unsaturated bonds in lipid moiety),⁸ which results in stronger hydrophobic interaction 239

240 between molecules and thus favours formation of large vesicles.

The diRL used in this study is not a pure compound comprising one species of molecules. Instead, it is a rhamnolipid mixture consisting of three homologues which are the same in structure of polar moiety (double rhamnose rings and a carboxylic group) while different in length of aliphatic chains ($Rha_2C_{10}C_{10}$, $Rha_2C_{10}C_{12:1}$ and

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245	$Rha_2C_{10}C_{12}$ with molar fractions of 0.70, 0.11 and 0.19, respectively). ²⁵ We speculate
246	that this multi-component nature of the diRL results in formation of aggregates at
247	concentrations below CMC_{st} . The strength of hydrophobic interactions between diRL
248	molecules with aliphatic chains of different lengths are not uniform, which may result
249	in a transitional state for aggregation-related behavior, e.g. formation of aggregates in
250	electrolyte solution before the solution surface is saturated with diRL (corresponding
251	to diRL concentration of $\mbox{CMC}_{\mbox{st}}$ and graduality in change of electrical conductivity
252	increasing rate. In the transitional state, increase in diRL solution concentration may
253	enhance partition of diRL molecules to aggregates and therefore increase the density
254	of the molecules in aggregate. Increase of solution pH results in enhanced dissociation
255	of diRL molecules. Both effects enhance the electrostatic repulsion between polar
256	moieties of diRL molecules in aggregates and hence the curvature of aggregates. As a
257	result, when diRL concentrations are lower than CMC_{st} (the transitional state) the
258	aggregate size decreases with increase of the concentration and solution pH.

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260 4 Conclusions

DLS and Cryo-TEM methods were used to study aggregation behavior of low-concentration diRL and the results demonstrated formation of aggregates at concentrations below CMC_{st} . The effect of diRL concentration and solution pH on aggregate size and zeta potential is significant when diRL concentration is lower than the CMC_{st} . The multicomponent nature of the diRL and consequently a transitional state are supposed to be responsible for these aggregation behaviors at low diRL

concentrations. Also, results of the study indicate that the surface-tension-based CMC may not be used as the concentration defining aggregate formation for multicomponent biosurfactants. This work is of importance for cost-effective application of rhamnolipid. Further researches should be focused on validating the transitional state speculation and characterizing the rhamnolipid aggregates in transitional state in more detail.

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274 Supplementary Information

Electrical conductivity versus diRL concentration profile, typical number PSD profiles generated by Malvern DTS Nano software, DLS diffusion coefficient versus diRL concentration profile, and the cryo-TEM images used for aggregate size distribution analysis are included in SI.

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Table 1 Gaussian regression parameters for DLS and cryo-TEM aggregate size distribution.

diRL Sample	DLS			cryo-TEM				
	μ^{a}	$\sigma^{2 b}$	\mathbb{R}^2	d^{c} (nm)	μ	σ^2	R^2	d (nm)
25µМ,рН 6.0	3.77	0.031	1.00	43.2	3.22	0.033	0.96	24.9
250µМ,рН 6.0	2.05	0.030	0.97	7.8	2.61	0.053	0.98	13.7
25µМ,рН 8.0	2.80	0.046	0.98	16.5	3.05	0.033	0.97	21.2
250µM,pH 8.0	2.06	0.027	0.98	7.8	2.61	0.044	1.00	13.6

^{*a*} mean of ln*d* obtained from Gaussian regression

^b variance of lnd obtained from Gaussian regression

^{*c*} the mean aggregate size obtained using $d = e^{\mu}$

Fig. 1 Surface tension versus diRL concentration in PBSS solution and determination of CMC_{st} .

Fig. 2 DLS size (a) and zeta potential (b) of aggregates as a function of diRL concentration and solution pH.

Fig. 3 Distribution of diRL aggregate size obtained using DLS and cryo-TEM methods and representative Cryo-TEM micrographs of diRL aggregates formed in PBSS solution. (a) 25 μ M, pH 6.0; (b) 250 μ M, pH 6.0; (c) 25 μ M, pH 8.0; (d) 250 μ M, pH 8.0.



Figure 1





Figure 2





Figure 3