Electronic Supplementary Information *for*

# **An insight into thermo-thickening behavior of wormlike micellar solutions based on ultra-long-chain surfactants**

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### **Characterization of five ultra-long-chain surfactants**

The purity of five ultra-long-chain surfactants was confirmed by high performance liquid chromatography (HPLC) (**Figures S1S5**).



Fig. S1. HPLC result of UC<sub>22</sub>DAI.



Fig. S2. HPLC result of UC<sub>22</sub>DAB.



Fig. S3. HPLC result of UC<sub>22</sub>DAS.



Fig. S4. HPLC result of UC<sub>18</sub>DAS.



Fig. S5. HPLC result of C<sub>18</sub>DAS.

## **Photographs of the solutions at characteristic temperature**

The five surfactant solutions were heated at different temperatures at least for 0.5 h. Then, we observed and photographed the appearance of five surfactant solutions at different temperatures. All the solution were optically clear and isotropic over the test temperature range (**Fig. S6**)



**Fig. S6**. Snapshots of the solutions of five ultra-long-chain surfactants at different temperatures.

### **The results and discussion of molecular dynamic simulation**



**Fig.** S7. Simulation snapshots of the equilibrated micellar structures of UC<sub>22</sub>DAB molecules at (A) 25 °C, (B) 40 °C, (C) 50 °C and (D) 70 °C. (E) The variation of asphericity ( $\alpha$ ), radius of gyration and the number of UC<sub>22</sub>DAB molecules inside micelle with temperature; (F) the calculated potential energy ( $\phi$ ) of  $UC_{22}DAB$  micelles at 50 °C.

To gain further understanding of the thermo-thickening behavior of  $UC_{22}DAB$  solution, MD simulations were performed at representative temperatures in Fig. 1A, 25, 40, 50, and 70 C, respectively, and the simulation snapshots of the equilibrated micellar structures are depicted in **Fig. S7**. The micelle in this work is artificially defined as a cluster with more than five  $UC_{22}DAB$  molecules. As shown in Fig. 7A, the  $UC_{22}DAB$  molecules self-assembles into two short rod-like micelles at 25 °C. At 40 and 50 °C, we find all the 30 UC<sub>22</sub>DAB molecules fuse into a large micelle (Figs. 7B and C). It just took 5 ns to create the large micelle at 50  $^{\circ}$ C, much earlier than that at 40 °C (17 ns), indicating that high temperature accelerates micellar aggregation. However, at 70 °C (Fig. 7D), the simulated 30 surfactants end up forming two micelles; furthermore, the combination of these two micelles does not happen even when we extend the simulation to 200 ns, which suggests that aggregation is a relatively unfavorable process at higher temperature.

We calculated the average radius of gyration  $(r<sub>g</sub>)$  of the UC<sub>22</sub>DAB micelles in the simulation at four different temperatures. It was determined that the micelles are well extended from 25 °C to 50 °C with  $r_g$  of around 0.91 nm, while a marked decrease of  $r_g$  (0.84 nm) is found at 70 °C, signifying the coil-like structure. The structural change of the surfactant corresponds well with the evolution of the micelles. Thus, "unbreakable" wormlike micelles were found below 50  $\degree$ C, as opposed to the short rod-like micelles and even spherical micelles formed by wormlike micelles fracture which were observed at  $70^{\circ}$ C. This smooth structure change from rod to spherical is better illustrated by a calculation of the micelle asphericity  $(a)$ , which is used to describe micellar morphology. The degree of asphericity ( $\alpha$ ) of micelles is defined in terms of the eigenvalue  $\lambda_i^2$ , which can be written as:

$$
\alpha = \frac{\langle (T_{\rm r}^2 - 3M) \rangle}{T_{\rm r}^2}
$$

(1)

with  $T_r = \lambda_1^2 + \lambda_2^2 + \lambda_3^2$ ,  $M = \lambda_1^2 \lambda_2^2 + \lambda_2^2 \lambda_3^2 + \lambda_1^2 \lambda_3^2$  and  $\langle \ldots \rangle$  stands for the ensemble average. For a perfect sphere  $\alpha$  equals 0, whereas  $\alpha = 1$  corresponds to the extreme of an infinitely thin rod. This result in Fig. S7E corresponds well to the estimate of the surfactant number counted in the biggest micelle, for which the larger aggregation number of surfactants observed from 25  $\degree$ C to 50  $\degree$ C results in a rod-like micelle with smaller asphericity. Note that, it is generally unfeasible to combine two spherical micelles, given the like-charged and hydrophilic oxygen atoms that cover the micelle surface. As a result, the micelles do not favour stepwise growth at  $70^{\circ}$ C.

The current results still cannot rationalize our simulations at 25  $\degree$ C, given the fact that the maximum micelle size (Fig. S7E) at this temperature is smaller, whereas the observed  $r_g$ and the asphericity factor at this temperature is indeed similar with those at 40 and 50  $^{\circ}$ C. We therefore examined the behaviour of the system potential energy ( $\phi$ ) measured at 50 °C (Fig. S7F). A sharp decrease of  $\phi$  with a magnitude of 400 kJ·mol<sup>-1</sup> occurs in the simulation which is mostly contributed from the energy gain in joining the hydrophobic interface of two micelles. Prior to this jump, we also find an energy barrier of  $150 \text{ kJ·mol}^{-1}$  before the equilibrated structure of the bigger micelle is reached. Possible reasons for this energy barrier are the electrostatic repulsions of the densely packed charged groups of the surfactant, and possibly more importantly, from a subtle effect of the self-adjustment of the micelle to extend its hydrophobic surface before joining with another micelle. Overcoming this energy barrier will be easier at high temperatures, whereas at lower temperatures (such as  $25 \text{ °C}$ ) the micelles have less flexibility and a high bending modulus. The simulation results indicate the presence of a range of temperatures between 40  $\degree$ C and 50  $\degree$ C where the underlying  $UC_{22}DAB$  have a better chance of forming larger micelles, and possibly of a denser wormlike micellar network.

#### **Additional Rheological results**



**Fig. S8.** Temperature dependence of apparent viscosity of UC<sub>22</sub>DAB aqueous solutions at different heating rate.



Fig. S9. Effect of concentration on thermo-thickening behavior of (A) UC<sub>22</sub>DAB and (C) UC<sub>22</sub>DAI at fixed shearing rate of 100 s<sup>-1</sup>; effect of shearing rate on thermo-thickening behavior of (B) UC<sub>22</sub>DAB and (D) UC<sub>22</sub>DAI at fixed concentration of 3 %.



Fig. S10. Temperature dependence on apparent viscosity of (A) series concentration  $UC_{22}DAS$  at 100 s<sup>-1</sup> and (B)  $3\%$  UC<sub>22</sub>DAS at various shear rates.



Fig. S11. Temperature dependence on apparent viscosity of (A) series concentration  $UC_{18}DAS$  at 100 s<sup>-1</sup> and (B) 40 %  $UC_{18}DAS$  at various shear rate.





**Fig.** S12. Rheo-SANS data of 3.0% UC<sub>22</sub>DAB at 100 s<sup>-1</sup>. For better visibility the  $I(q)$  vs *q* curves are scaled vertically by factors of 2 (45 °C), 4 (50 °C), 8 (60 °C) and 12 (70 °C). The solid lines are the best fits to cylinder model.



Fig. S13. SAXS data of 3% (A)UC<sub>22</sub>DAB and (B)UC<sub>22</sub>DAI at various temperatures. For better visibility, the  $I(q)$  vs q curves of high temperatures ( $> 25 \degree$ C) in A and B are scaled vertically by factors of 2, 4, 6, 8, 10, 20 and 30. The red solid lines are the best fits to worm model.



Fig. S14. SAXS data of 3% (A)UC<sub>22</sub>DAB, (B)UC<sub>22</sub>DAI and (C) UC<sub>22</sub>DAS solutions at various temperatures. For better visibility, the  $I(q)$  vs  $q$  curves of high temperatures (> 25 °C) in A, B and C are scaled vertically by factors of 2, 4, 6, 8, 10, 20 and 30. The red solid lines are the best fits to worm model.



Fig. S15. SAXS data of (A) 1% UC<sub>22</sub>DAS and (B) 20% UC<sub>18</sub>DAS solutions at various temperatures. For better visibility, the  $I(q)$  vs *q* curves of high temperatures (> 25 °C) in A and B are scaled vertically by factors of 2, 4, 6, 8, 10, 20 and 30. The red solid lines are the best fits to worm model.



Fig. S16. SAXS data of 10% (A) C<sub>18</sub>DAS and (B) UC<sub>18</sub>DAS solutions at various temperatures. For better visibility, the  $I(q)$  vs q curves of high temperatures (> 25 °C) in A and B are scaled vertically by factors of 2, 4, 6, 8, 10, 20 and 30. The red solid lines are the best fits to worm model.

using cylinder model								
$T (^{\circ}C)$	Radius (nm)	Distribution of Radius	Persistence Length (nm)	<b>Fit Error</b> (nm)	Volume Fraction	Reduced $\chi^2$		
30	2.5	0.146	84.9	2.1	0.073	2.4		
45	2.5	0.096	52.1	1.7	0.095	6.9		
50	2.4	0.080	55.7	2.0	0.111	13.2		
60	2.4	0.058	61.4	2.1	0.105	14.0		
70	2.4	0.085	47.4	1.5	0.125	12.7		

Table S1. Structural parameters obtained from Rheo-SANS data on UC<sub>22</sub>DAB solution by curve fitting

**Table S2.** Characteristic parameters for 10% C<sub>18</sub>DAS solution at different shear rate

Shear rate $(s^{-1})$	$\eta_{\rm ini}$ (mPa·s)	$\eta_{\text{max}}$ (mPa·s)	$\sqrt{\circ}$	$(^{\circ}C)$ $\mathbf{c}$	
	. . ხ	210	42.5		63.6
10	. . 5	443	42.5	70.9	22.7
100	6	231	42.6	71.1	13.1
300	6	195	42.6	72.1	10.9
500	$\overline{\phantom{0}}$ . .	141	42.5	72.1	ر.,

 $\eta_{\text{ini}}$ : the initial viscosity;

 $\eta_{\text{max}}$ : the maximum viscosity;

 $T_s$ : the temperature at which viscosity starts to increase;

*T*<sub>c</sub>: the temperature at which viscosity reaches maximum;

*k*: the temperature sensitive factor.



Fig. S17. Cryo-TEM images of the 10 % UC<sub>18</sub>DAS solution at various temperatures.

The SAXS data was fitted to "worm-chain" model by SASfit software. The scattering vector *q* is defined as:

$$
q = (4\pi/\lambda)\sin\theta \tag{1}
$$

in which  $\lambda$  is the wavelength of the X-rays, and 2 $\theta$  is the scattering angle. The scattering intensity *I*(*q*) is generally described by:

$$
I(q) = nP(q)S(q) \tag{2}
$$

where *n* is the total number of particles, and *P*(q) and *S*(q) are the form and structure factors.

The form factors for wormlike micelles are approximated by the form factor of the Kholodenko-worm where the scattering length density profile across the wormlike segments are described by those of a rodlike micelle. The corresponding function are given by:

$$
P_{\text{core}}\left(q, R_{\text{core}}l, L\right) = P_{\text{worm}}\left(q, l, L\right) P_{\text{cs}}(q, R_{\text{core}}l, R_g) \tag{3}
$$

The contribution of the wormlike conformation of the micelle  $P_{\text{worm}}(q,l,L)$  is described by the formula of Kholodenko for wormlike structures. The contribution of the cross-section  $P_{cs}$ is the same as for rodlike micelles and given by

$$
P_{\rm cs}(q, R_{\rm core}, d, R_{\rm g}) = \left[\frac{2J_I(qR_{\rm core})}{qR_{\rm core}}\right]^2
$$
\n(4)

$$
Si (x) = \int_{0}^{x} t^{-1} \sin t dt
$$
\n
$$
P_{\text{brush}}(q, R_{g}) = 2 \frac{\exp (-x) - I + x}{x^{2}} \text{ with } x = R_{g}^{2} q^{2}
$$
\n(6)

$$
S_{\text{brush - core}}(q, R_{\text{core}}, l, L, R_{\text{g}}, d) = \psi(q, R_{\text{g}}) \times \frac{2J_I(qR_{\text{core}})}{qR_{\text{core}}} J_0 \left[ q \left( r_{\text{core}} + dR_{\text{g}} \right) \right] P_{\text{worm}}(q, l, L) \tag{7}
$$

$$
S_{\text{brush - brush}}\left(q, R_{\text{core}}l, L, d, R_{g}\right) = \psi^{2}\left(qR_{g}\right)J_{0}^{2}\left[q\left(r_{\text{core}} + dR_{g}\right)\right]P_{\text{worm}}\left(q, l, L\right) \tag{8}
$$

The excess scattering lengths and aggregation number are given by:

$$
\beta_{\text{core}} = V_{\text{core}} \left( \eta_{\text{core}} - \eta_{\text{solv}} \right) \tag{9}
$$

$$
\beta_{\text{brush}} = V_{\text{brush}} \left( \eta_{\text{brush}} - \eta_{\text{solv}} \right) \tag{10}
$$

$$
N_{\text{agg}} = \left(I - x_{\text{solv,core}}\right) \pi R_{\text{core}}^2 L/V_{\text{core}} \tag{11}
$$

The parameters are:

*R*<sub>core</sub>: core radius

 $V_{\text{core}}$ : molecular volume of single block unit in the micellar core

*V*brush: molecular volume of single block unit in the micellar corona

 $\eta_{\text{core}}$ : scattering length density of spherical core

 $\eta_{\text{break}}$ : scattering length density of the block unit in the corona

 $\eta_{\text{solv}}$ : scattering length density of solvent

 $xsolv<sub>core</sub>: amount of solvent in core$ 

*R*g: gyration radius of polymer chains in the corona

*l*: Kuhn length of the wormlike of the micelle

*L*: contour length of the wormlike of the micelle

The description of logistic function in Fig. 5A:

$$
y = A_2 + \frac{A_1 - A_2}{1 + \left(\frac{x}{x_0}\right)^p}
$$
 (12)

It can be converted into:

$$
\frac{y - A_2}{A_1 - A_2} = \frac{1}{1 + \left(\frac{x}{A_0}\right)^p}
$$
(13)

where, the  $A_1$  is the minimum value of logistic function,  $A_2$  is the maximum value,  $x_0$  is the center and the *p* is the power index. The result of fitting with logistic function is shown in Fig. S18, here, the  $\eta$ , T,  $\eta_{\text{min}}$ ,  $\eta_{\text{max}}$ ,  $T_0$  are *y*, *x*,  $A_1$ ,  $A_2$ ,  $x_0$  respectively. Thus, function (19) can be written as:

$$
\frac{\eta - \eta_{\text{max}}}{\eta_{\text{min}} - \eta_{\text{max}}} = \frac{1}{1 + \left(\frac{T}{T_0}\right)^p}
$$
(14)

(15)

where, the  $\eta_{\text{min}}$  is the minimum viscosity,  $\eta_{\text{max}}$  is the maximum viscosity, respectively,  $T_0$  is the center temperature and the  $p$  is the power index. And the viscosity at  $T_0$  satisfies the following formula:



Fig. S18 Temperature dependence of apparent viscosity of  $10\%$  C<sub>18</sub>DAS. The solid line is fitted by logistic function.

Sample	$T (^{\circ}C)$	$R_{\rm core}(nm)$	$V_{\rm core}$	$V_{\text{brush}}$	$R_{\rm g}$ (nm)	$l$ (nm)	$L$ (nm)
3%	25	2.5	0.6	0.1331	1.4	52	$>100$
UC22DAB	33	2.5	0.6	0.121	1.6	52	$>100\,$
(without	38	2.5	$0.6\,$	0.1331	1.4	55	$>100\,$
NaCl)	47	2.5	0.6	0.11	1.6	58	$>100$
	53	2.5	0.6	0.1331	1.4	60	$>100$
	56	2.5	0.6	0.1331	1.4	60	$>100$
	65	2.5	0.6	0.1331	1.4	55	$>100\,$
	73	2.5	0.6	0.1331	1.4	51	$>100\,$
3%	25	2.9	0.6	0.132	1.3	55	$>100$
UC22DAI	33	2.9	$0.6\,$	0.132	1.3	57	$>100\,$
(without	38	2.9	0.6	0.132	1.3	59	$>100\,$
NaCl)	47	2.9	0.6	0.132	1.3	59	$>100\,$
	53	2.9	0.6	0.132	1.2	60	$>100\,$
	56	2.9	0.6	0.132	1.3	61	$>100\,$
	65	2.9	0.6	0.132	1.3	56	$>100\,$
	73	2.9	0.6	0.132	1.3	49	$>100\,$
3%	25	2.5	0.6	0.1461	1.44	46	$>100\,$
UC22DAB	33	2.5	0.6	0.1461	1.44	46	$>100$
	38	2.5	0.6	0.1461	1.44	51	$>100$
	47	2.5	0.6	0.1461	1.44	53	$>100$
	53	2.5	0.6	0.1461	1.44	56	$>100$
	56	2.5	0.6	0.1461	1.44	57	$>100\,$
	65	2.5	$0.6\,$	0.1461	1.44	50	$>100\,$
	73	2.5	0.6	0.1461	1.44	48	$>100\,$
$3\%$	25	2.9	0.6	0.16	0.73	63	$>100$
UC22DAI	33	2.9	0.6	0.16	0.73	60	$>100$
	38	2.9	0.6	0.16	0.73	59	$>100\,$
	47	2.9	0.6	0.16	0.73	55	$>100$
	53	2.9	0.6	0.16	0.73	54	$>100\,$
	56	2.9	0.6	0.16	0.73	54	$>100\,$
	65	2.9	0.6	0.16	0.73	52	$>100\,$
	73	2.9	0.6	0.16	0.73	50	$>100\,$
3%	25	2.8	0.6	0.1947	1.3	50	$>100$
UC22DAS	33	2.8	0.6	0.1947	1.3	51	$>100$
	38	2.8	0.6	0.1947	1.3	52	>100
	47	2.8	0.6	0.1947	1.3	51	$>100$
	53	2.8	0.6	0.1947	1.3	50	$>100$
	56	2.8	0.6	0.1947	1.3	51	$>100$
	65	2.8	0.6	0.1947	1.3	48	$>100$
	73	2.8	0.6	0.1947	1.3	45	$>100$
$1\%$	25	2.8	0.6	0.2848	1.19	37	$>100$
UC22DAS	33	2.8	0.6	0.2848	1.19	40	$>100$
	38	2.8	0.6	0.2848	1.19	42	>100
	47	2.8	0.6	0.2848	1.19	43	$>100$
	53	2.8	0.6	0.2848	1.19	43	$>100$
	56	2.8	0.6	0.2848	1.19	44	$>100$
	65	2.8	0.6	0.2848	1.19	42	$>100$
	73	2.8	0.6	0.2848	1.19	38	$>100$
20%	25	2.3	0.6	0.1089	0.1	1.8	4.5
UC18DAS	33	2.3	0.6	0.1089	0.1	1.7	$\overline{4}$
	38	2.3	0.6	0.1089	0.1	1.7	4.1
	47	2.3	0.6	0.1089	0.1	1.7	3.8

**Table S3.** Fitting parameters for the size distribution and formfactor of "worm-chain" model

