

## Supporting Information

### Molecular Packing of Surface Active Ionic Liquids in a Deep Eutectic Solvent: A Small Angle X-ray Scattering (SAXS) Study

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## **<sup>1</sup>H NMR results (600MHz, DMSO)**

C<sub>12</sub>mimBr 0.86-0.90 (3H), 1.21-1.38 (18H), 2.09-2.32 (2H), 4.12-4.16 (3H), 4.30-4.35 (2H), 7.32-7.40 (1H), 7.45-7.55 (1H), 10.32-10.43 (1H).

C<sub>14</sub>mimBr 0.86-0.90 (3H), 1.21-1.38 (22H), 2.00-2.10 (2H), 4.12-4.16 (3H), 4.30-4.35 (2H), 7.31-7.36 (1H), 7.43-7.51 (1H), 10.39-10.49 (1H).

C<sub>16</sub>mimBr 0.87-0.91 (3H), 1.20-1.340 (26H), 1.96-2.00 (2H), 4.12-4.16 (3H), 4.31-4.35 (2H), 7.27-7.30 (1H), 7.37-7.43 (1H), 10.49-10.57 (1H).

## **SAXS analysis**

### **Micelles**

The scattering vector  $q$  is defined as

$$q = (4\pi/\lambda)\sin\theta/2$$

in which  $\lambda$  is the wavelength of the X-ray and  $\theta$  is the scattering angle. The scattering intensity  $I(q)$  for monodisperse, homogeneous, and spherical particles is generally described by:

$$I(q) = nP(q)S(q)$$

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

### ***Generalized indirect Fourier transformation (GIFT) method***

For the GIFT method, it was done with the PCG software (version 4.05.12). The smeared data were used and desmearing process was included in the GIFT software. The pair distance distribution function (PDDF)  $p(r)$  could be calculated by the Fourier transformation of  $P(q)$ , which is defined as

$$P(q) = 4\pi \int_0^{\infty} p(r) \frac{\sin(qr)}{qr} dr$$

The size and shape of micelles could be obtained from PDDF curves.

For the electron density of the shell and the solvent is close, parameters obtained from PDDF curves correspond to the core of micelles, which is made of alkyl chains of surfactants. Considering the charge screening effects of ChG, a hard sphere  $S(q)$  with Percus-Yevick closure relation would be good. No polydispersity was taken into consideration.

### ***Model-fitting method***

For the model-fitting method, it was done with the SASfit software (version 0.94.7). The data were desmeared with the beam length by the SAXSquant software

(version 4.1.1.8319). Several models like sphere, ellipsoid and cylinder have been tried and the ellipsoid model give good fitting results. The hard sphere structure factor with Percus-Yevick closure relation was adopted for the interaction between micelles. A constant was added in the background.

### Calculation of the structural parameters of LLC

#### The H<sub>1</sub> phase

The lattice parameter (D) of the normal hexagonal liquid crystalline phase is obtained according to the equation.

$$D = \frac{4\pi}{\sqrt{3}q_1}$$

With the alkyl chains arranged inside the cylinders, i.e. the H<sub>1</sub> phase, the structural parameters are given by following equations.

$$R = D \sqrt{\frac{\sqrt{3}}{2\pi(1 + \frac{\rho_a(1-\omega)}{\rho_s \omega})}}$$

$$S = \frac{2M_a}{\rho_a RN}$$

where  $\rho_a$  and  $\rho_s$  are the density of C<sub>n</sub>mimBr and solvent ChG ( $\rho_a = 1.05, 1.04, 1.03$  g/cm<sup>3</sup> for C<sub>12</sub>mimBr, C<sub>14</sub>mimBr, C<sub>16</sub>mimBr, respectively;  $\rho_s = 1.18$  g/cm<sup>3</sup> for ChG), R is the radius of the normal cylinder-like aggregates, S is the corresponding area per molecule of surfactants at the hydrophilic/hydrophobic interface, M<sub>a</sub> is the molar mass of surfactants (331.33, 359.38, 387.44 g/mol for C<sub>12</sub>mimBr, C<sub>14</sub>mimBr, C<sub>16</sub>mimBr, respectively),  $\omega$  is the weight fraction of surfactant in the binary system, N is the Avogadro's number ( $6.022 \times 10^{23}$  mol<sup>-1</sup>).

#### The V<sub>1</sub> phase

The lattice parameter (D) of the normal bicontinuous cubic liquid crystalline phase is obtained according to the equation.

$$D = \frac{2\pi\sqrt{h^2 + l^2 + k^2}}{q}$$

The structural parameters are given by following equations,

$$1 - \Phi_a = 2\sigma\left(\frac{l}{D}\right) + \frac{4}{3}\pi\chi\left(\frac{l}{D}\right)^3$$

$$S = 2v_a \frac{\sigma D^2 + 2\pi\chi l^2}{\Phi_a D^3}$$

$$R = 0.248D - l \quad (\text{Ia3d})$$

where  $\Phi_a$  is the surfactant volume percentage,  $l$  is the solvent thickness,  $v_a$  is the volume of the surfactant,  $\chi$  is Euler-Poincaré characteristic (-8 for Ia3d),  $\sigma$  is ratio of the minimal surface in a unit cell to the quantity (unit cell volume)<sup>2/3</sup> (3.091 for Ia3d),  $R$  is the radius of the solvophobic domains.

### **The $L_\alpha$ phase**

The lattice parameter ( $D$ ) of the lamellar liquid crystalline phase is obtained according to the equation.

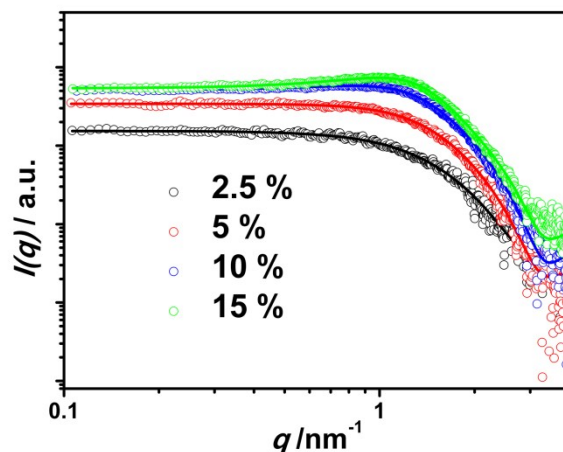
$$D = \frac{2\pi}{q_1}$$

$$d_s = D(1 - \Phi_a)$$

$$d_a = D - d_s$$

$$S = \frac{v_a}{d_a}$$

where  $d_s$  and  $d_a$  is thickness of the solvent and solvophobic layer .



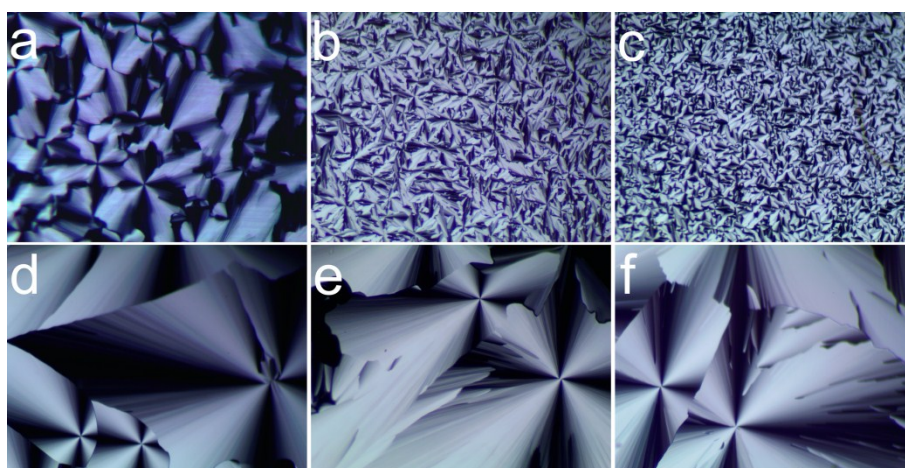
**Fig. S1** SAXS results (model-fitting method) of the C<sub>12</sub>mimBr/ChG system at 30 °C and different concentrations.

Open symbols for experimental and lines for fitting curves.

**Table S1** Parameters of micelles in the C<sub>12</sub>mimBr/ChG systems at 30 °C and different concentrations.

Concentration / %	a / nm	$\varepsilon$	R <sub>HS</sub> / nm	$\varphi$
2.5	1.28	1.66	2.49	0.0175
5	1.27	1.73	2.12	0.0436
10	1.26	1.82	2.17	0.0910
15	1.25	1.92	2.15	0.129

a, the equal semi-axis;  $\varepsilon$ , the axis ratio,  $\varepsilon a$ , the principle semi-axis; R<sub>HS</sub>, hard sphere repulsion radius;  $\varphi$ , volume fraction.

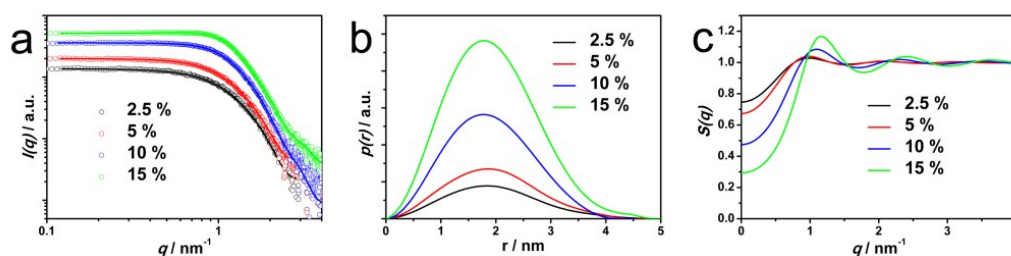


**Fig. S2** POM images of C<sub>12</sub>mimBr/ChG samples at 30 °C and different concentrations.

a) 60 %; b) 65%; c) 70 %; d) 75 %; e) 80 %; f) 85%.

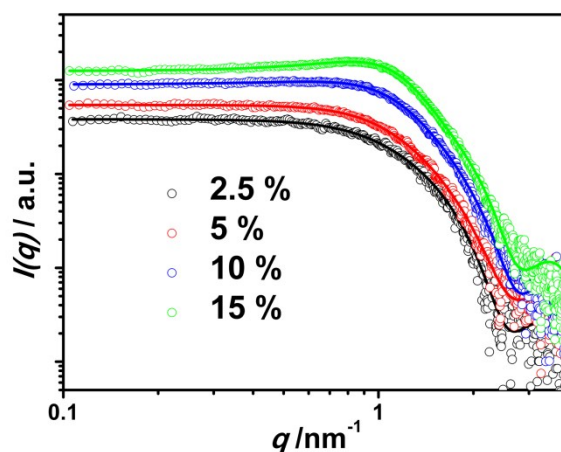
**Table S2** Structure parameters of H<sub>1</sub> phases in the C<sub>12</sub>mimBr/ChG system at 30 °C and different concentrations.

Concentrations / %	D / nm	R / nm	d / nm	S / nm <sup>2</sup>
60	4.37	1.82	0.73	0.577
65	4.22	1.82	0.58	0.575
70	4.15	1.85	0.45	0.565
75	4.08	1.88	0.32	0.557
80	3.97	1.88	0.21	0.556
85	3.86	1.88	0.10	0.557



**Fig. S3** SAXS results (GIFT method) of the C<sub>14</sub>mimBr/ChG system at 30 °C and different concentrations.

a) SAXS curves, open symbols for experimental and lines for fitting curves; b) PDDF curves; c)  $S(q)$  curves.

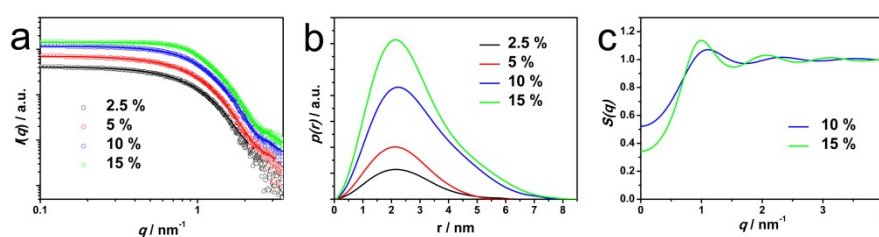


**Fig. S4** SAXS results (model-fitting method) of the C<sub>14</sub>mimBr/ChG system at 30 °C and different concentrations.

Open symbols for experimental and lines for fitting curves.

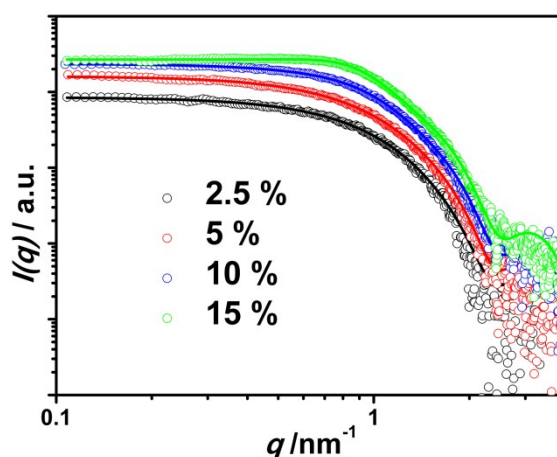
**Table S3** Parameters of micelles in the C<sub>14</sub>mimBr/ChG systems at 30 °C and different concentrations.

Concentration / %	a / nm	$\epsilon$	R <sub>HS</sub> / nm	$\phi$
2.5	1.59	1.56	3.26	0.0224
5	1.57	1.79	2.86	0.0400
10	1.53	1.87	2.62	0.0791
15	1.51	2.32	2.55	0.144



**Fig. S5** SAXS results (GIFT method) of the C<sub>16</sub>mimBr/ChG system at 30 °C and different concentrations.

a) SAXS curves, open symbols for experimental and lines for fitting curves; b) PDDF curves; c)  $S(q)$  curves.



**Fig. S6** SAXS results (model-fitting method) of the C<sub>16</sub>mimBr/ChG system at 30 °C and different concentrations.

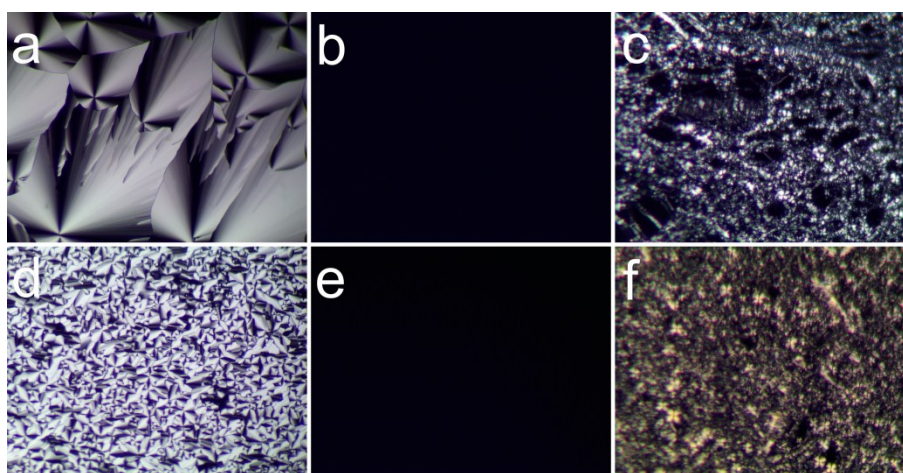
Open symbols for experimental and lines for fitting curves.

**Table S4** Parameters of micelles in the C<sub>16</sub>mimBr/ChG systems at 30 °C and different concentrations.

Concentration / %	a / nm	$\epsilon$	R <sub>HS</sub> / nm	$\phi$
2.5	1.76	1.93	-	-
5	1.73	2.07	-	-
10	1.70	3.24	3.02	0.0799
15	1.68	3.49	3.27	0.135

**Table S5** Parameters of micelles in the C<sub>n</sub>mimBr/ChG systems at a 5% concentration and 30 °C.

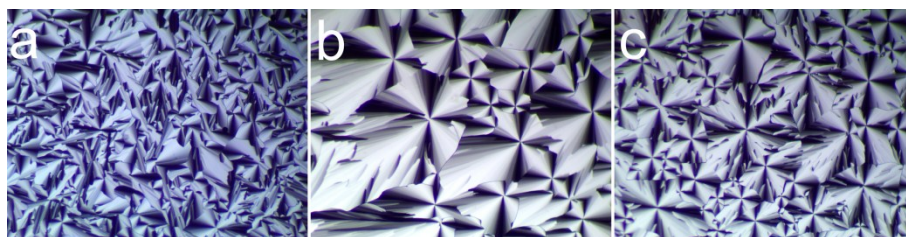
SAIL	a / nm	$\epsilon$
C <sub>12</sub> mimBr	1.27	1.73
C <sub>14</sub> mimBr	1.57	1.79
C <sub>16</sub> mimBr	1.73	2.07



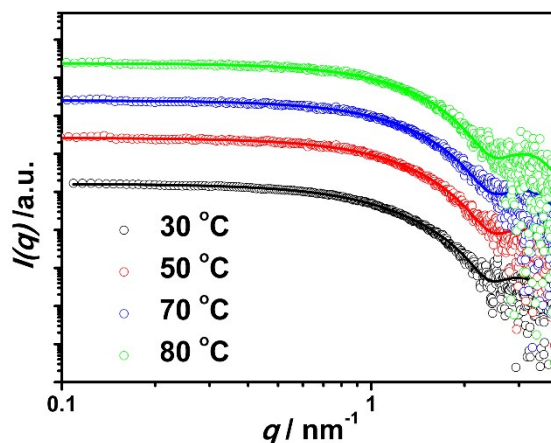
**Fig. S7** Representative POM images of different samples.

- a) C<sub>14</sub>mimBr/ChG, 65 %, 30 °C; b) C<sub>14</sub>mimBr/ChG, 80 %, 50 °C;  
 c) C<sub>14</sub>mimBr/ChG, 90 %, 60 °C; d) C<sub>16</sub>mimBr/ChG, 50 %, 50 °C;  
 e) C<sub>16</sub>mimBr/ChG, 75 %, 50 °C; f) C<sub>16</sub>mimBr/ChG, 90 %, 60 °C.





**Fig. S8** POM images of 65 %  $C_n$ mimBr/ChG samples at 50 °C.  
a)  $C_{12}$ mimBr; b)  $C_{14}$ mimBr; c)  $C_{16}$ mimBr.



**Fig. S9** SAXS results (model-fitting method) of the 5 %  $C_{16}$ mimBr/ChG sample at different temperatures.

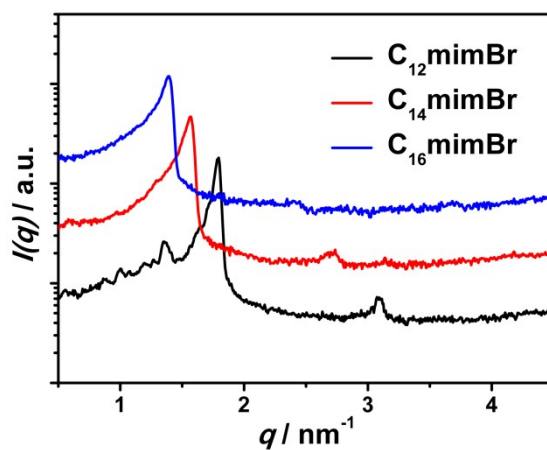
Open symbols for experimental and lines for fitting curves.

**Table S6** Micellar Parameters of the 5 %  $C_{16}$ mimBr/ChG sample at different temperatures.

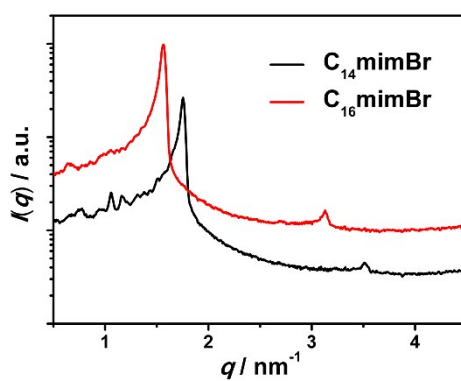
Temperature / °C	a / nm	$\epsilon$
30	1.73	2.07
50	1.71	1.78
70	1.70	1.71
80	1.70	1.67

**Table S7** Phase sequence of SAILs in different solvents.

Solvent	$C_{12}$ mimBr	$C_{14}$ mimBr	$C_{16}$ mimBr
ChG	$L_1/H_1/V_1$	$L_1/H_1/V_1/L_\alpha$	$L_1/H_1/V_1/L_\alpha$
water	$L_1/H_1/V_1/L_\alpha$	$L_1/H_1/V_1/L_\alpha$	$L_1/H_1/V_1/L_\alpha$
EAN	$L_1/H_1/V_1$	$L_1/H_1/V_1/L_\alpha$	$L_1/H_1/V_1/L_\alpha$



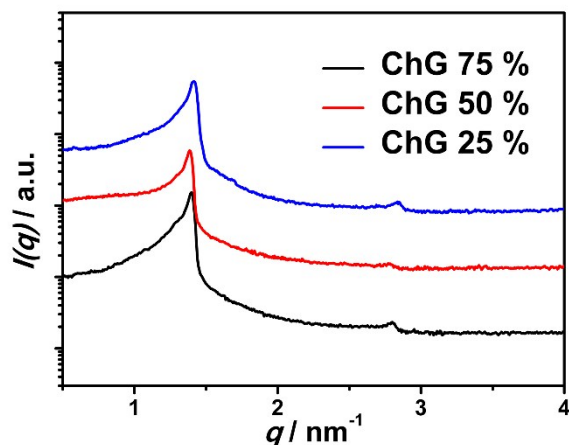
**Fig. S10** SAXS results of 65 %  $C_n$ mimBr/water samples at 50 °C.



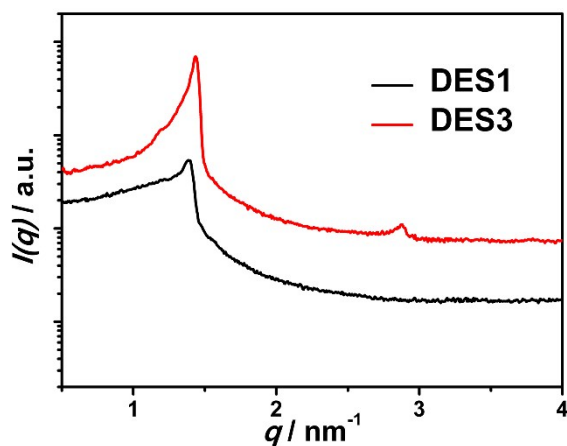
**Fig. S11** SAXS results of 65 %  $C_n$ mimBr/EAN samples at 50 °C.

**Table S8** Structure parameters of  $H_1$  phases at 65 % and 50 °C.

Surfactant	Solvent	D / nm	R / nm	d / nm	S / nm <sup>2</sup>
$C_{12}$ mimBr	water	4.03	1.69	0.65	0.620
$C_{14}$ mimBr	water	4.60	1.93	0.74	0.595
$C_{16}$ mimBr	water	5.16	2.17	0.82	0.576
$C_{14}$ mimBr	EAN	4.12	1.79	0.54	0.640
$C_{16}$ mimBr	EAN	4.62	2.01	0.60	0.621



**Fig. S12** SAXS results of 65 %  $C_{16}mimBr$ /solvent samples at 50 °C. ChG and water mixed solvents with different ChG weight percentage.



**Fig. S13** SAXS results of 65 %  $C_{16}mimBr$ /DES samples at 50 °C. DES1, ChCl/G molar ratio 1/1; DES3, ChCl/G molar ratio 1/3.

**Table S9** Structure parameters of 65 %  $C_{16}mimBr$ /solvent  $H_1$  phases at 50 °C.

Solvent	D / nm	R / nm	d / nm	S / nm <sup>2</sup>
ChG 75 %	5.16	2.22	0.72	0.563
ChG 50 %	5.19	2.21	0.77	0.565
ChG 25 %	5.15	2.18	0.79	0.569
DES1	5.17	2.23	0.71	0.561
DES3	5.01	2.17	0.67	0.575