

Supporting Information  
to the paper “Character of localization and microenvironment of the solvatochromic  
Reichardt’s betaine dye in SDS and CTAB micelles: MD simulation study”

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Table S1. Average numbers of surfactant C and H within 0.4 nm of different parts of the dye molecule.

system	Py <i>p</i> -Ph	Py <i>o</i> -Ph	Ph <i>o</i> -Ph	$\Delta$ Py <i>o</i> -Ph	$\Delta$ Ph <i>o</i> -Ph	O
$^+D^-$ in SDS	$26.3 \pm 1.2$	$21.3 \pm 1.0$	$14.9 \pm 0.4$	$17.1 \pm 3$	$15.3 \pm 0.9$	$2.4 \pm 0.05$
$^+D^-$ in CTAB	$26.4 \pm 0.8$	$23.2 \pm 0.5$	$17.9 \pm 0.6$	$15.1 \pm 1.2$	$21.5 \pm 1.2$	$3.9 \pm 0.10$
$^+DH$ in SDS	$16.4 \pm 0.01$	$21.2 \pm 0.2$	$22.2 \pm 0.5$	$15.8 \pm 1.0$	$16.1 \pm 1.0$	$8.3 \pm 0.3$
$^+DH$ in CTAB	$18.5 \pm 1.1$	$19.8 \pm 0.5$	$20.7 \pm 1.0$	$14.8 \pm 1.5$	$14.8 \pm 2$	$7.1 \pm 0.8$

Table S2. Maximum numbers of surfactant C and H atoms found within 0.4 nm of different parts of the dye molecule (averaged over 10 ps intervals).

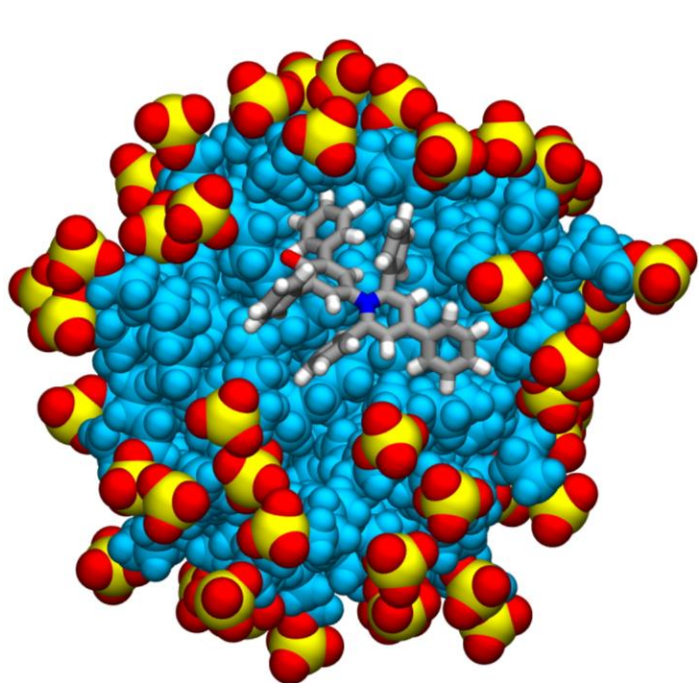
system	Py <i>p</i> -Ph	Py <i>o</i> -Ph	Ph <i>o</i> -Ph	O
$^+D^-$ in SDS	49.2	46.7	44.6	10.9
$^+D^-$ in CTAB	48.4	49.7	46.1	12.9
$^+DH$ in SDS	44.7	47.9	48.8	17.7
$^+DH$ in CTAB	48.2	47.3	49.0	17.1

Table S3. Average numbers of surfactant C and H atoms within 0.4 nm of different parts of  $^+D^-$  in SDS solution for runs starting from the different initial configurations.

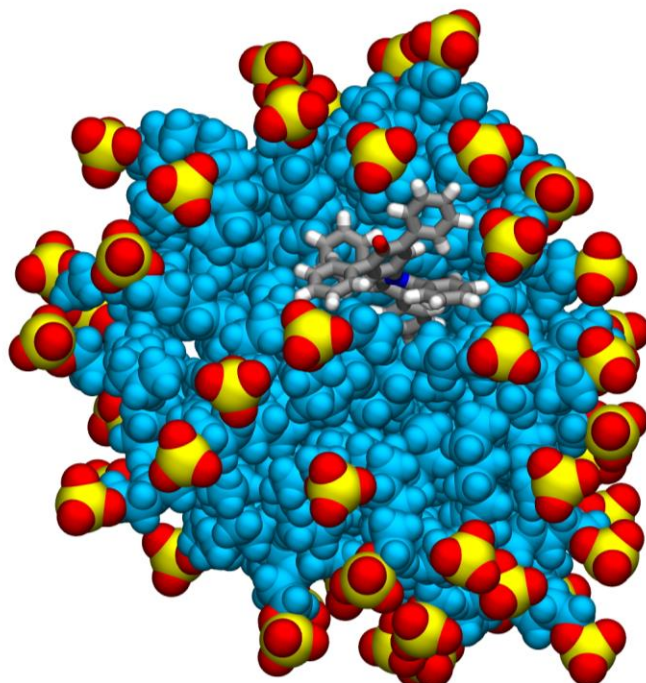
initial configurations	Py <i>p</i> -Ph	Py <i>o</i> -Ph	Ph <i>o</i> -Ph	$\Delta$ Py <i>o</i> -Ph	$\Delta$ Ph <i>o</i> -Ph	O
dye in bulk water	$25.7 \pm 1.6$	$21.4 \pm 0.4$	$15.1 \pm 0.5$	$17.3 \pm 0.9$	$15.9 \pm 0.3$	$2.5 \pm 0.09$
dye in micelle core	$26.3 \pm 1.2$	$21.3 \pm 1.0$	$14.9 \pm 0.4$	$17.1 \pm 3$	$15.3 \pm 0.9$	$2.4 \pm 0.05$

Table S4. Average numbers of various atoms in microenvironment of  $^+D^-$  and its O atom on SDS micelles for runs starting from the different initial configurations.

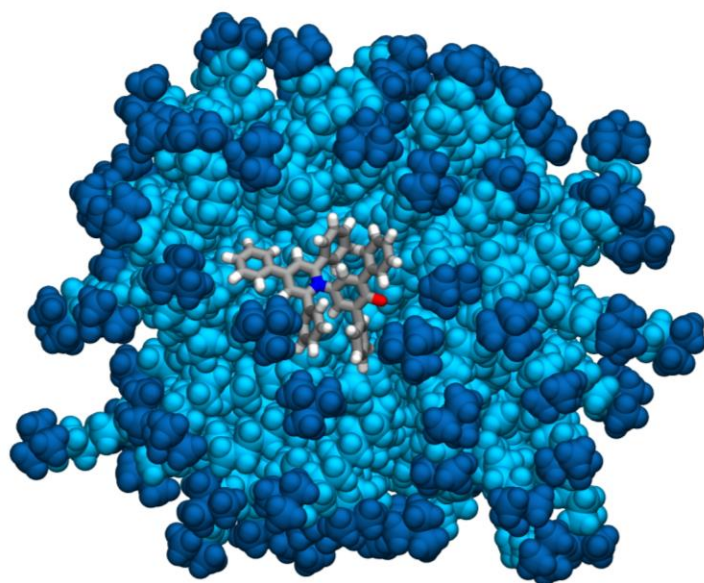
initial configurations	micelle core	water	headgroups	counterions	total
the dye molecule					
dye in bulk water	$119.1 \pm 1.6$	$89.5 \pm 0.4$	$8.6 \pm 0.3$	$0.37 \pm 0.02$	$217.6 \pm 2.3$
dye in micelle core	$120 \pm 5$	$89 \pm 5$	$8.4 \pm 0.7$	$0.37 \pm 0.03$	$218 \pm 11$
the O atom					
dye in bulk water	$2.5 \pm 0.18$	$10.3 \pm 0.17$	$0.028 \pm 0.006$	$0.082 \pm 0.006$	$12.8 \pm 0.4$
dye in micelle core	$2.6 \pm 0.17$	$10.29 \pm 0.04$	$0.032 \pm 0.007$	$0.10 \pm 0.02$	$13.0 \pm 0.3$



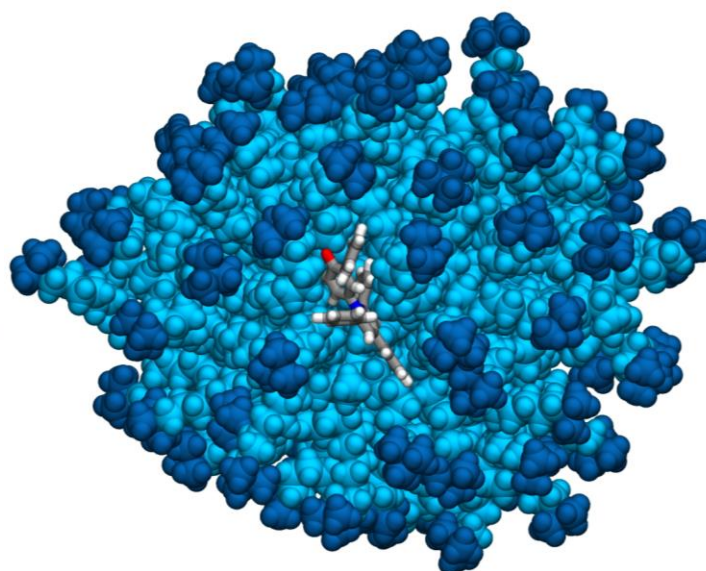
A



B



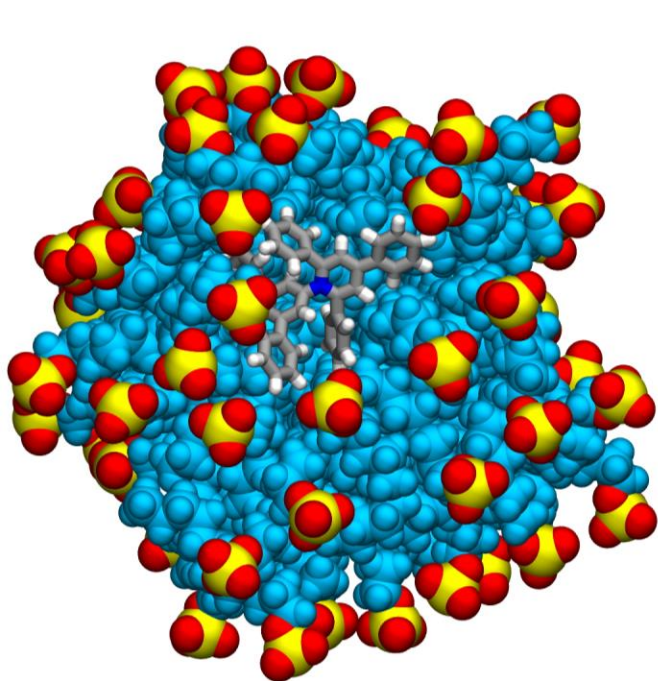
C



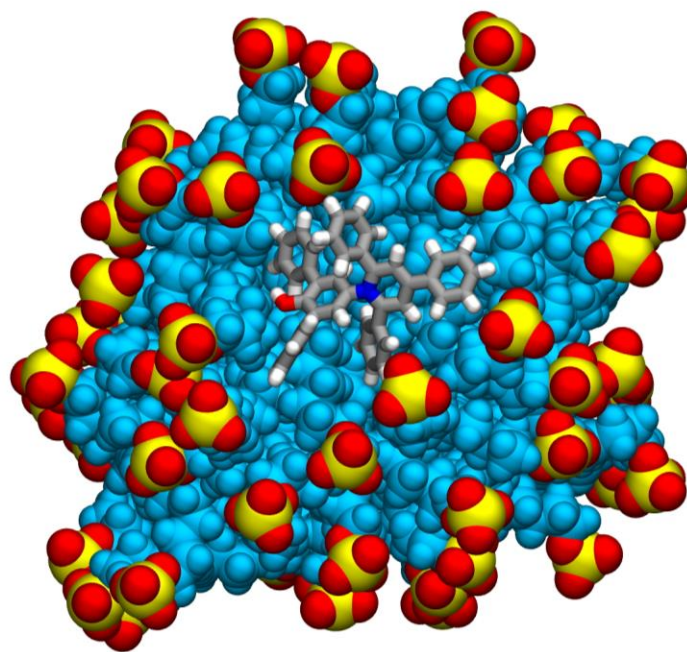
D

Figure S1. Snapshots from MD trajectories showing typical localizations and orientations of the neutral form of RD on SDS (A, B) and CTAB (C, D) micelles.

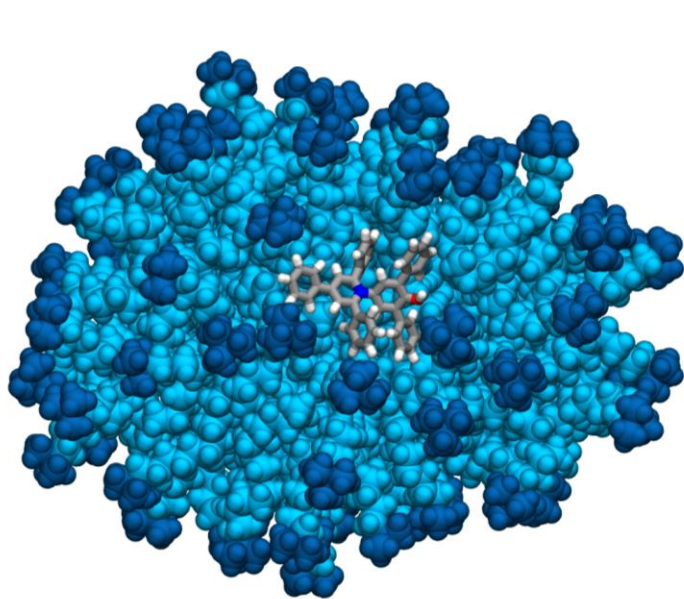




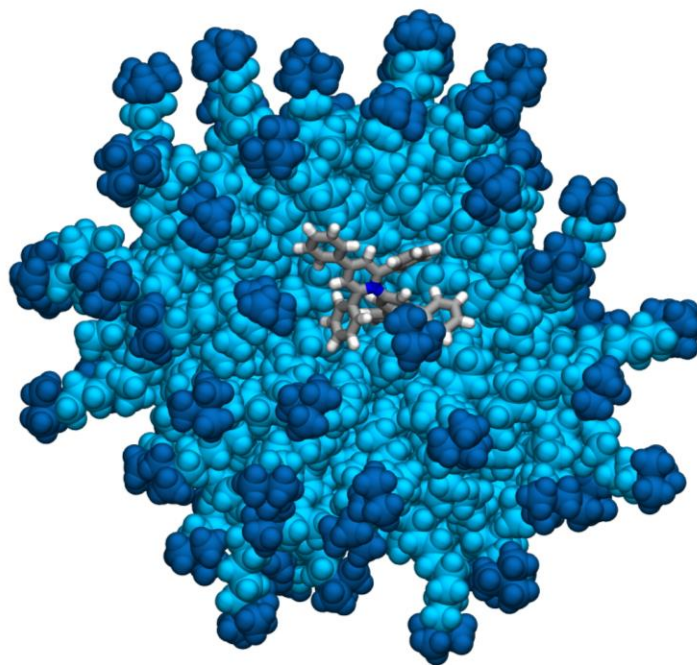
A



B



C



D

Figure S2. Snapshots from MD trajectories showing typical localizations and orientations of the protonated form of RD on SDS (A, B) and CTAB (C, D) micelles.

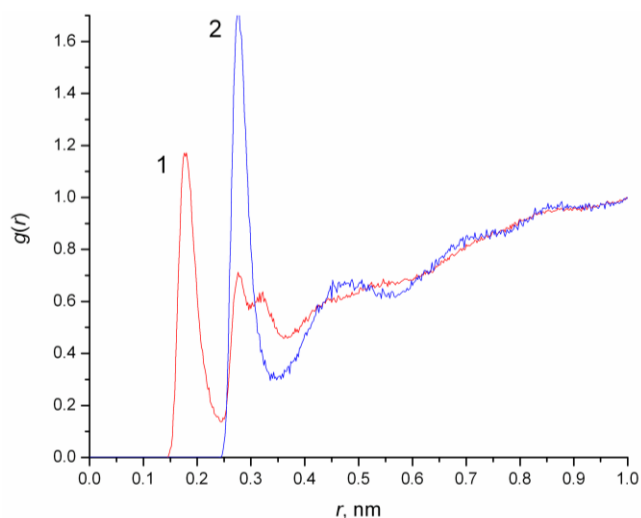


Figure S3. Radial distribution functions between the  $^+D^-$  O atom and either both water O and H atoms (red curve, 1) or water O atoms only (blue curve, 2) in pure water solution.

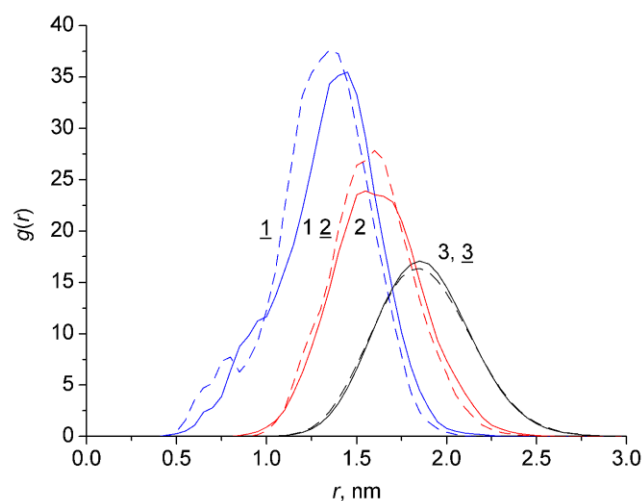


Figure S4. Radial distribution functions micelle COM — RD N (blue curves, 1) and micelle COM — RD O (red curves, 2) in comparison with RDFs micelle COM — surfactant S (N) (black curves, 3). Solid lines are for the runs starting from the initial configurations with the solubilized dye, dashed lines with underlined numbers are for the runs starting from the initial configurations with the dye placed in bulk water.

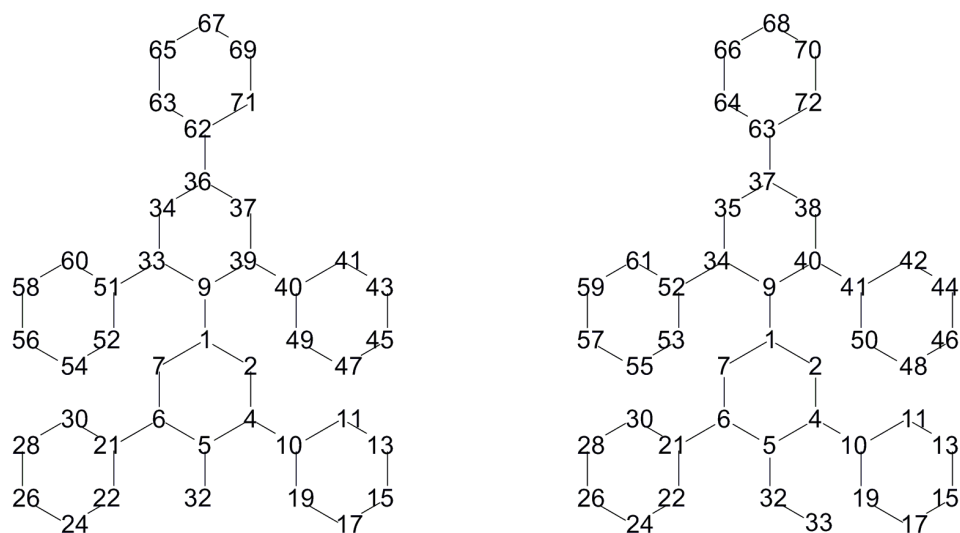


Figure S5. Scheme of numeration of atoms in potential models of the neutral and protonated forms of the RD. Hydrogen atoms in rings are not shown, they always follow the carbon atoms they are bonded to.

\*\*\*\*\* File C-C\_long\_dihs.itp \*\*\*\*\*

; Krzysztof Murzyn, Maciej Bratek, and Marta Pasenkiewicz-Gierula  
; Refined OPLS All-Atom Force Field Parameters for n-Pentadecane,  
; Methyl Acetate, and Dimethyl Phosphate  
; J. Phys. Chem. B 2013, 117, 16388-16396

#define HC_CT_CT_HC_long	3	0.3138	0.9414	0.0000	-1.2552	0.0000	0.0000
#define CT_CT_CT_CT_long	3	2.209155	4.543825	0.77822	-7.5312	0.0000	0.0000

\*\*\*\*\* File Ph-Ph\_params.itp \*\*\*\*\*

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[ angletypes ]

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C!	NC	C!	1	117.000	585.760

; Markus K. Dahlgren, Patric Schyman, Julian Tirado-Rives, and William L. Jorgensen  
; Characterization of Biaryl Torsional Energetics and its Treatment in OPLS All-Atom Force Fields  
; J. Chem. Inf. Model., 2013, 53 (5), 1191-1199

[ dihedraltypes ]

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C!	CA	CA	CA	3	30.33400	0.00000	-30.33400	0.00000	0.00000	0.00000
X	C!	NC	X	3	30.33400	0.00000	-30.33400	0.00000	0.00000	0.00000
X	NC	C!	X	3	30.33400	0.00000	-30.33400	0.00000	0.00000	0.00000
X	C!	CA	X	3	30.33400	0.00000	-30.33400	0.00000	0.00000	0.00000
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C!	CA	OH	HO	3	7.03749	0.00000	-7.03749	0.00000	0.00000	0.00000

\*\*\*\*\* File CTA.itp \*\*\*\*\*

; V.S. Farafonov, A.V. Lebed. Kharkov Univ. Bull., Chem. Ser. 2016,  
; 27, 25-30. [http://chembull.univer.kharkov.ua/archiv/2016\\_2/03.pdf](http://chembull.univer.kharkov.ua/archiv/2016_2/03.pdf)

[ moleculetype ]

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CTA 3

[ atoms ]

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24	opls_140	1	CTA	H71	8	0.06	
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51	50	53	56	3
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47	50	53	54	3
47	50	53	55	3
52	50	53	54	HC_CT_CT_HC_long
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50	53	56	58	3
54	53	56	57	HC_CT_CT_HC_long
54	53	56	58	HC_CT_CT_HC_long
54	53	56	59	3
55	53	56	57	HC_CT_CT_HC_long
55	53	56	58	HC_CT_CT_HC_long
55	53	56	59	3
57	56	59	60	HC_CT_CT_HC_long
57	56	59	62	HC_CT_CT_HC_long
57	56	59	61	HC_CT_CT_HC_long
53	56	59	60	3
53	56	59	62	3
53	56	59	61	3
58	56	59	60	HC_CT_CT_HC_long
58	56	59	62	HC_CT_CT_HC_long
58	56	59	61	HC_CT_CT_HC_long
1	14	17	20	3
1	14	17	18	3
1	14	17	19	3
23	20	17	14	CT_CT_CT_CT_long
17	20	23	26	CT_CT_CT_CT_long
29	26	23	20	CT_CT_CT_CT_long
23	26	29	32	CT_CT_CT_CT_long
35	32	29	26	CT_CT_CT_CT_long
29	32	35	38	CT_CT_CT_CT_long
41	38	35	32	CT_CT_CT_CT_long
35	38	41	44	CT_CT_CT_CT_long
47	44	41	38	CT_CT_CT_CT_long
41	44	47	50	CT_CT_CT_CT_long
44	47	50	53	CT_CT_CT_CT_long
47	50	53	56	CT_CT_CT_CT_long
50	53	56	59	CT_CT_CT_CT_long



\*\*\*\*\* File RD.itp \*\*\*\*\*

[ moleculetype ]

RD 3

[ atoms ]

1	opls_916	1	RD	C1	1	0.0249
2	opls_145	1	RD	C2	2	-0.1793
3	opls_146	1	RD	H2	2	0.0461
4	opls_145B	1	RD	C3	3	-0.1221
5	opls_166	1	RD	C4	4	0.4152
6	opls_145B	1	RD	C5	5	-0.1221
7	opls_145	1	RD	C6	6	-0.1793
8	opls_146	1	RD	H6	6	0.0461
9	opls_520	1	RD	N	7	0.1992
10	opls_145B	1	RD	C7	8	0.1032
11	opls_145	1	RD	C8	9	-0.1197
12	opls_146	1	RD	H8	9	0.1044
13	opls_145	1	RD	C9	10	-0.1337
14	opls_146	1	RD	H9	10	0.0974
15	opls_145	1	RD	C10	11	-0.1122
16	opls_146	1	RD	H10	11	0.0937
17	opls_145	1	RD	C11	12	-0.1337
18	opls_146	1	RD	H11	12	0.0974
19	opls_145	1	RD	C12	13	-0.1197
20	opls_146	1	RD	H12	13	0.1044
21	opls_145B	1	RD	C13	14	0.1032
22	opls_145	1	RD	C14	15	-0.1197
23	opls_146	1	RD	H14	15	0.1044
24	opls_145	1	RD	C15	16	-0.1337
25	opls_146	1	RD	H15	16	0.0974
26	opls_145	1	RD	C16	17	-0.1122
27	opls_146	1	RD	H16	17	0.0937
28	opls_145	1	RD	C17	18	-0.1337
29	opls_146	1	RD	H17	18	0.0974
30	opls_145	1	RD	C18	19	-0.1197
31	opls_146	1	RD	H18	19	0.1044
32	opls_167	1	RD	O	20	-0.566
33	opls_145B	1	RD	C19	21	0.0877
34	opls_145	1	RD	C20	22	-0.1749
35	opls_146	1	RD	H20	22	0.0768
36	opls_145B	1	RD	C21	23	0.1031
37	opls_145	1	RD	C22	24	-0.1749
38	opls_146	1	RD	H21	24	0.0768
39	opls_145B	1	RD	C23	25	0.0877
40	opls_145B	1	RD	C24	26	0.1156
41	opls_145	1	RD	C25	27	-0.1208
42	opls_146	1	RD	H25	27	0.1065
43	opls_145	1	RD	C26	28	-0.1219
44	opls_146	1	RD	H26	28	0.1216
45	opls_145	1	RD	C27	29	-0.0776
46	opls_146	1	RD	H27	29	0.1165
47	opls_145	1	RD	C28	30	-0.1219

48	opls_146	1	RD	H28	30	0.1216
49	opls_145	1	RD	C29	31	-0.1208
50	opls_146	1	RD	H29	31	0.1065
51	opls_145B	1	RD	C30	32	0.1156
52	opls_145	1	RD	C31	33	-0.1208
53	opls_146	1	RD	H31	33	0.1065
54	opls_145	1	RD	C32	34	-0.1219
55	opls_146	1	RD	H32	34	0.1216
56	opls_145	1	RD	C33	35	-0.0776
57	opls_146	1	RD	H33	35	0.1165
58	opls_145	1	RD	C34	36	-0.1219
59	opls_146	1	RD	H34	36	0.1216
60	opls_145	1	RD	C35	37	-0.1208
61	opls_146	1	RD	H35	37	0.1065
62	opls_145B	1	RD	C36	38	0.0969
63	opls_145	1	RD	C37	39	-0.1311
64	opls_146	1	RD	H37	39	0.1291
65	opls_145	1	RD	C38	40	-0.1146
66	opls_146	1	RD	H38	40	0.1169
67	opls_145	1	RD	C39	41	-0.0679
68	opls_146	1	RD	H39	41	0.1118
69	opls_145	1	RD	C40	42	-0.1146
70	opls_146	1	RD	H40	42	0.1169
71	opls_145	1	RD	C41	43	-0.1311
72	opls_146	1	RD	H41	43	0.1291

[ bonds ]

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60	58	1
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[ angles ]

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1	7	8	1
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33	51	60	1

[ dihedrals ]

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\*\*\*\*\* File RDH.itp \*\*\*\*\*

[ moleculetype ]

RDH 3

[ atoms ]

1	opls_916	1	RD	C1	1	0.0289
2	opls_145	1	RD	C2	2	-0.1710
3	opls_146	1	RD	H2	2	0.0970
4	opls_145B	1	RD	C3	3	-0.0410
5	opls_166	1	RD	C4	4	0.2523
6	opls_145B	1	RD	C5	5	-0.0410
7	opls_145	1	RD	C6	6	-0.1710
8	opls_146	1	RD	H6	6	0.0970
9	opls_520	1	RD	N	7	0.1769
10	opls_145B	1	RD	C7	8	0.1010
11	opls_145	1	RD	C8	9	-0.1223
12	opls_146	1	RD	H8	9	0.1040
13	opls_145	1	RD	C9	10	-0.1159
14	opls_146	1	RD	H9	10	0.1231
15	opls_145	1	RD	C10	11	-0.0888
16	opls_146	1	RD	H10	11	0.1172
17	opls_145	1	RD	C11	12	-0.1159
18	opls_146	1	RD	H11	12	0.1231
19	opls_145	1	RD	C12	13	-0.1223
20	opls_146	1	RD	H12	13	0.1040
21	opls_145B	1	RD	C13	14	0.1010
22	opls_145	1	RD	C14	15	-0.1223
23	opls_146	1	RD	H14	15	0.1040
24	opls_145	1	RD	C15	16	-0.1159
25	opls_146	1	RD	H15	16	0.1231
26	opls_145	1	RD	C16	17	-0.0888
27	opls_146	1	RD	H16	17	0.1172
28	opls_145	1	RD	C17	18	-0.1159
29	opls_146	1	RD	H17	18	0.1231
30	opls_145	1	RD	C18	19	-0.1223
31	opls_146	1	RD	H18	19	0.1040
32	opls_167	1	RD	O	20	-0.5019
33	opls_168	1	RD	HO	20	0.3723
34	opls_145B	1	RD	C19	21	0.0899
35	opls_145	1	RD	C20	22	-0.1677
36	opls_146	1	RD	H20	22	0.0978
37	opls_145B	1	RD	C21	23	0.1300
38	opls_145	1	RD	C22	24	-0.1677
39	opls_146	1	RD	H21	24	0.0978
40	opls_145B	1	RD	C23	25	0.0899
41	opls_145B	1	RD	C24	26	0.1096
42	opls_145	1	RD	C25	27	-0.1264
43	opls_146	1	RD	H25	27	0.1126
44	opls_145	1	RD	C26	28	-0.1212
45	opls_146	1	RD	H26	28	0.1328
46	opls_145	1	RD	C27	29	-0.0637
47	opls_146	1	RD	H27	29	0.1255
48	opls_145	1	RD	C28	30	-0.1212



49	opls_146	1	RD	H28	30	0.1328
50	opls_145	1	RD	C29	31	-0.1264
51	opls_146	1	RD	H29	31	0.1126
52	opls_145B	1	RD	C30	32	0.1096
53	opls_145	1	RD	C31	33	-0.1264
54	opls_146	1	RD	H31	33	0.1126
55	opls_145	1	RD	C32	34	-0.1212
56	opls_146	1	RD	H32	34	0.1328
57	opls_145	1	RD	C33	35	-0.0637
58	opls_146	1	RD	H33	35	0.1255
59	opls_145	1	RD	C34	36	-0.1212
60	opls_146	1	RD	H34	36	0.1328
61	opls_145	1	RD	C35	37	-0.1264
62	opls_146	1	RD	H35	37	0.1126
63	opls_145B	1	RD	C36	38	0.0701
64	opls_145	1	RD	C37	39	-0.1256
65	opls_146	1	RD	H37	39	0.1315
66	opls_145	1	RD	C38	40	-0.1145
67	opls_146	1	RD	H38	40	0.1274
68	opls_145	1	RD	C39	41	-0.0466
69	opls_146	1	RD	H39	41	0.1220
70	opls_145	1	RD	C40	42	-0.1145
71	opls_146	1	RD	H40	42	0.1274
72	opls_145	1	RD	C41	43	-0.1256
73	opls_146	1	RD	H41	43	0.1315

[ bonds ]

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61	59	1
11	13	1

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38	37	35	34	3
38	37	35	36	3
63	37	35	34	3
63	37	35	36	3
4	6	5	32	4

improper\_O\_C\_X\_Y



\*\*\*\*\* File topol\_CTAB\_neu.top \*\*\*\*\*

```
#include "oplsaa.ff/forcefield.itp"
#include "C-C_params.itp"
#include "Ph-Ph_params.itp"
#include "CTA.itp"
#include "oplsaa.ff/ions.itp"
#include "oplsaa.ff/spc.itp"
#include "RD.itp"
```

```
[ system ]
; Name
RD in CTAB micelle
```

```
[ molecules ]
; Compound      #mols
CTA              80
BR               80
RD               1
SOL             28571
```

\*\*\*\*\* File topol\_SDS\_prot.top \*\*\*\*\*

```
#include "oplsaa.ff/forcefield.itp"
#include "C-C_params.itp"
#include "Ph-Ph_params.itp"
#include "DS.itp"
; DS.itp is provided in the Supporting information to
; V.S. Farafonov, A.V. Lebed. Jour. Chem. Theor. Comput. 2017, 13, 2742-2750
#include "oplsaa.ff/ions.itp"
#include "oplsaa.ff/spc.itp"
#include "RDH.itp"
```

```
[ system ]
; Name
RDH in SDS micelle
```

```
[ molecules ]
; Compound      #mols
DS              60
Na              60
RDH             1
CL              1
SOL            16123
```