

Supporting information: A coil-to-globule transition capable coarse-grained model for poly(N-isopropylacrylamide)

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This file contents the comparison of the radial distribution functions (RDFs) and probability density functions (PDFs) obtained from coarse-grained and atomistic simulations. Also, in the second section we are including the topology files needed to simulate a 35-mer with the Gromacs package. These files include definitions of the bonded and non-bonded parameters and must be linked to the corresponding *topology.top* file.

I. RADIAL DISTRIBUTION AND PROBABILITY DENSITY FUNCTIONS

Each figure compares the outcomes from the coarse-grained (dashed lines) and the atomistic (continuous lines) models. Blue lines correspond to $T = 288$ K and red lines to $T = 328$ K, each one being below and above $T_c = (302.1 \pm 1.6)$ K, respectively. The insets show in highlighted colors the reference bead types. For all cases, to make the RDFs and PDFs comparable, we are considering the center of mass of the AA groups in correspondence to the centers of the CG beads. All data from CG simulations are obtained through 10 μ s trajectories and discarding the first 8 μ s.

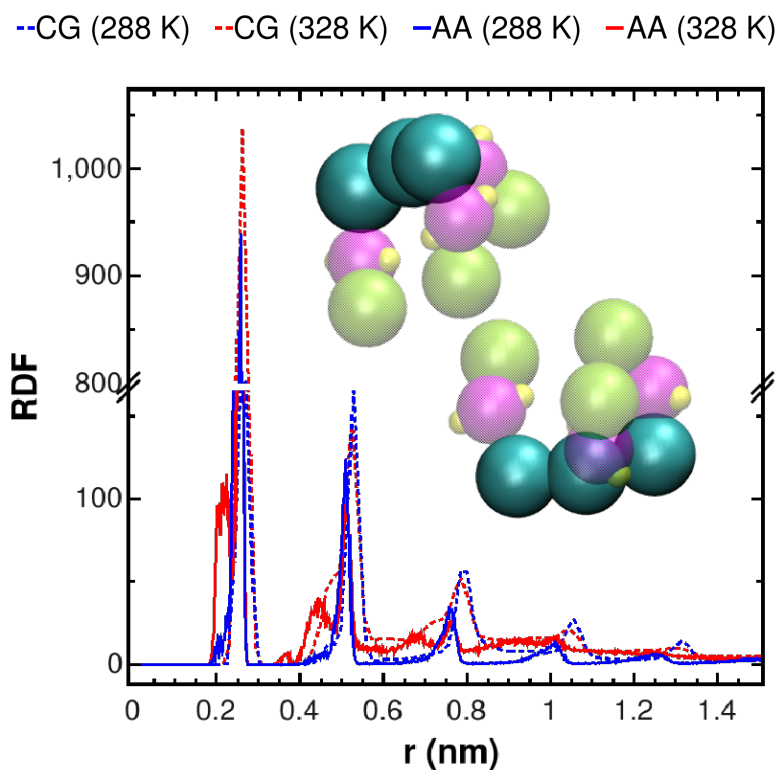


FIG. S1: Radial distribution function for the SC1-SC1 (backbone-backbone) pair.

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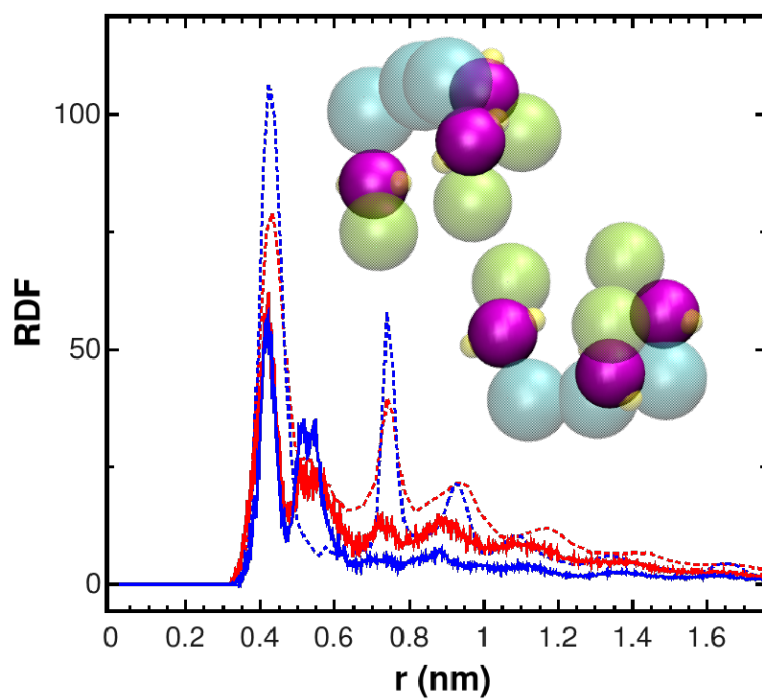


FIG. S2: Radial distribution function for the SNda-SNda (amide-amide) pair.

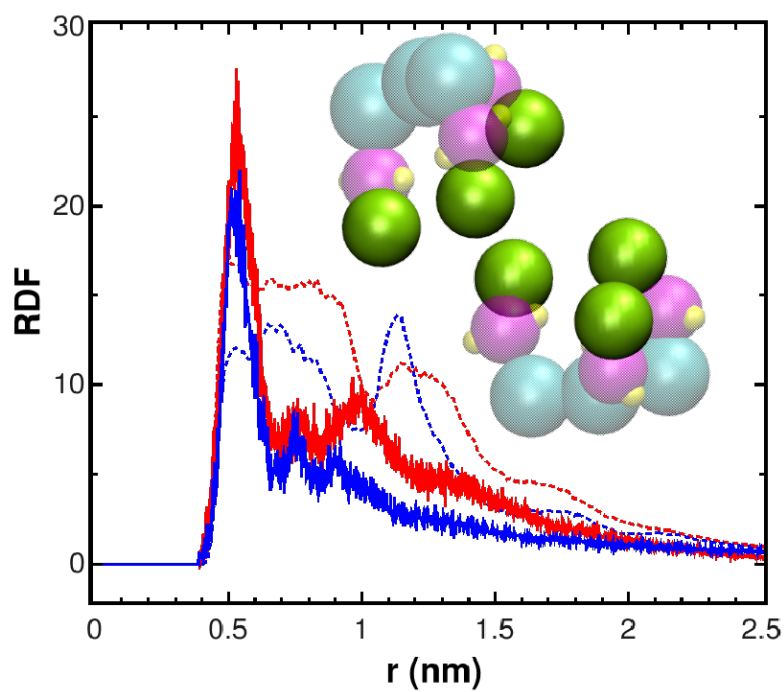


FIG. S3: Radial distribution function for the SC3-SC3 (isopropyl-isopropyl) pair.

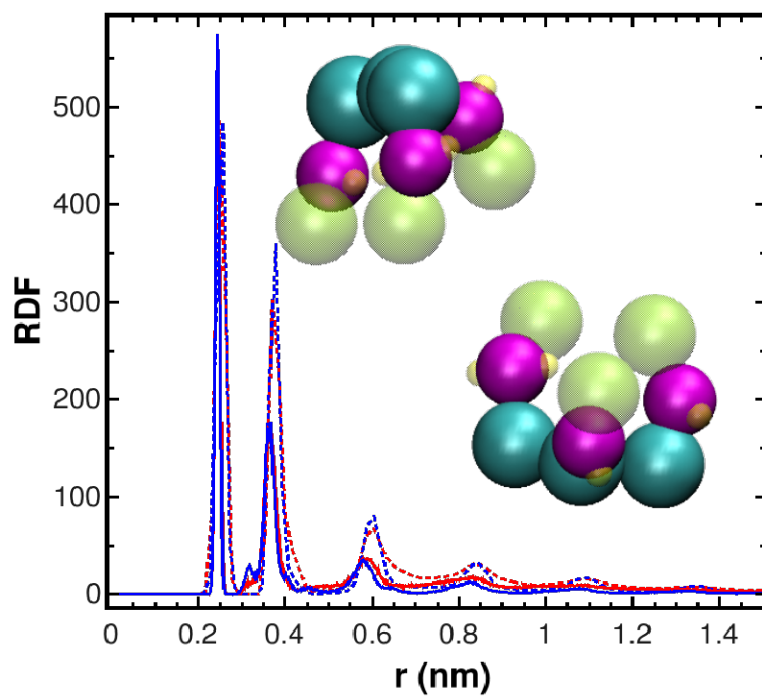


FIG. S4: Radial distribution function for the SC1-SNda (backbone-amide) pair.

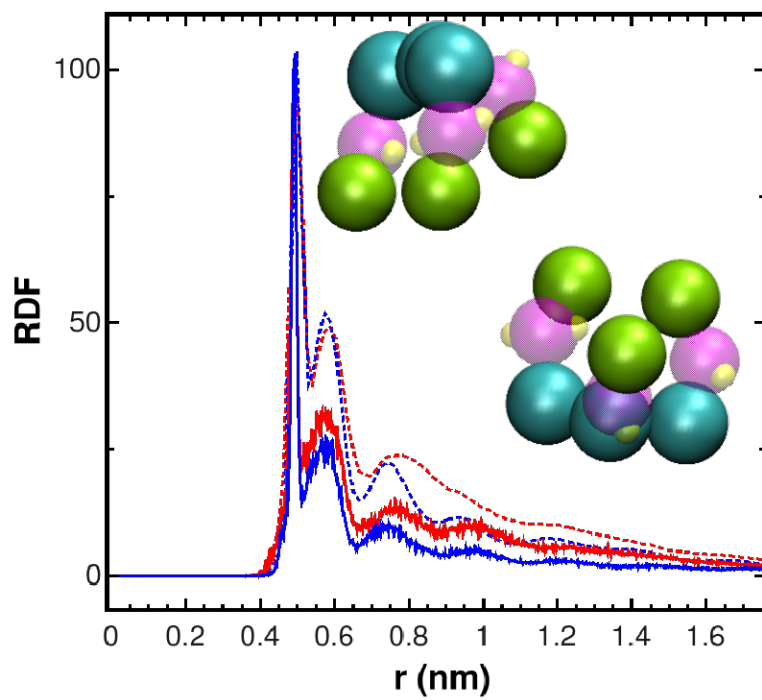


FIG. S5: Radial distribution function for the SC1-SC3 (backbone-isopropyl) pair.

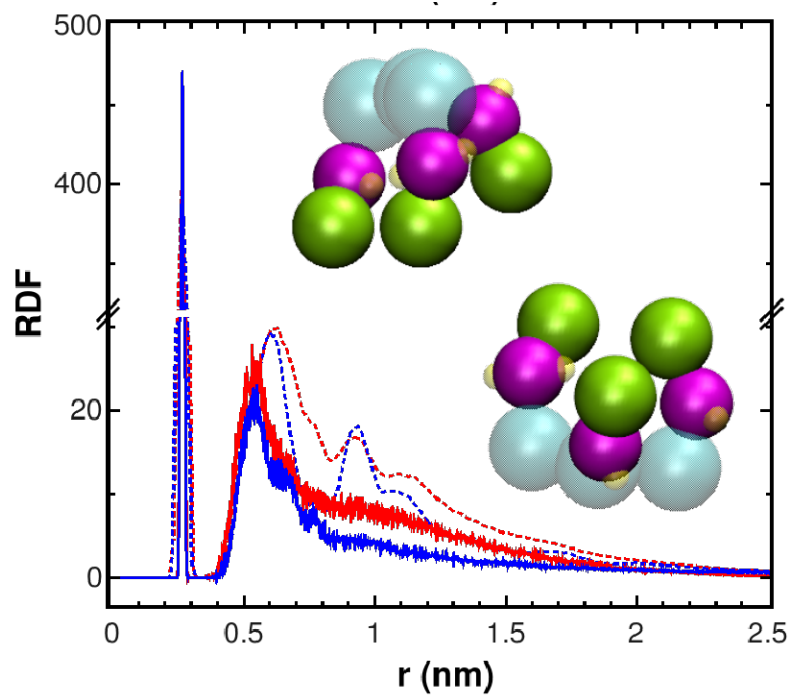


FIG. S6: Radial distribution function for the SNda-SC3 (amide-isopropyl) pair.

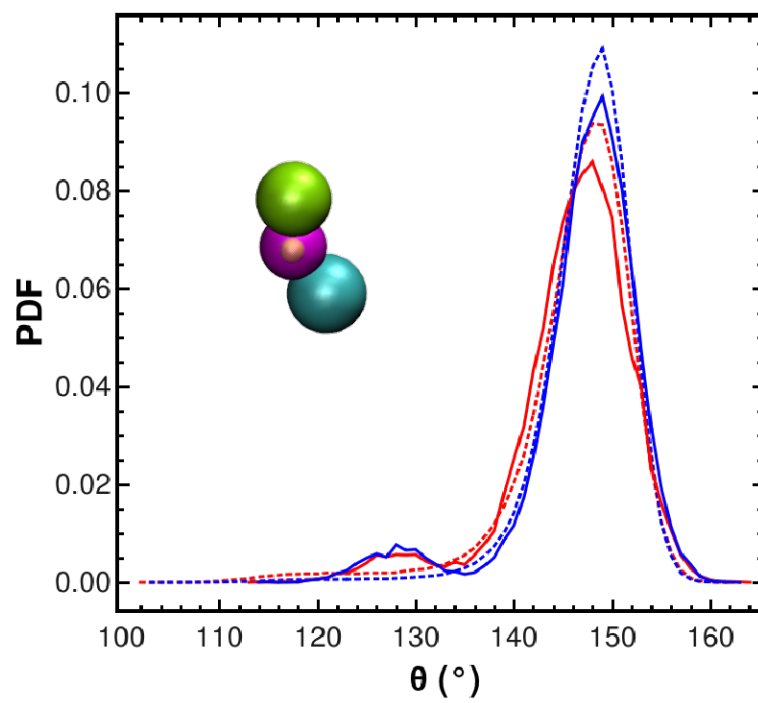


FIG. S7: Probability density function for the angle formed by the bonded particles SC3-SNda-SC1 (isopropyl-amide-backbone).

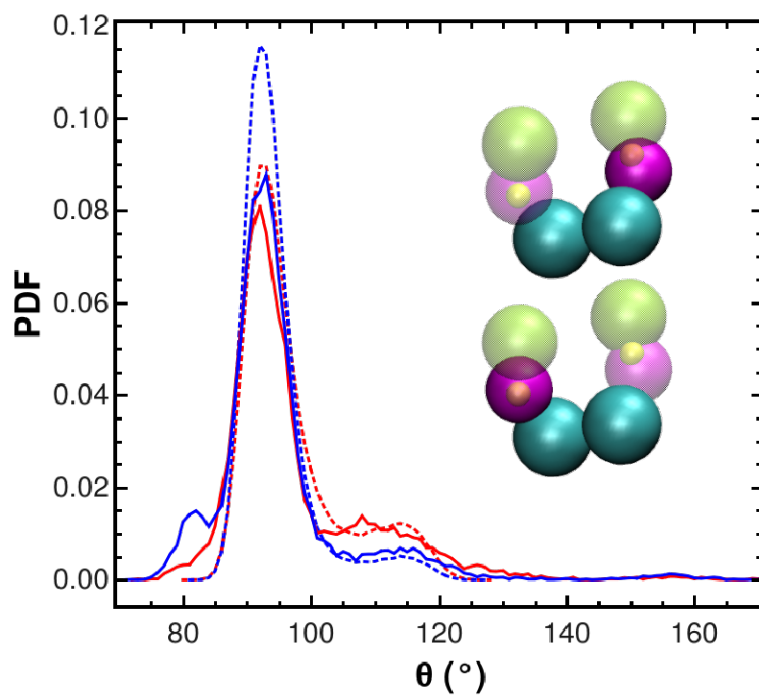


FIG. S8: Probability density function for the angle formed by the bonded particles SNda-SC1-SC1 (amide-backbone-backbone).

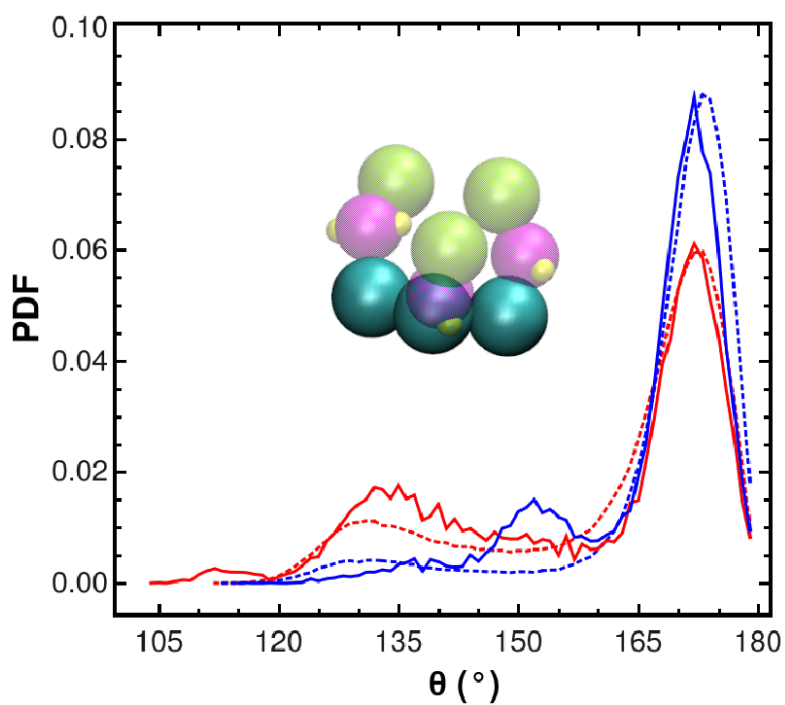


FIG. S9: Probability density function for the angle formed by the bonded particles SC1-SC1-SC1 (backbone-backbone-backbone).

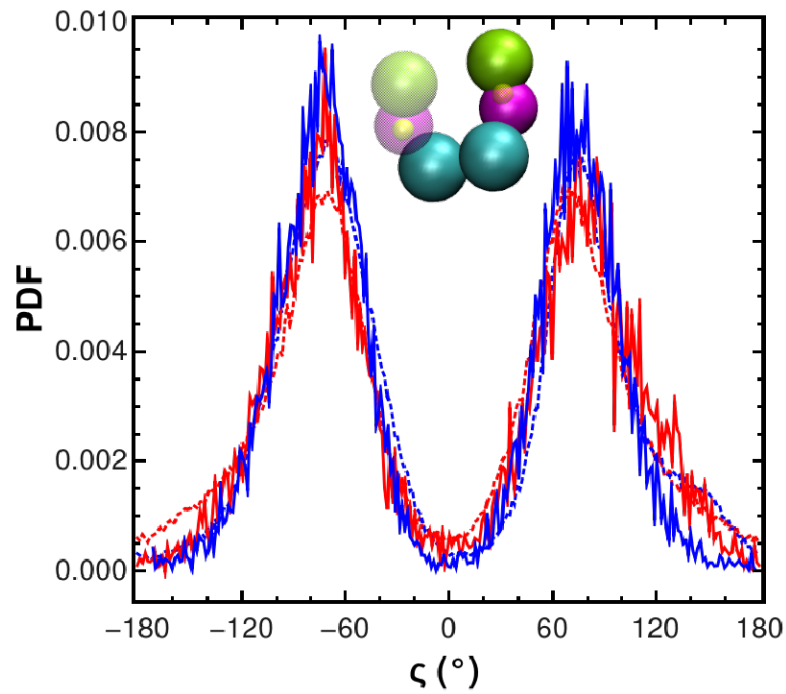


FIG. S10: Probability density function for the proper dihedral angle formed by the bonded particles SC3-SNda-SC1-SC1 (isopropyl-amide-backbone-backbone). It is important that this dihedral can jump from one configuration to the other. This allows the dipole moment of the SNda bead to align according to the local electrical field.

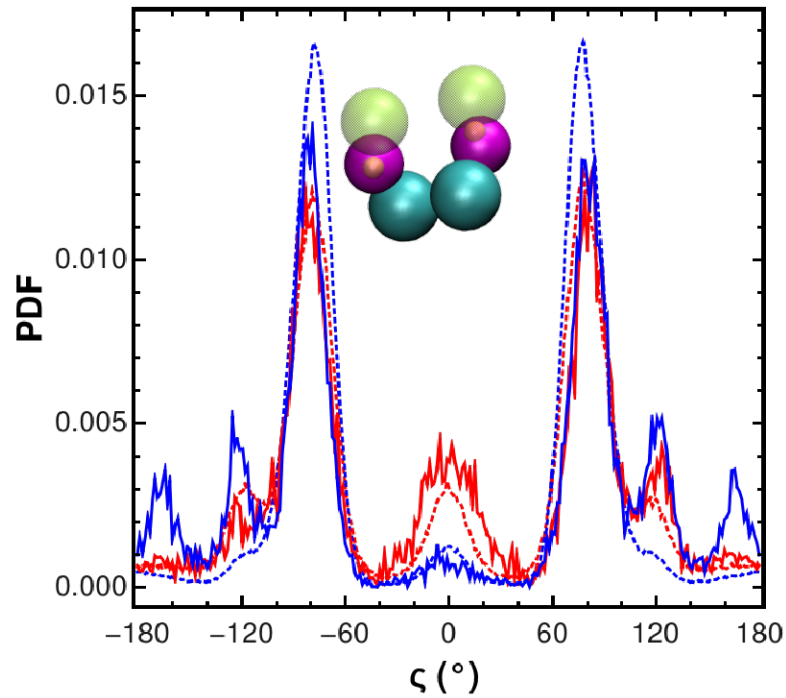


FIG. S11: Probability density function of the proper dihedral angle formed by the bonded particles SNda-SC1-SC1-SNda (amide-backbone-backbone-amide).

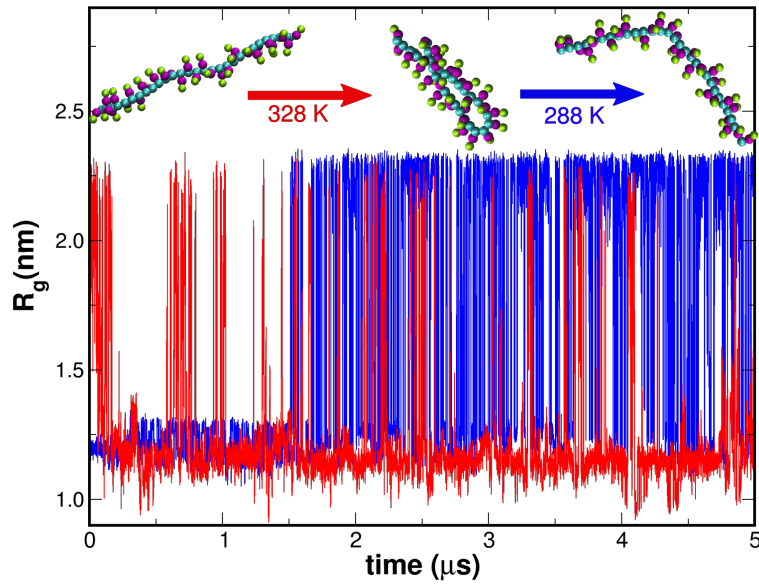


FIG. S12: Time evolution of the radius of gyration depicting the coil-to-globule reversibility. The red curve corresponds to 328 K when starting from a set of identical coil configurations. The blue curve corresponds to 288 K when starting from the final state of the red curve (a set of identical globule configurations). Both curves are obtained through REMD simulations.

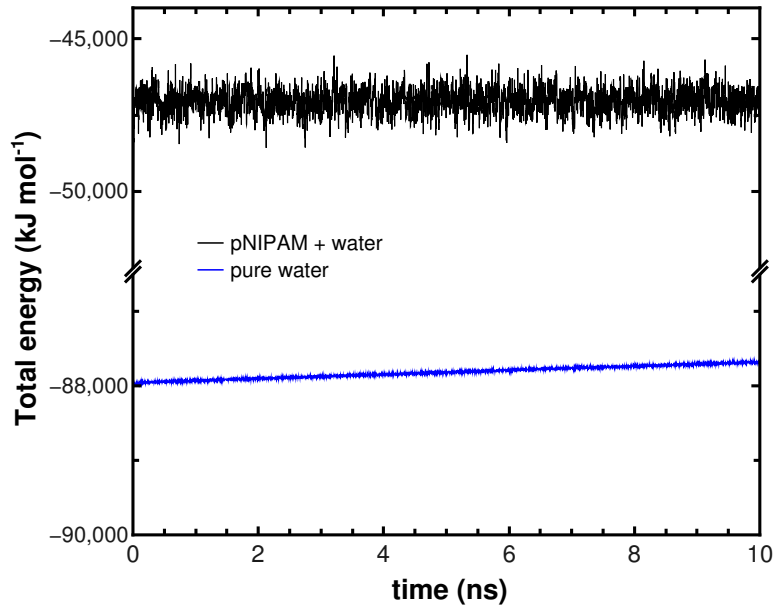


FIG. S13: Time evolution of the total energy at the microcanonical ensemble (298 K) for pure water (blue) and a single 30-mer in water (black). In both cases, the systems were firstly equilibrated employing the NPT ensemble.

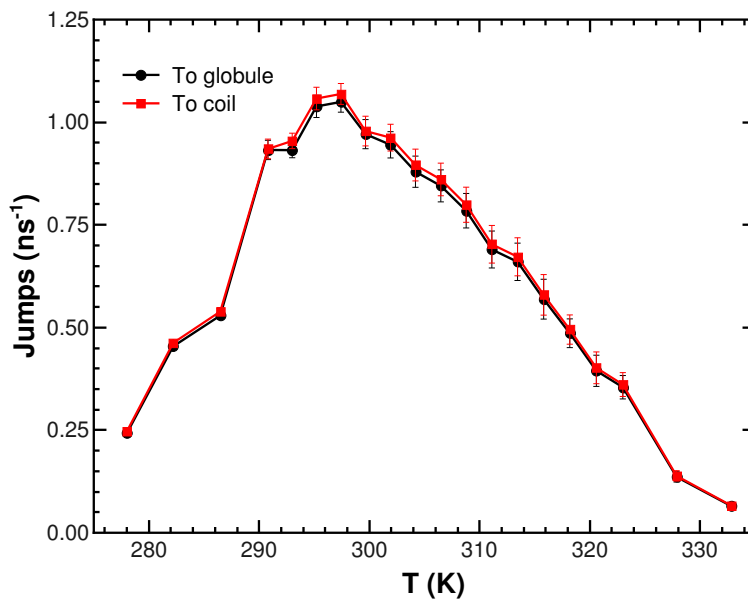


FIG. S14: Number of coil-to-globule (black) and globule-to-coil (red) jumps per ns detected along the REMD simulations as a function of temperature. Note that both curves are very similar at each temperature. Equilibrium of the whole set of replicas implies that the overall number of coil-to-globule jumps must be equal to the overall number of globule-to-coil ones. However, we observe equilibrium at all temperatures. Note that the curve peaks relatively close to the transition temperature.

II. TOPOLOGY FILES

The following topology files can be used to simulate a 35-mer of a CG pNIPAM. This *pnipam.itp* file must be attached to the corresponding *topology.top* file.

pnipam.itp

```
[ moleculetype ]
; molname nrexcl
PNIPA 1

[ atoms ]
;id type resnr residu atom chgrp charge mass
1 SC3 1 PNIPA SC3 1 0 43
2 SNda 1 PNIPA SNda 2 0 0
3 D 1 PNIPA D 3 0.48 21.5
4 D 1 PNIPA D 4 -0.48 21.5
5 SC1 1 PNIPA SC1 5 0 43
6 SC3 2 PNIPA SC3 1 0 43
7 SNda 2 PNIPA SNda 2 0 0
8 D 2 PNIPA D 3 0.48 21.5
9 D 2 PNIPA D 4 -0.48 21.5
10 SC1 2 PNIPA SC1 5 0 43
11 SC3 3 PNIPA SC3 1 0 43
12 SNda 3 PNIPA SNda 2 0 0
13 D 3 PNIPA D 3 0.48 21.5
14 D 3 PNIPA D 4 -0.48 21.5
15 SC1 3 PNIPA SC1 5 0 43
16 SC3 4 PNIPA SC3 1 0 43
17 SNda 4 PNIPA SNda 2 0 0
18 D 4 PNIPA D 3 0.48 21.5
```

19 D	4	PNIPA	D	4 -0.48 21.5
20 SC1	4	PNIPA	SC1	5 0 43
21 SC3	5	PNIPA	SC3	1 0 43
22 SNda	5	PNIPA	SNda	2 0 0
23 D	5	PNIPA	D	3 0.48 21.5
24 D	5	PNIPA	D	4 -0.48 21.5
25 SC1	5	PNIPA	SC1	5 0 43
26 SC3	6	PNIPA	SC3	1 0 43
27 SNda	6	PNIPA	SNda	2 0 0
28 D	6	PNIPA	D	3 0.48 21.5
29 D	6	PNIPA	D	4 -0.48 21.5
30 SC1	6	PNIPA	SC1	5 0 43
31 SC3	7	PNIPA	SC3	1 0 43
32 SNda	7	PNIPA	SNda	2 0 0
33 D	7	PNIPA	D	3 0.48 21.5
34 D	7	PNIPA	D	4 -0.48 21.5
35 SC1	7	PNIPA	SC1	5 0 43
36 SC3	8	PNIPA	SC3	1 0 43
37 SNda	8	PNIPA	SNda	2 0 0
38 D	8	PNIPA	D	3 0.48 21.5
39 D	8	PNIPA	D	4 -0.48 21.5
40 SC1	8	PNIPA	SC1	5 0 43
41 SC3	9	PNIPA	SC3	1 0 43
42 SNda	9	PNIPA	SNda	2 0 0
43 D	9	PNIPA	D	3 0.48 21.5
44 D	9	PNIPA	D	4 -0.48 21.5
45 SC1	9	PNIPA	SC1	5 0 43
46 SC3	10	PNIPA	SC3	1 0 43
47 SNda	10	PNIPA	SNda	2 0 0
48 D	10	PNIPA	D	3 0.48 21.5
49 D	10	PNIPA	D	4 -0.48 21.5
50 SC1	10	PNIPA	SC1	5 0 43
51 SC3	11	PNIPA	SC3	1 0 43
52 SNda	11	PNIPA	SNda	2 0 0
53 D	11	PNIPA	D	3 0.48 21.5
54 D	11	PNIPA	D	4 -0.48 21.5
55 SC1	11	PNIPA	SC1	5 0 43
56 SC3	12	PNIPA	SC3	1 0 43
57 SNda	12	PNIPA	SNda	2 0 0
58 D	12	PNIPA	D	3 0.48 21.5
59 D	12	PNIPA	D	4 -0.48 21.5
60 SC1	12	PNIPA	SC1	5 0 43
61 SC3	13	PNIPA	SC3	1 0 43
62 SNda	13	PNIPA	SNda	2 0 0
63 D	13	PNIPA	D	3 0.48 21.5
64 D	13	PNIPA	D	4 -0.48 21.5
65 SC1	13	PNIPA	SC1	5 0 43
66 SC3	14	PNIPA	SC3	1 0 43
67 SNda	14	PNIPA	SNda	2 0 0
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69 D	14	PNIPA	D	4 -0.48 21.5
70 SC1	14	PNIPA	SC1	5 0 43
71 SC3	15	PNIPA	SC3	1 0 43
72 SNda	15	PNIPA	SNda	2 0 0
73 D	15	PNIPA	D	3 0.48 21.5
74 D	15	PNIPA	D	4 -0.48 21.5
75 SC1	15	PNIPA	SC1	5 0 43

76	SC3	16	PNIPA	SC3	1 0 43
77	SNda	16	PNIPA	SNda	2 0 0
78	D	16	PNIPA	D	3 0.48 21.5
79	D	16	PNIPA	D	4 -0.48 21.5
80	SC1	16	PNIPA	SC1	5 0 43
81	SC3	17	PNIPA	SC3	1 0 43
82	SNda	17	PNIPA	SNda	2 0 0
83	D	17	PNIPA	D	3 0.48 21.5
84	D	17	PNIPA	D	4 -0.48 21.5
85	SC1	17	PNIPA	SC1	5 0 43
86	SC3	18	PNIPA	SC3	1 0 43
87	SNda	18	PNIPA	SNda	2 0 0
88	D	18	PNIPA	D	3 0.48 21.5
89	D	18	PNIPA	D	4 -0.48 21.5
90	SC1	18	PNIPA	SC1	5 0 43
91	SC3	19	PNIPA	SC3	1 0 43
92	SNda	19	PNIPA	SNda	2 0 0
93	D	19	PNIPA	D	3 0.48 21.5
94	D	19	PNIPA	D	4 -0.48 21.5
95	SC1	19	PNIPA	SC1	5 0 43
96	SC3	20	PNIPA	SC3	1 0 43
97	SNda	20	PNIPA	SNda	2 0 0
98	D	20	PNIPA	D	3 0.48 21.5
99	D	20	PNIPA	D	4 -0.48 21.5
100	SC1	20	PNIPA	SC1	5 0 43
101	SC3	21	PNIPA	SC3	1 0 43
102	SNda	21	PNIPA	SNda	2 0 0
103	D	21	PNIPA	D	3 0.48 21.5
104	D	21	PNIPA	D	4 -0.48 21.5
105	SC1	21	PNIPA	SC1	5 0 43
106	SC3	22	PNIPA	SC3	1 0 43
107	SNda	22	PNIPA	SNda	2 0 0
108	D	22	PNIPA	D	3 0.48 21.5
109	D	22	PNIPA	D	4 -0.48 21.5
110	SC1	22	PNIPA	SC1	5 0 43
111	SC3	23	PNIPA	SC3	1 0 43
112	SNda	23	PNIPA	SNda	2 0 0
113	D	23	PNIPA	D	3 0.48 21.5
114	D	23	PNIPA	D	4 -0.48 21.5
115	SC1	23	PNIPA	SC1	5 0 43
116	SC3	24	PNIPA	SC3	1 0 43
117	SNda	24	PNIPA	SNda	2 0 0
118	D	24	PNIPA	D	3 0.48 21.5
119	D	24	PNIPA	D	4 -0.48 21.5
120	SC1	24	PNIPA	SC1	5 0 43
121	SC3	25	PNIPA	SC3	1 0 43
122	SNda	25	PNIPA	SNda	2 0 0
123	D	25	PNIPA	D	3 0.48 21.5
124	D	25	PNIPA	D	4 -0.48 21.5
125	SC1	25	PNIPA	SC1	5 0 43
126	SC3	26	PNIPA	SC3	1 0 43
127	SNda	26	PNIPA	SNda	2 0 0
128	D	26	PNIPA	D	3 0.48 21.5
129	D	26	PNIPA	D	4 -0.48 21.5
130	SC1	26	PNIPA	SC1	5 0 43
131	SC3	27	PNIPA	SC3	1 0 43
132	SNda	27	PNIPA	SNda	2 0 0

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133 D    27  PNIPA  D    3 0.48 21.5
134 D    27  PNIPA  D    4 -0.48 21.5
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137 SNda  28  PNIPA  SNda 2 0 0
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139 D    28  PNIPA  D    4 -0.48 21.5
140 SC1   28  PNIPA  SC1  5 0 43
141 SC3   29  PNIPA  SC3  1 0 43
142 SNda  29  PNIPA  SNda 2 0 0
143 D    29  PNIPA  D    3 0.48 21.5
144 D    29  PNIPA  D    4 -0.48 21.5
145 SC1   29  PNIPA  SC1  5 0 43
146 SC3   30  PNIPA  SC3  1 0 43
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149 D    30  PNIPA  D    4 -0.48 21.5
150 SC1   30  PNIPA  SC1  5 0 43
151 SC3   31  PNIPA  SC3  1 0 43
152 SNda  31  PNIPA  SNda 2 0 0
153 D    31  PNIPA  D    3 0.48 21.5
154 D    31  PNIPA  D    4 -0.48 21.5
155 SC1   31  PNIPA  SC1  5 0 43
156 SC3   32  PNIPA  SC3  1 0 43
157 SNda  32  PNIPA  SNda 2 0 0
158 D    32  PNIPA  D    3 0.48 21.5
159 D    32  PNIPA  D    4 -0.48 21.5
160 SC1   32  PNIPA  SC1  5 0 43
161 SC3   33  PNIPA  SC3  1 0 43
162 SNda  33  PNIPA  SNda 2 0 0
163 D    33  PNIPA  D    3 0.48 21.5
164 D    33  PNIPA  D    4 -0.48 21.5
165 SC1   33  PNIPA  SC1  5 0 43
166 SC3   34  PNIPA  SC3  1 0 43
167 SNda  34  PNIPA  SNda 2 0 0
168 D    34  PNIPA  D    3 0.48 21.5
169 D    34  PNIPA  D    4 -0.48 21.5
170 SC1   34  PNIPA  SC1  5 0 43
171 SC3   35  PNIPA  SC3  1 0 43
172 SNda  35  PNIPA  SNda 2 0 0
173 D    35  PNIPA  D    3 0.48 21.5
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175 SC1   35  PNIPA  SC1  5 0 43

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142 145 1 0.242 25000
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167 170 1 0.242 25000
170 175 1 0.254 20000
171 172 1 0.268 20500
172 175 1 0.242 25000

```

```
[ angles ]
```

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; ai aj ak funct angle fc
```

```

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1 2 4 2 119.00 150.00
3 2 4 2 161.00 150.00
3 2 5 2 90.00 150.00
4 2 5 2 90.00 150.00
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5 10 7 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
5 10 15 6 166.158 6.718 -40.527 -1 176.41 179.444
6 7 10 6 138.52 3.63 -29.06 -32.86 350.6 650.93
6 7 8 2 61.00 200.00
6 7 9 2 119.00 150.00
8 7 9 2 161.00 150.00
8 7 10 2 90.00 150.00
9 7 10 2 90.00 150.00
7 10 15 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
10 15 12 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
10 15 20 6 166.158 6.718 -40.527 -1 176.41 179.444
11 12 15 6 138.52 3.63 -29.06 -32.86 350.6 650.93
11 12 13 2 61.00 200.00
11 12 14 2 119.00 150.00
13 12 14 2 161.00 150.00
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15 20 17 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
15 20 25 6 166.158 6.718 -40.527 -1 176.41 179.444
16 17 20 6 138.52 3.63 -29.06 -32.86 350.6 650.93
16 17 18 2 61.00 200.00
16 17 19 2 119.00 150.00
18 17 19 2 161.00 150.00
18 17 20 2 90.00 150.00
19 17 20 2 90.00 150.00
17 20 25 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
20 25 22 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
20 25 30 6 166.158 6.718 -40.527 -1 176.41 179.444
21 22 25 6 138.52 3.63 -29.06 -32.86 350.6 650.93

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 21 22 24 2 119.00 150.00
 23 22 24 2 161.00 150.00
 23 22 25 2 90.00 150.00
 24 22 25 2 90.00 150.00
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 25 30 35 6 166.158 6.718 -40.527 -1 176.41 179.444
 26 27 30 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 26 27 28 2 61.00 200.00
 26 27 29 2 119.00 150.00
 28 27 29 2 161.00 150.00
 28 27 30 2 90.00 150.00
 29 27 30 2 90.00 150.00
 27 30 35 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 30 35 32 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 30 35 40 6 166.158 6.718 -40.527 -1 176.41 179.444
 31 32 35 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 31 32 33 2 61.00 200.00
 31 32 34 2 119.00 150.00
 33 32 34 2 161.00 150.00
 33 32 35 2 90.00 150.00
 34 32 35 2 90.00 150.00
 32 35 40 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 35 40 37 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 35 40 45 6 166.158 6.718 -40.527 -1 176.41 179.444
 36 37 40 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 36 37 38 2 61.00 200.00
 36 37 39 2 119.00 150.00
 38 37 39 2 161.00 150.00
 38 37 40 2 90.00 150.00
 39 37 40 2 90.00 150.00
 37 40 45 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 40 45 42 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 40 45 50 6 166.158 6.718 -40.527 -1 176.41 179.444
 41 42 45 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 41 42 43 2 61.00 200.00
 41 42 44 2 119.00 150.00
 43 42 44 2 161.00 150.00
 43 42 45 2 90.00 150.00
 44 42 45 2 90.00 150.00
 42 45 50 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 45 50 47 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 45 50 55 6 166.158 6.718 -40.527 -1 176.41 179.444
 46 47 50 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 46 47 48 2 61.00 200.00
 46 47 49 2 119.00 150.00
 48 47 49 2 161.00 150.00
 48 47 50 2 90.00 150.00
 49 47 50 2 90.00 150.00
 47 50 55 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 50 55 52 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 50 55 60 6 166.158 6.718 -40.527 -1 176.41 179.444
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 53 52 54 2 161.00 150.00

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 56 57 59 2 119.00 150.00
 58 57 59 2 161.00 150.00
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 59 57 60 2 90.00 150.00
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 60 65 62 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 60 65 70 6 166.158 6.718 -40.527 -1 176.41 179.444
 61 62 65 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 61 62 63 2 61.00 200.00
 61 62 64 2 119.00 150.00
 63 62 64 2 161.00 150.00
 63 62 65 2 90.00 150.00
 64 62 65 2 90.00 150.00
 62 65 70 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 65 70 67 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 65 70 75 6 166.158 6.718 -40.527 -1 176.41 179.444
 66 67 70 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 66 67 68 2 61.00 200.00
 66 67 69 2 119.00 150.00
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 70 75 80 6 166.158 6.718 -40.527 -1 176.41 179.444
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 71 72 74 2 119.00 150.00
 73 72 74 2 161.00 150.00
 73 72 75 2 90.00 150.00
 74 72 75 2 90.00 150.00
 72 75 80 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 75 80 77 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 75 80 85 6 166.158 6.718 -40.527 -1 176.41 179.444
 76 77 80 6 138.52 3.63 -29.06 -32.86 350.6 650.93
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 76 77 79 2 119.00 150.00
 78 77 79 2 161.00 150.00
 78 77 80 2 90.00 150.00
 79 77 80 2 90.00 150.00
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 80 85 82 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 80 85 90 6 166.158 6.718 -40.527 -1 176.41 179.444
 81 82 85 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 81 82 83 2 61.00 200.00
 81 82 84 2 119.00 150.00
 83 82 84 2 161.00 150.00
 83 82 85 2 90.00 150.00
 84 82 85 2 90.00 150.00
 82 85 90 6 98.09 3.95 37.9 -38.2 -575.7 1400.7

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 85 90 95 6 166.158 6.718 -40.527 -1 176.41 179.444
 86 87 90 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 86 87 88 2 61.00 200.00
 86 87 89 2 119.00 150.00
 88 87 89 2 161.00 150.00
 88 87 90 2 90.00 150.00
 89 87 90 2 90.00 150.00
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 90 95 92 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
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 91 92 95 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 91 92 93 2 61.00 200.00
 91 92 94 2 119.00 150.00
 93 92 94 2 161.00 150.00
 93 92 95 2 90.00 150.00
 94 92 95 2 90.00 150.00
 92 95 100 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 95 100 97 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
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 96 97 100 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 96 97 98 2 61.00 200.00
 96 97 99 2 119.00 150.00
 98 97 99 2 161.00 150.00
 98 97 100 2 90.00 150.00
 99 97 100 2 90.00 150.00
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 100 105 102 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
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 104 102 105 2 90.00 150.00
 102 105 110 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
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 106 107 108 2 61.00 200.00
 106 107 109 2 119.00 150.00
 108 107 109 2 161.00 150.00
 108 107 110 2 90.00 150.00
 109 107 110 2 90.00 150.00
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 110 115 112 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
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 111 112 115 6 138.52 3.63 -29.06 -32.86 350.6 650.93
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 111 112 114 2 119.00 150.00
 113 112 114 2 161.00 150.00
 113 112 115 2 90.00 150.00
 114 112 115 2 90.00 150.00
 112 115 120 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 115 120 117 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 115 120 125 6 166.158 6.718 -40.527 -1 176.41 179.444
 116 117 120 6 138.52 3.63 -29.06 -32.86 350.6 650.93

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 116 117 119 2 119.00 150.00
 118 117 119 2 161.00 150.00
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 119 117 120 2 90.00 150.00
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 120 125 122 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
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 121 122 124 2 119.00 150.00
 123 122 124 2 161.00 150.00
 123 122 125 2 90.00 150.00
 124 122 125 2 90.00 150.00
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 131 132 135 6 138.52 3.63 -29.06 -32.86 350.6 650.93
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 139 137 140 2 90.00 150.00
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 140 145 150 6 166.158 6.718 -40.527 -1 176.41 179.444
 141 142 145 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 141 142 143 2 61.00 200.00
 141 142 144 2 119.00 150.00
 143 142 144 2 161.00 150.00
 143 142 145 2 90.00 150.00
 144 142 145 2 90.00 150.00
 142 145 150 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 145 150 147 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
 145 150 155 6 166.158 6.718 -40.527 -1 176.41 179.444
 146 147 150 6 138.52 3.63 -29.06 -32.86 350.6 650.93
 146 147 148 2 61.00 200.00
 146 147 149 2 119.00 150.00
 148 147 149 2 161.00 150.00

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148 147 150 2 90.00 150.00
149 147 150 2 90.00 150.00
147 150 155 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
150 155 152 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
150 155 160 6 166.158 6.718 -40.527 -1 176.41 179.444
151 152 155 6 138.52 3.63 -29.06 -32.86 350.6 650.93
151 152 153 2 61.00 200.00
151 152 154 2 119.00 150.00
153 152 154 2 161.00 150.00
153 152 155 2 90.00 150.00
154 152 155 2 90.00 150.00
152 155 160 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
155 160 157 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
155 160 165 6 166.158 6.718 -40.527 -1 176.41 179.444
156 157 160 6 138.52 3.63 -29.06 -32.86 350.6 650.93
156 157 158 2 61.00 200.00
156 157 159 2 119.00 150.00
158 157 159 2 161.00 150.00
158 157 160 2 90.00 150.00
159 157 160 2 90.00 150.00
157 160 165 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
160 165 162 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
160 165 170 6 166.158 6.718 -40.527 -1 176.41 179.444
161 162 165 6 138.52 3.63 -29.06 -32.86 350.6 650.93
161 162 163 2 61.00 200.00
161 162 164 2 119.00 150.00
163 162 164 2 161.00 150.00
163 162 165 2 90.00 150.00
164 162 165 2 90.00 150.00
162 165 170 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
165 170 167 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
165 170 175 6 166.158 6.718 -40.527 -1 176.41 179.444
166 167 170 6 138.52 3.63 -29.06 -32.86 350.6 650.93
166 167 168 2 61.00 200.00
166 167 169 2 119.00 150.00
168 167 169 2 161.00 150.00
168 167 170 2 90.00 150.00
169 167 170 2 90.00 150.00
167 170 175 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
170 175 172 6 98.09 3.95 37.9 -38.2 -575.7 1400.7
171 172 175 6 138.52 3.63 -29.06 -32.86 350.6 650.93
171 172 173 2 61.00 200.00
171 172 174 2 119.00 150.00
173 172 174 2 161.00 150.00
173 172 175 2 90.00 150.00
174 172 175 2 90.00 150.00

[ dihedrals ] ; ai aj ak funct C0 C1 C2 C3 C4 C5 1 2 5 10 3 1.35 4.30 1.3 -5.8 1.30 1.60
2 5 10 7 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
6 7 10 15 3 1.35 4.30 1.3 -5.8 1.30 1.60
7 10 15 12 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
11 12 15 20 3 1.35 4.30 1.3 -5.8 1.30 1.60
12 15 20 17 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
16 17 20 25 3 1.35 4.30 1.3 -5.8 1.30 1.60
17 20 25 22 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
21 22 25 30 3 1.35 4.30 1.3 -5.8 1.30 1.60
22 25 30 27 3 -1.20 8.5 17.7 -27.7 -17.1 25.5

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 27 30 35 32 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 31 32 35 40 3 1.35 4.30 1.3 -5.8 1.30 1.60
 32 35 40 37 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 36 37 40 45 3 1.35 4.30 1.3 -5.8 1.30 1.60
 37 40 45 42 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 41 42 45 50 3 1.35 4.30 1.3 -5.8 1.30 1.60
 42 45 50 47 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 46 47 50 55 3 1.35 4.30 1.3 -5.8 1.30 1.60
 47 50 55 52 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
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 56 57 60 65 3 1.35 4.30 1.3 -5.8 1.30 1.60
 57 60 65 62 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 61 62 65 70 3 1.35 4.30 1.3 -5.8 1.30 1.60
 62 65 70 67 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 66 67 70 75 3 1.35 4.30 1.3 -5.8 1.30 1.60
 67 70 75 72 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 71 72 75 80 3 1.35 4.30 1.3 -5.8 1.30 1.60
 72 75 80 77 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 76 77 80 85 3 1.35 4.30 1.3 -5.8 1.30 1.60
 77 80 85 82 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 81 82 85 90 3 1.35 4.30 1.3 -5.8 1.30 1.60
 82 85 90 87 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 86 87 90 95 3 1.35 4.30 1.3 -5.8 1.30 1.60
 87 90 95 92 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 91 92 95 100 3 1.35 4.30 1.3 -5.8 1.30 1.60
 92 95 100 97 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 96 97 100 105 3 1.35 4.30 1.3 -5.8 1.30 1.60
 97 100 105 102 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
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 106 107 110 115 3 1.35 4.30 1.3 -5.8 1.30 1.60
 107 110 115 112 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 111 112 115 120 3 1.35 4.30 1.3 -5.8 1.30 1.60
 112 115 120 117 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 116 117 120 125 3 1.35 4.30 1.3 -5.8 1.30 1.60
 117 120 125 122 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 121 122 125 130 3 1.35 4.30 1.3 -5.8 1.30 1.60
 122 125 130 127 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 126 127 130 135 3 1.35 4.30 1.3 -5.8 1.30 1.60
 127 130 135 132 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 131 132 135 140 3 1.35 4.30 1.3 -5.8 1.30 1.60
 132 135 140 137 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 136 137 140 145 3 1.35 4.30 1.3 -5.8 1.30 1.60
 137 140 145 142 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 141 142 145 150 3 1.35 4.30 1.3 -5.8 1.30 1.60
 142 145 150 147 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 146 147 150 155 3 1.35 4.30 1.3 -5.8 1.30 1.60
 147 150 155 152 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 151 152 155 160 3 1.35 4.30 1.3 -5.8 1.30 1.60
 152 155 160 157 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 156 157 160 165 3 1.35 4.30 1.3 -5.8 1.30 1.60
 157 160 165 162 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 161 162 165 170 3 1.35 4.30 1.3 -5.8 1.30 1.60
 162 165 170 167 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
 166 167 170 175 3 1.35 4.30 1.3 -5.8 1.30 1.60

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167 170 175 172 3 -1.20 8.5 17.7 -27.7 -17.1 25.5
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[ constraints ]
; i j funct length
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8 9 1 0.226
13 14 1 0.226
18 19 1 0.226
23 24 1 0.226
28 29 1 0.226
33 34 1 0.226
38 39 1 0.226
43 44 1 0.226
48 49 1 0.226
53 54 1 0.226
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63 64 1 0.226
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83 84 1 0.226
88 89 1 0.226
93 94 1 0.226
98 99 1 0.226
103 104 1 0.226
108 109 1 0.226
113 114 1 0.226
118 119 1 0.226
123 124 1 0.226
128 129 1 0.226
133 134 1 0.226
138 139 1 0.226
143 144 1 0.226
148 149 1 0.226
153 154 1 0.226
158 159 1 0.226
163 164 1 0.226
168 169 1 0.226
173 174 1 0.226
```

The following **[nonbond_params]** must replace the corresponding lines of the original Martini force field file.
martini.itp

```
[ nonbond_params ]

;;; self terms
; i j f c6 c12
SNda SNda 1 0.30788E-01 0.78995E-04
SC3 SC3 1 0.73073E-01 0.60678E-03
SC1 SC1 1 0.60685E-01 0.38361E-03

;;; cross terms
; i j f c6 c12
SNda SC3 1 0.43179E-01 0.35855E-03
SNda SC1 1 0.19660E-01 0.80530E-04
SC3 SC1 1 0.58156E-01 0.36762E-03
```

```
;;; water cross terms
POL SNda 1 0.57344E-01 0.23488E-03
POL SC3 1 0.32768E-01 0.13421E-03
POL SC1 1 0.28149E-01 0.99052E-04
;;;
```