

**Conformational and Entropy Analyses of
Extended Molecular Dynamics Simulations of α -, β - and γ -Cyclodextrin
and of the β -Cyclodextron/Nabumetone Complex**

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Supporting Information

Figure S1. Histograms of the ϕ and ψ dihedral angles computed for all the glucose units during the production phase of the MD simulations of native α -, β -, and γ -CDs.

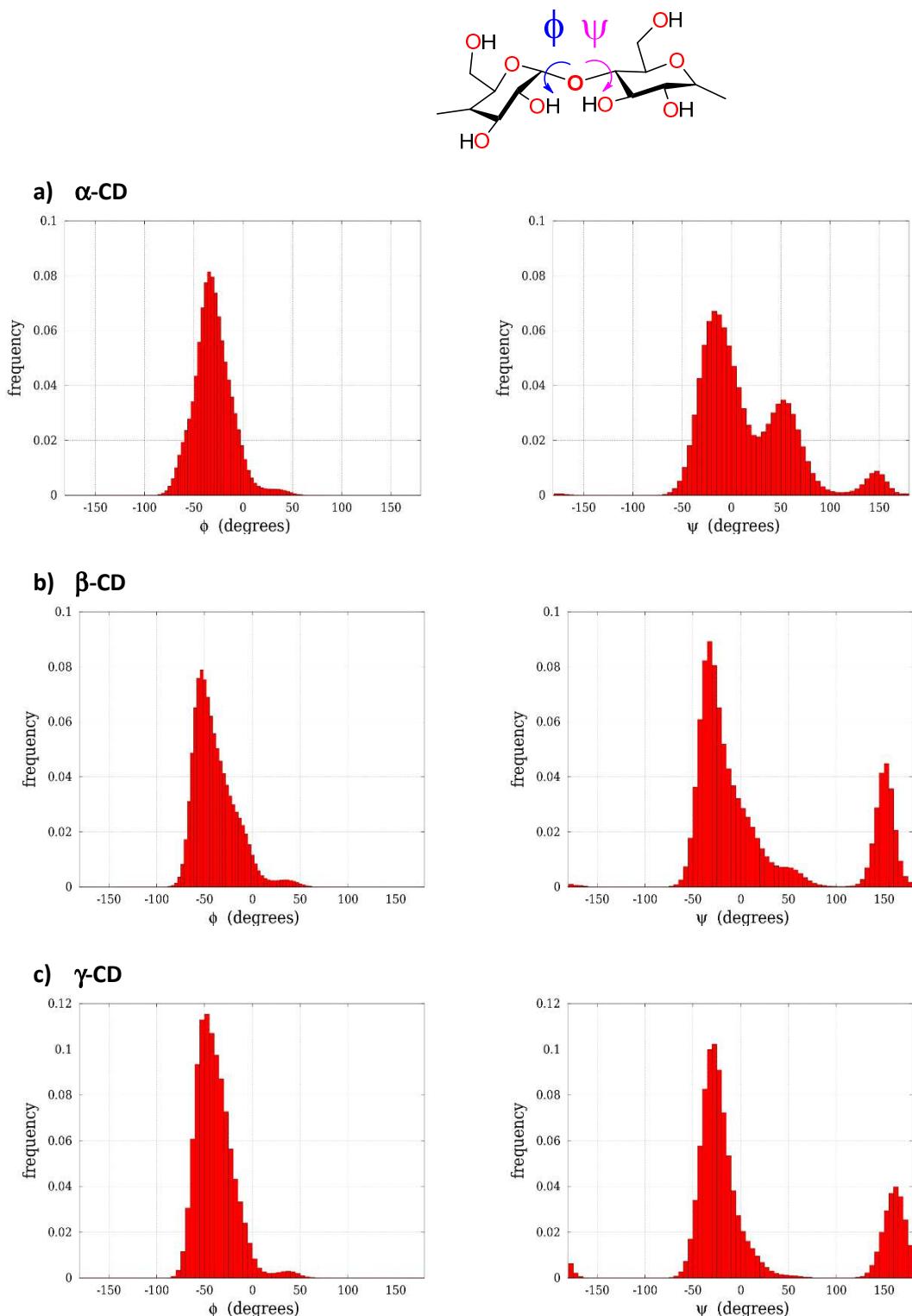


Figure S2. Time evolution of the radius of accessibility of the internal cavity of the CDs during the MD simulations.

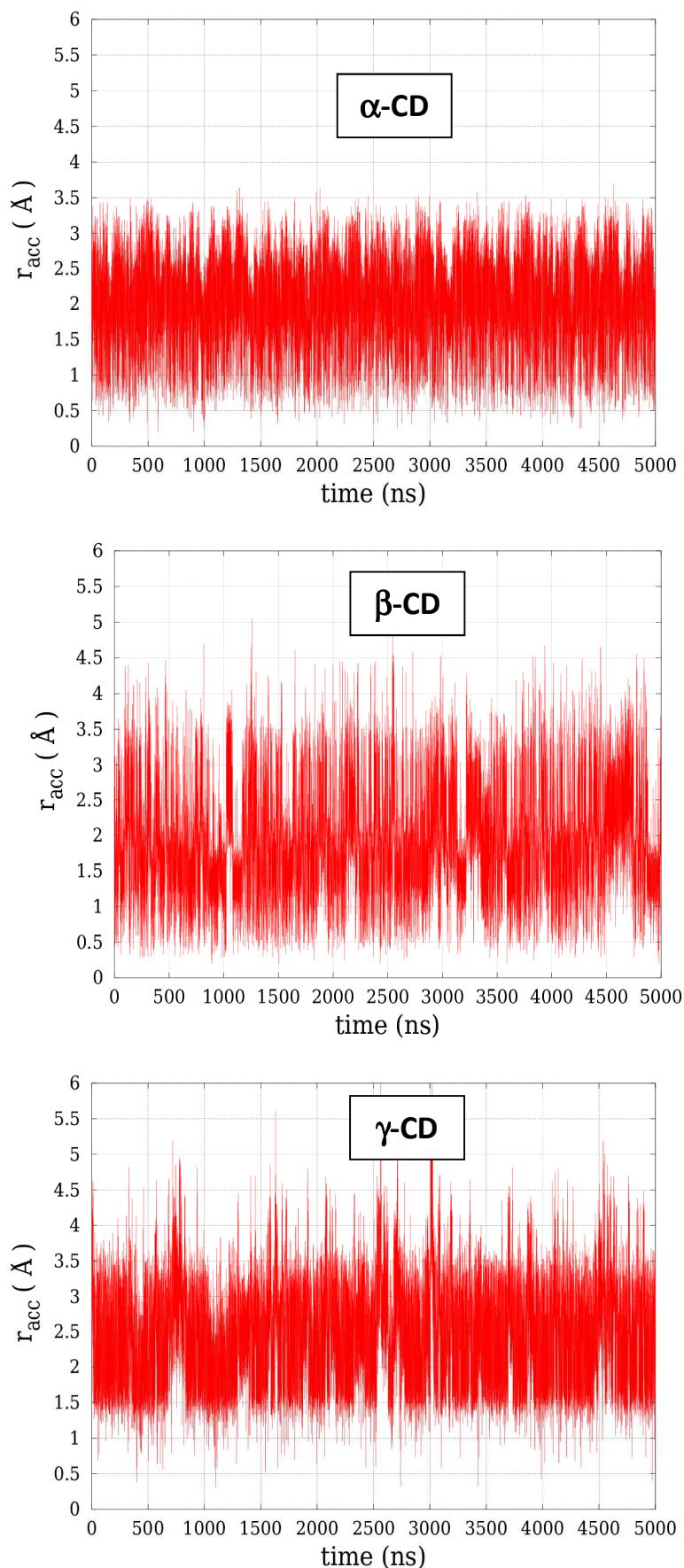


Figure S3. Correlation plots between r_{acc} values and the number of water molecules for CD structures showing inclusion complexes with water molecules.

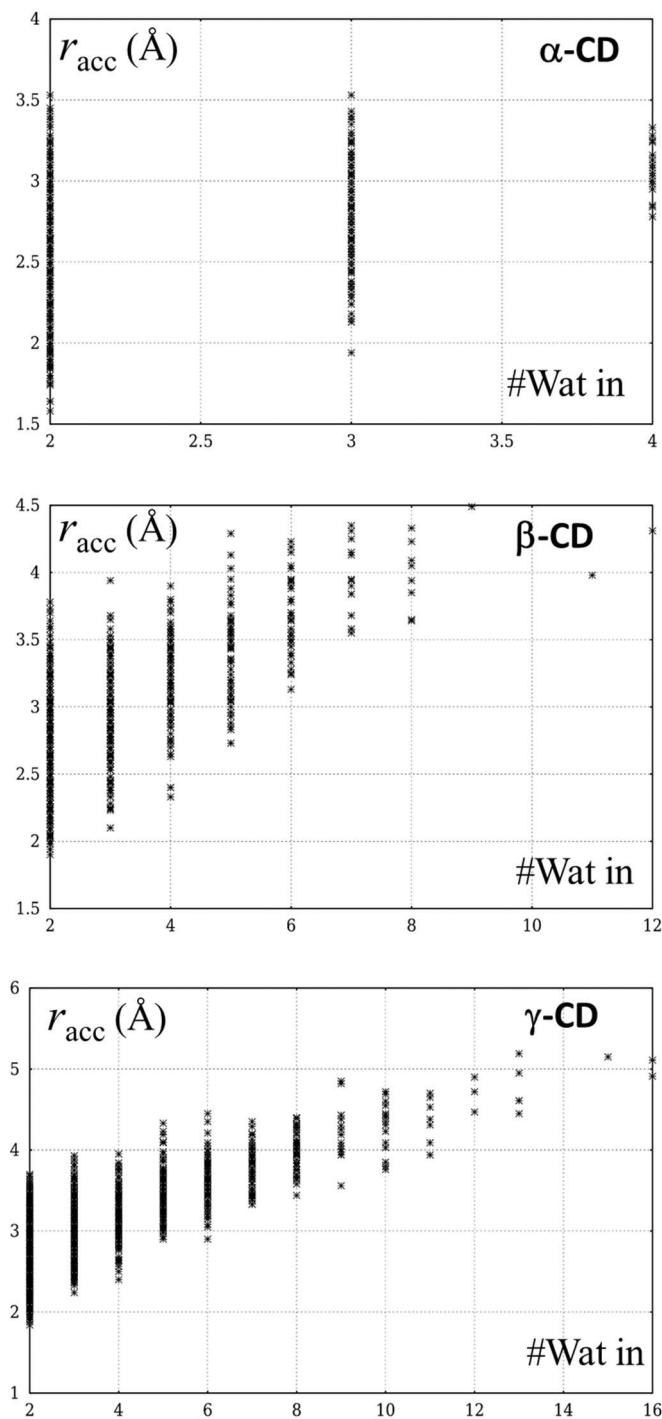
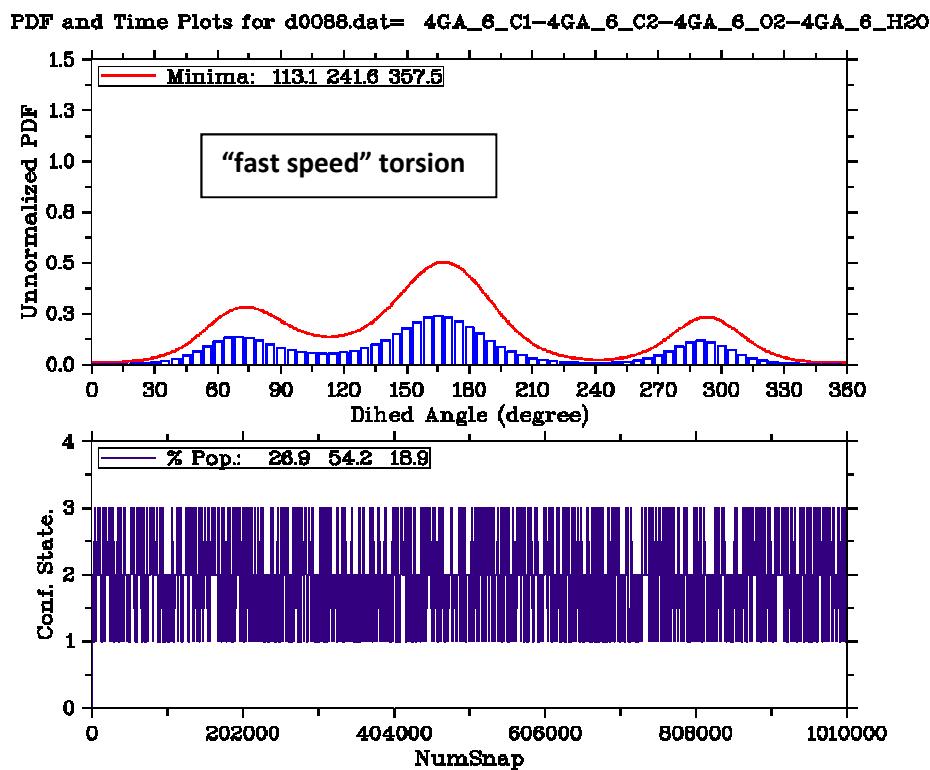


Figure S4. In the upper plot, probability density function for selected dihedral angles of γ -CD (a-d), β -CD (e) and α -CD (f), as obtained from a histogram representation and a Von-Mises kernel estimator (red curve). The lower plot shows the time evolution of the associated discrete variables. Note that plots (d-f) correspond to conformational motions around the glycosidic C4-O4 bond although the definition of the corresponding dihedral angle does not match that of the ψ angle.

(a)



(b)

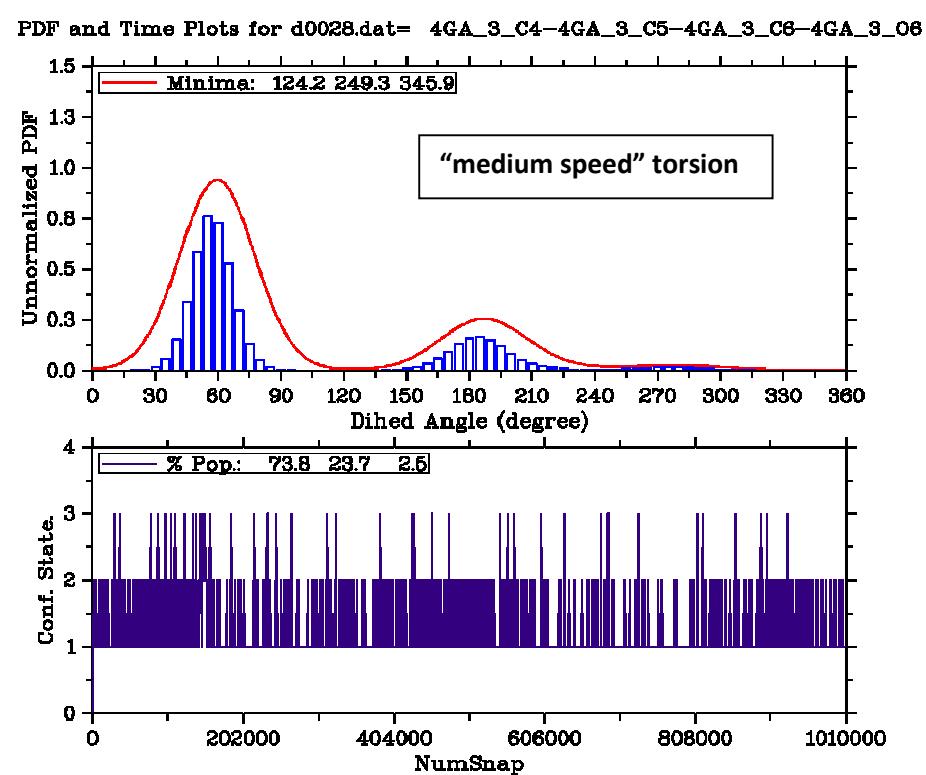
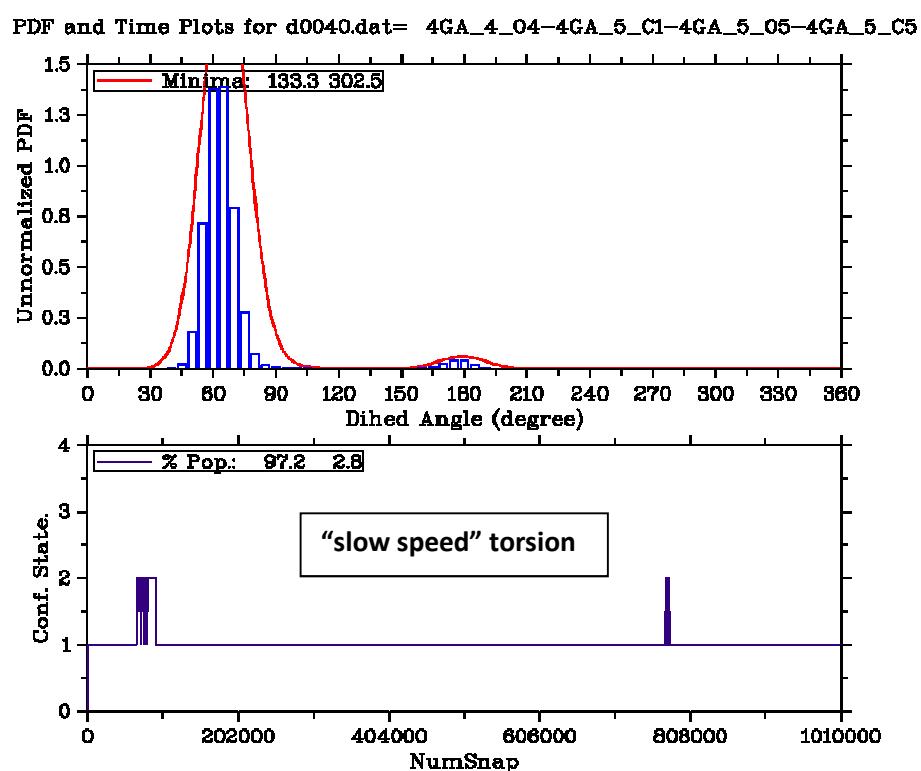


Figure S4 (cont).

(c)



(d)

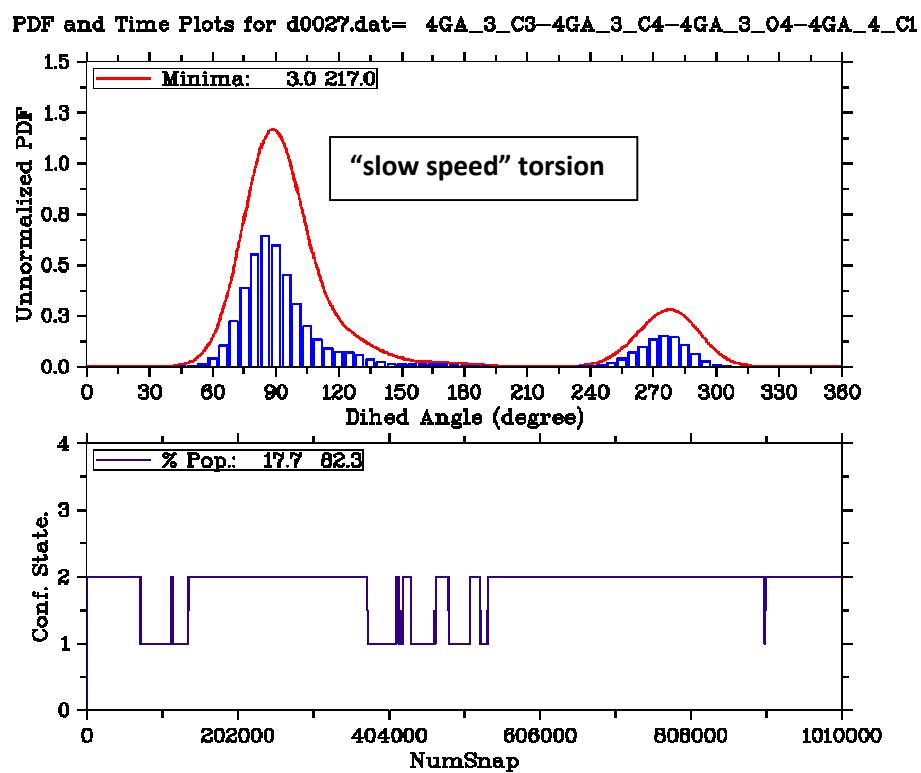
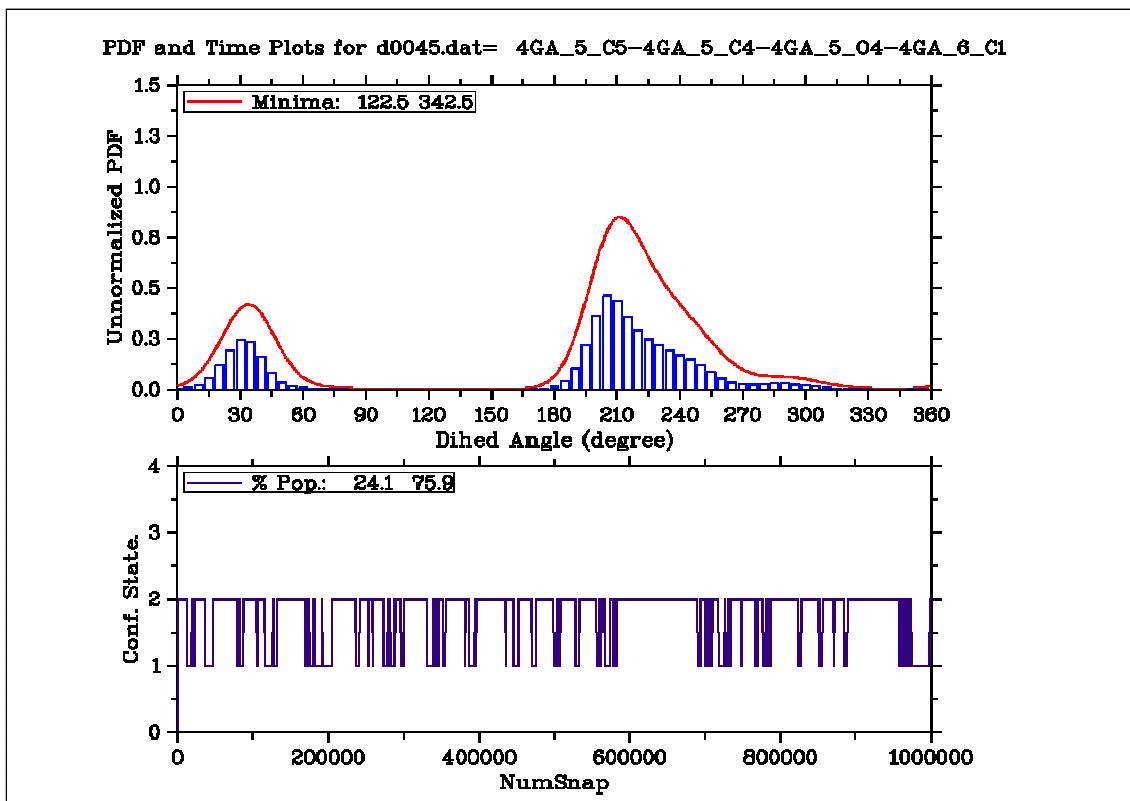


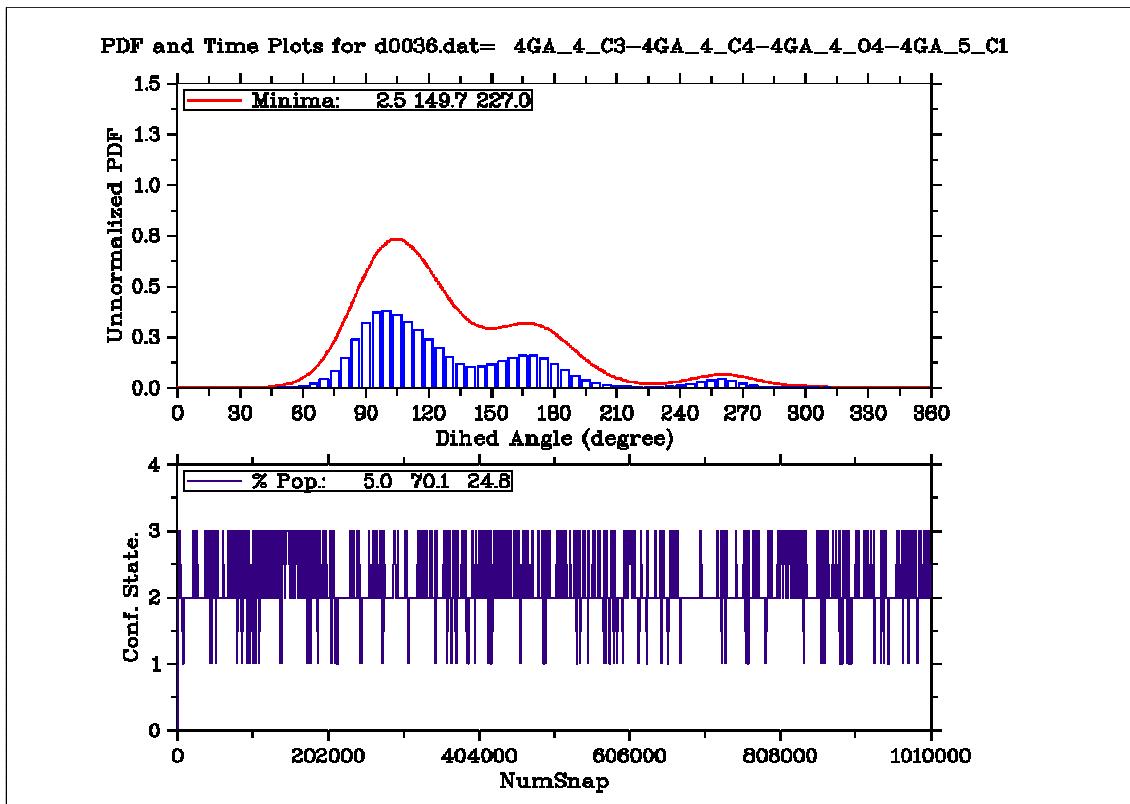
Figure S4

(cont.).

(e)



(f)



mol2 file containing the initial geometry of the nabumetone ligand, GAFF atom types and HF/6-31G* RESP charges.

```
@<TRIPOS>MOLECULE
LIG
 33      34      1      0      0
SMALL
rc
@<TRIPOS>ATOM
 1 C1      5.9420   -0.0700    1.3530 c3      1 LIG      -0.247874
 2 C2      5.0270   -0.1780    0.1510 c      1 LIG      0.602829
 3 C3      3.5490    0.0590    0.4040 c3      1 LIG      -0.028607
 4 C4      2.6850   -0.0680   -0.8540 c3      1 LIG      -0.028958
 5 C5      1.2200    0.1810   -0.5700 ca      1 LIG      -0.002054
 6 C6      0.3320   -0.8420   -0.4430 ca      1 LIG      -0.192815
 7 C7     -1.0390   -0.6100   -0.1520 ca      1 LIG      0.089967
 8 C8     -1.9710   -1.6730   -0.0110 ca      1 LIG      -0.266465
 9 C9     -3.2760   -1.4250    0.2680 ca      1 LIG      -0.140046
10 C10    -3.7290   -0.0910    0.4210 ca      1 LIG      0.243615
11 O1     -5.0370    0.1330    0.7290 os      1 LIG      -0.356695
12 C11    -5.9290    0.1040   -0.3570 c3      1 LIG      0.033626
13 C12    -2.8640    0.9480    0.2960 ca      1 LIG      -0.248428
14 C13    -1.4950    0.7110    0.0090 ca      1 LIG      0.098450
15 C14    -0.5580    1.7700   -0.1310 ca      1 LIG      -0.225442
16 C15     0.7450    1.5130   -0.4100 ca      1 LIG      -0.158629
17 O2      5.4510   -0.4360   -0.9340 o       1 LIG      -0.539464
18 H1      6.9630   -0.2610    1.0520 hc      1 LIG      0.064493
19 H2      5.6490   -0.7850    2.1160 hc      1 LIG      0.064493
20 H3      5.8720    0.9190    1.7940 hc      1 LIG      0.064493
21 H4      3.4370    1.0440    0.8530 hc      1 LIG      0.010801
22 H5      3.2160   -0.6440    1.1640 hc      1 LIG      0.010801
23 H6      2.8180   -1.0560   -1.2760 hc      1 LIG      0.032317
24 H7      3.0430    0.6320   -1.6020 hc      1 LIG      0.032317
25 H8      0.6620   -1.8600   -0.5660 ha      1 LIG      0.144455
26 H9     -1.6260   -2.6860   -0.1230 ha      1 LIG      0.166423
27 H10    -3.9800   -2.2280    0.3930 ha      1 LIG      0.142664
28 H11    -6.9150    0.2890    0.0440 h1      1 LIG      0.059444
29 H12    -5.9190   -0.8620   -0.8510 h1      1 LIG      0.059444
30 H13    -5.6820    0.8730   -1.0820 h1      1 LIG      0.059444
31 H14    -3.2220    1.9530    0.4270 ha      1 LIG      0.153949
32 H15    -0.8980    2.7850   -0.0180 ha      1 LIG      0.154145
33 H16     1.4360    2.3300   -0.5220 ha      1 LIG      0.147309

@<TRIPOS>BOND
 1   1   2  1
 2   1   18 1
 3   1   19 1
 4   1   20 1
 5   2   3  1
 6   2   17 2
 7   3   4  1
 8   3   21 1
 9   3   22 1
10   4   5  1
11   4   23 1
12   4   24 1
13   5   6 ar
14   5   16 ar
15   6   7 ar
16   6   25 1
17   7   8 ar
18   7   14 ar
19   8   9 ar
20   8   26 1
21   9   10 ar
22   9   27 1
23  10   11 1
24  10   13 ar
25  11   12 1
26  12   28 1
27  12   29 1
28  12   30 1
29  13   14 ar
30  13   31 1
31  14   15 ar
32  15   16 ar
33  15   32 1
34  16   33 1

@<TRIPOS>SUBSTRUCTURE
 1 LIG      1 TEMP      0 ****      0 ROOT
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