

## Supplementary Material

for

### **Porous organic cage as an inhibitor of A $\beta$ <sub>42</sub> peptide: a simulation study**

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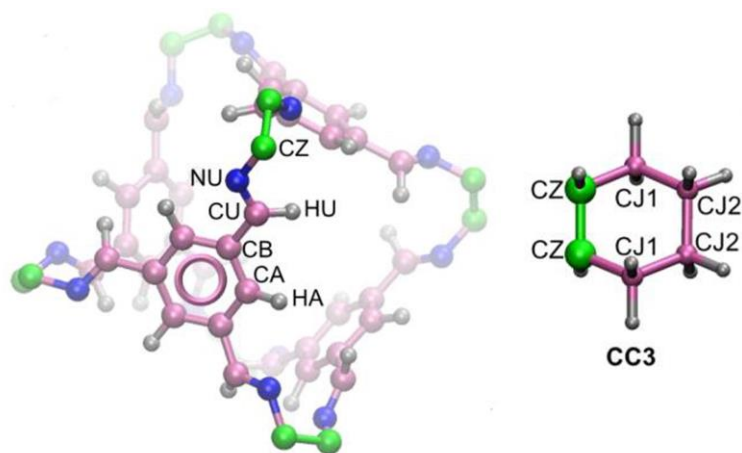
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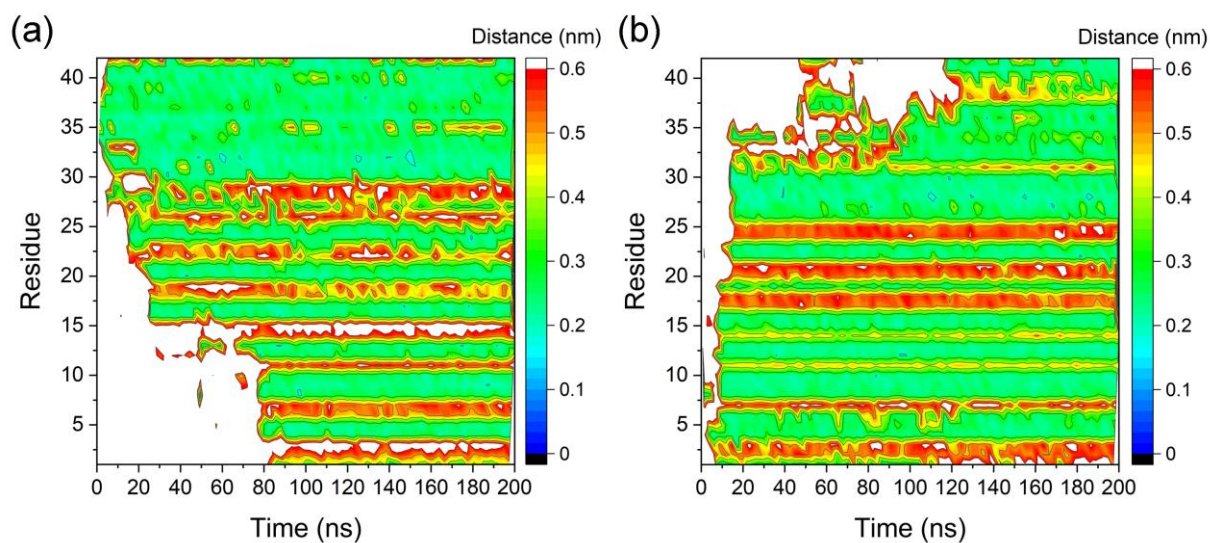
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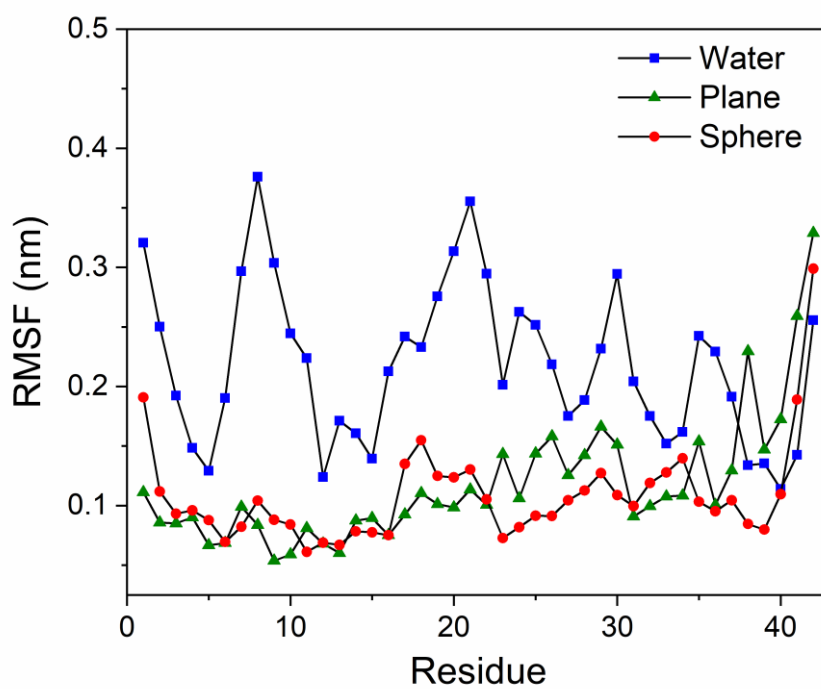
**Figure S1.** Atom types in CC3<sup>1,2</sup>.

**Table S1.** Non-bonded parameters in CC3<sup>1,2</sup>.

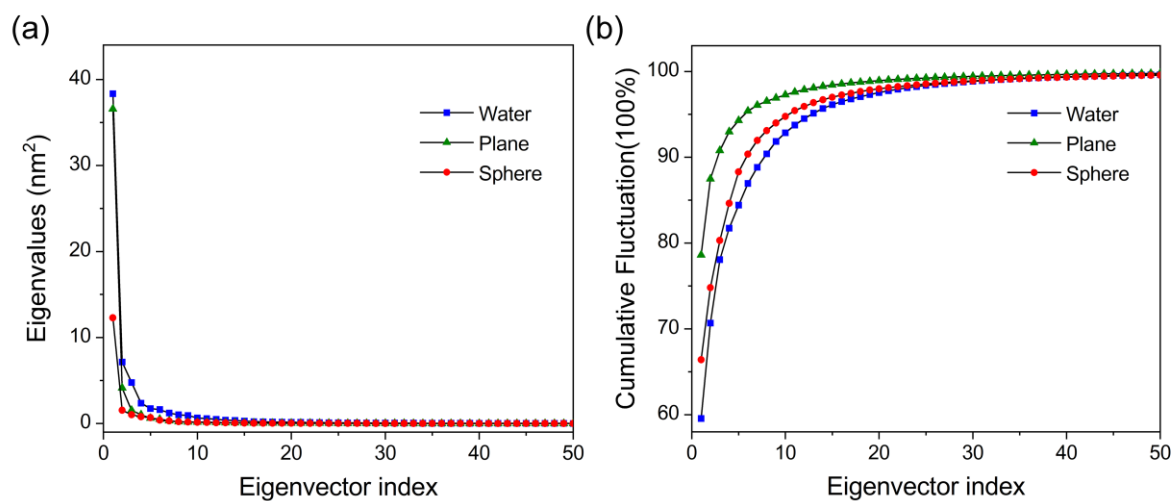
Atom type	$\sigma$ (nm)	$\epsilon$ (kJ/mol)	Charge(e)
CA	0.355	0.2929	0.00
HA	0.242	0.1255	0.06
CB	0.355	0.2929	-0.16
CU	0.350	0.3347	0.47
HU	0.250	0.2092	0.00
NU	0.325	0.7113	-0.62
CZ	0.350	0.2761	0.22
HZ	0.250	0.1255	0.03
CJ1	0.350	0.2761	-0.13
HJ1	0.250	0.1255	0.05
CJ2	0.350	0.2761	0.07
HJ2	0.250	0.1255	-0.02



**Figure S2.** The residue contact maps of Aβ<sub>42</sub> peptide adsorbed on CC3 surfaces (a) plane and (b) sphere.



**Figure S3.** RMSF of C<sub>α</sub> atoms of Aβ<sub>42</sub> peptide averaged over the last 40 ns trajectory.



**Figure S4.** The (a) eigenvalues (nm<sup>2</sup>) and (b) cumulative percentages (%) of the first 50 eigenvectors derived from PCA analysis of the Aβ<sub>42</sub> peptide backbone atoms in water and on the CC3 crystals versus eigenvector indices.

## References

1. M. J. Bojdys, T. Hasell, N. Severin, K. E. Jelfs, J. P. Rabe and A. I. Cooper, *Chem. Commun.*, 2012, **48**, 11948-11950.
2. X. Kong and J. Jiang, *Phys. Chem. Chem. Phys.*, 2017, **19**, 18178-18185.