

Supporting Information

Folding Dynamics of 3,4,3-LI(1,2-HOPO) in Its Free and Bound State with U⁴⁺ Implicated by MD Simulations

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Table S1. The number of clusters sampled in the simulations of *t*-HOPO, *t*-HOPO⁴⁻ and U^{IV}-*t*-HOPO complex systems, respectively, using three cutoff values. The proportions of main clusters in each system are also shown.

	<i>t</i> -HOPO			<i>t</i> -HOPO ⁴⁻			U ^{IV} - <i>t</i> -HOPO complex		
Cutoff (nm)	0.05	0.18	0.35	0.05	0.18	0.35	0.05	0.18	0.35
Num of clusters	4643	244	13	2394	44	4	26	1	1
The proportion of main clusters (%)									
Cluster1	0.40	5.50	59.08	2.82	59.74	97.68	51.10	100	100
Cluster2	0.32	4.08	17.36	1.70	6.98	1.62	23.98	/	/
Cluster3	0.22	3.18	10.84	1.28	5.80	0.64	16.30	/	/
Cluster4	0.20	2.96	5.02	1.16	4.00	0.06	3.90	/	/
Cluster5	0.16	2.84	4.04	1.16	3.52	/	1.34	/	/
Cluster6	0.16	2.60	1.22	0.92	2.66	/	0.86	/	/

Table S2. The compositions of the systems to simulate *t*-HOPO, *t*-HOPO⁴⁻ and U^{IV}-*t*-HOPO in aqueous phase.

Systems	N _{HOPO}	N _{U4+}	N _{Na}	N _{water}	Sampling time (ns)	Box size (nm ³)
<i>t</i> -HOPO	1	0	0	1559	100	3.844*3.678*3.391
<i>t</i> -HOPO ⁴⁻	1	0	4	1541	100	3.842*3.699*3.308
U ^{IV} - <i>t</i> -HOPO	1	1	0	1500	100	3.459*3.602*3.683

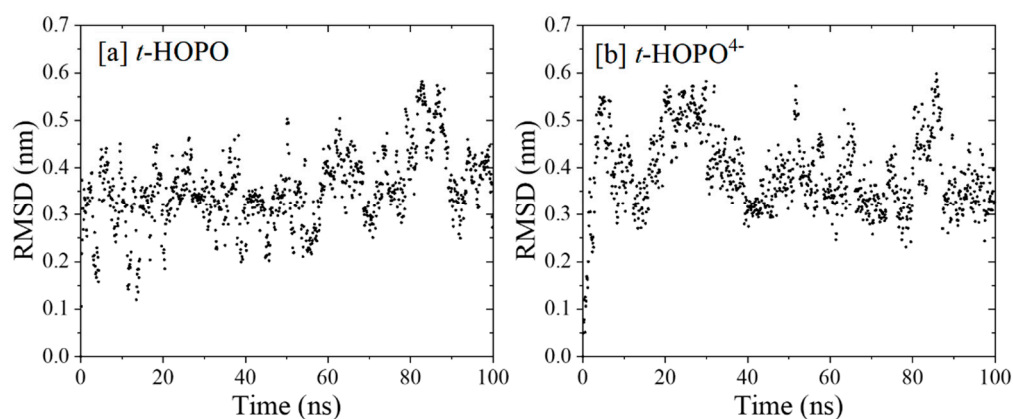


Figure S1. Atomic positional RMSD of all atoms of *t*-HOPO (a) and *t*-HOPO⁴⁻ (b). The structure of the first frame in each simulation was used as the reference.

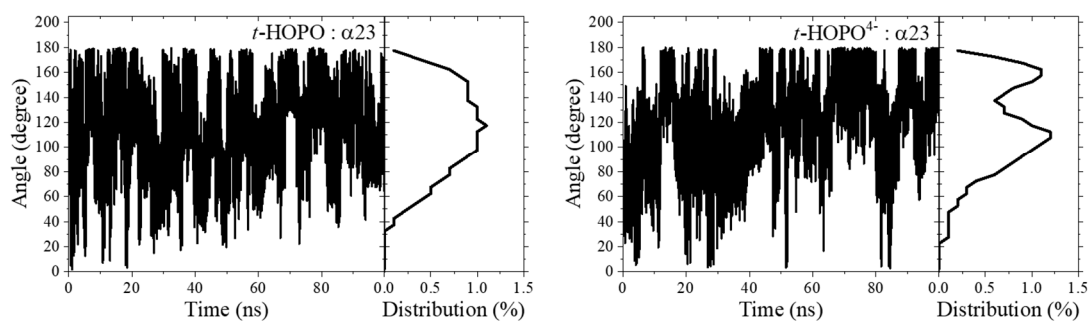


Figure S2. The time evolution of the angle between Plane2 and Plane3 (α_{23}) and their populations in the simulations of *t*-HOPO in its neutral (left) and deprotonated (right) states.

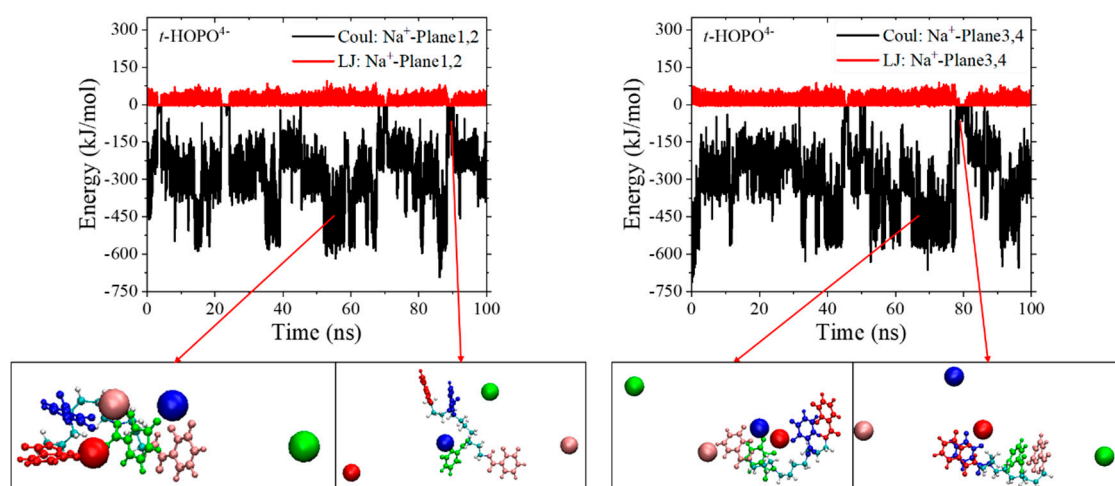


Figure S3. The time evolution of the interaction between Na⁺ counterion and the m-HOPO units in the simulations of *t*-HOPO⁴⁻ and the typical conformations in the simulation. Color scheme: Plane1 and Na1 in red, Plane2 and Na2 in blue, Plane3 and Na3 in green, Plane4 and Na4 in pink.

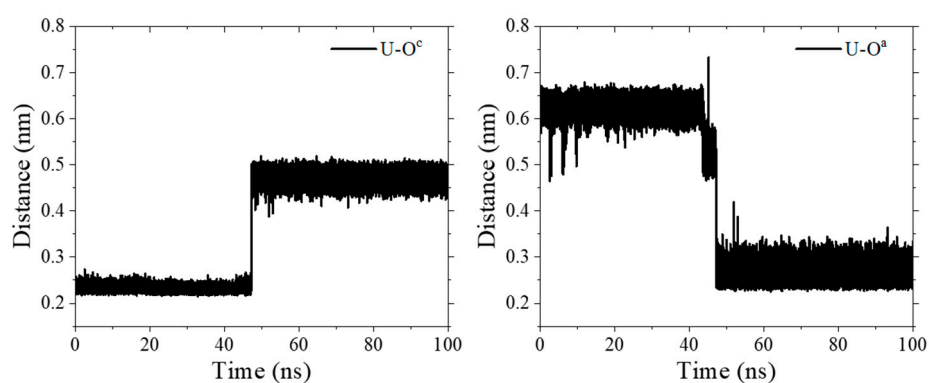


Figure S4. The time evolution of distances between U^{4+} and O^c , U^{4+} and O^a .

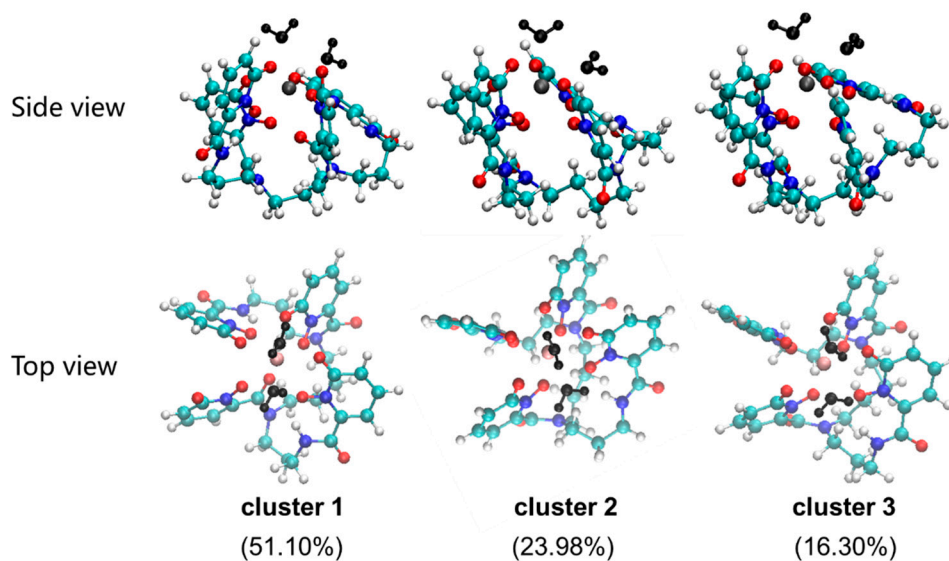


Figure S5. The central structures of the top three clusters in the simulations of U^{IV} -*t*-HOPO complex with side view (top) and top view (below). Color scheme: N atoms in blue, O atoms in red, water molecules in black, C atoms in cyan, H atoms in white, and U atom in brown.

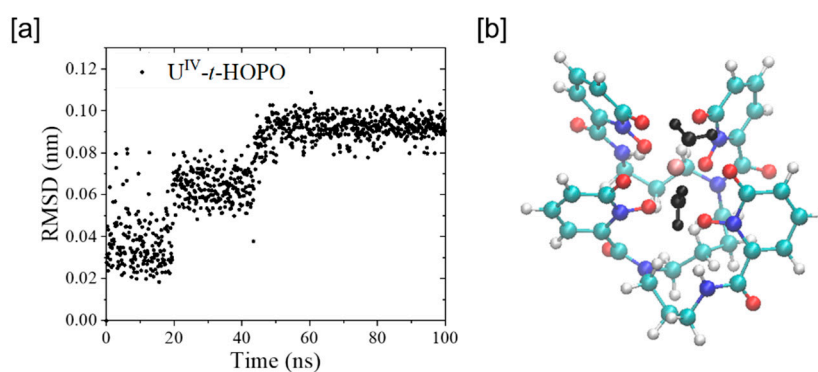


Figure S6. (a) Atomic positional RMSD of all atoms of U^{IV} -*t*-HOPO complex with the structure of first frame in the simulation used as the reference. (b) The structure of first frame in the simulation.