

Supporting Information

Host-Guest Interactions Induced Emission Enhancement of Amphiphilic AIEgens: A Computational Study

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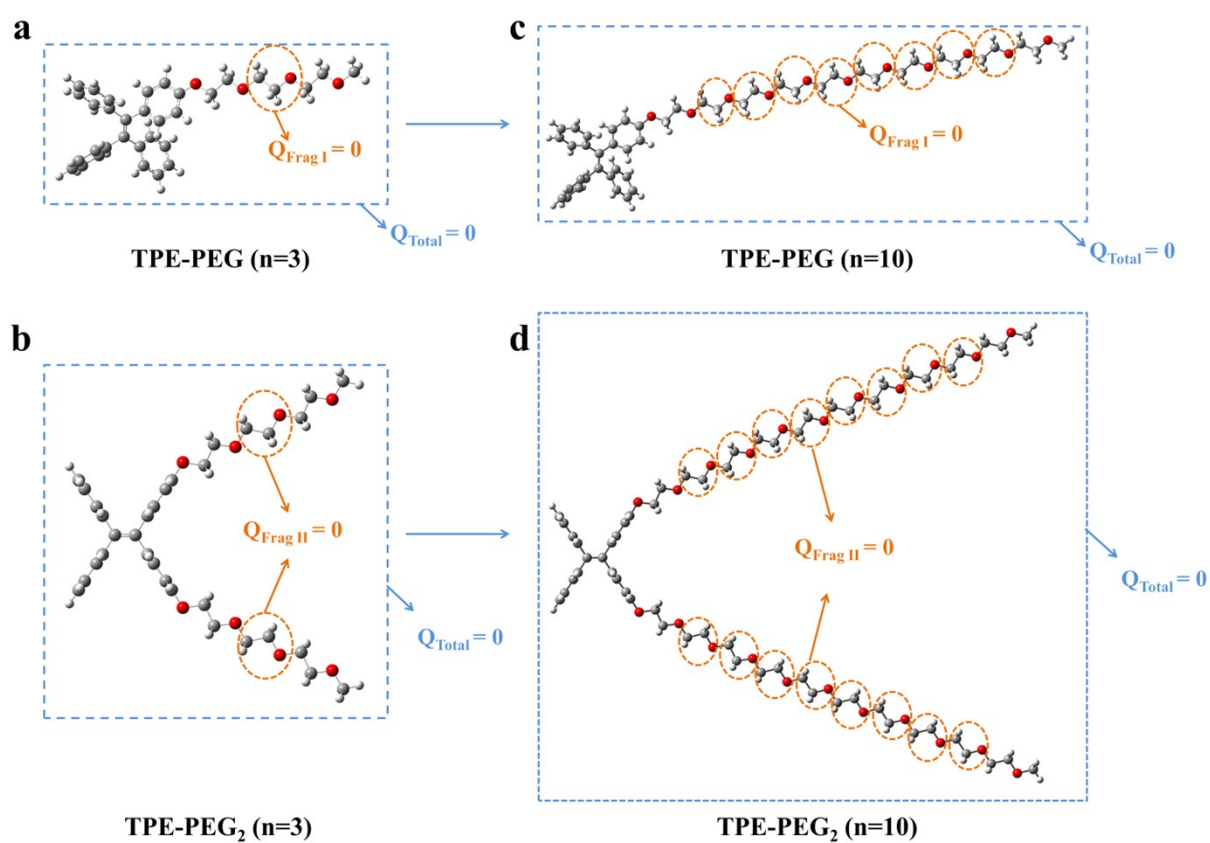


Fig. S1. The optimized structures of TPE-PEG and TPE-PEG₂ to obtain the partial charges of modeled guest molecules (G-1, G-2 and G-3) by using restrained electrostatic potential (RESP) fit method.

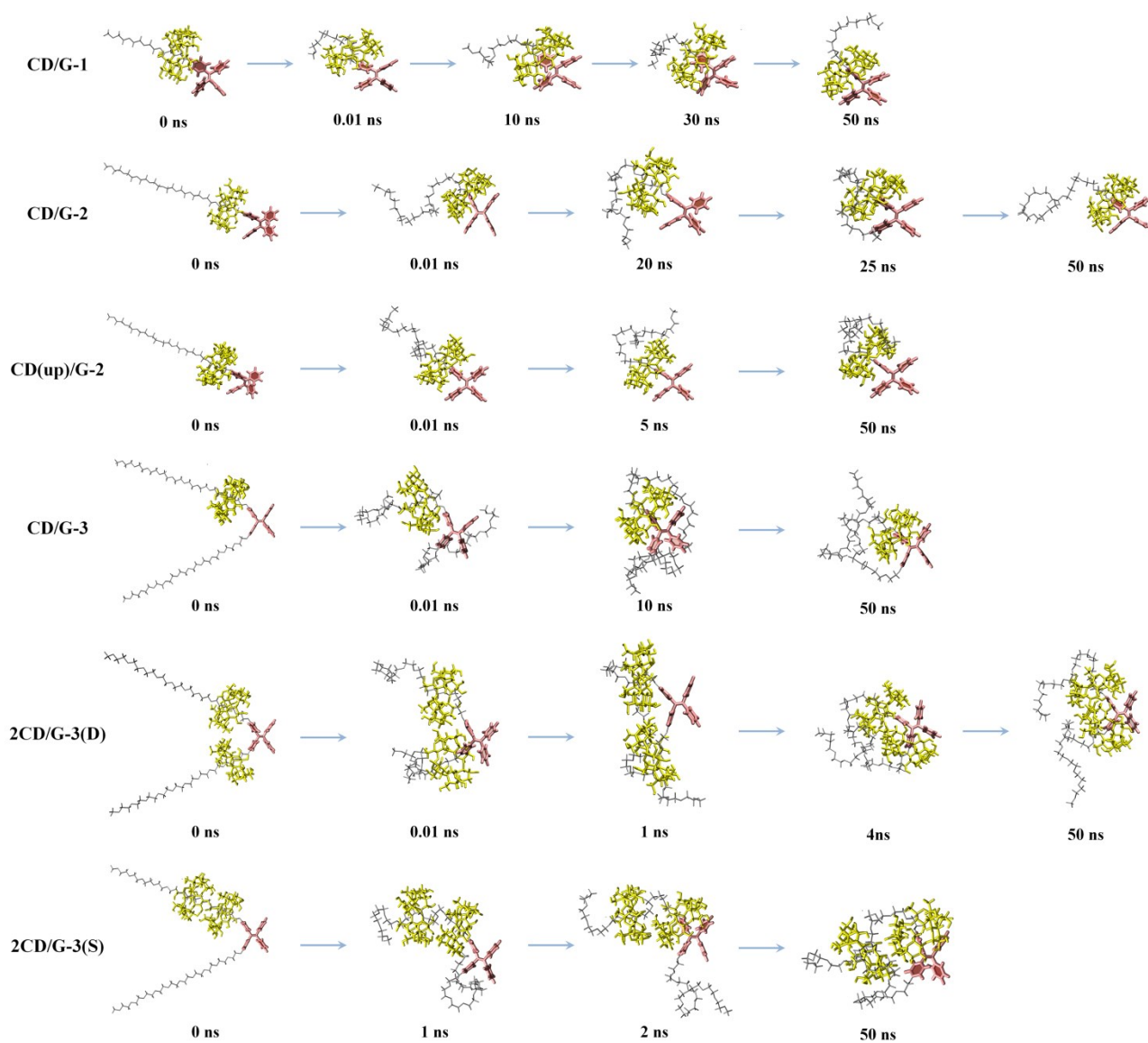


Fig. S2. Snapshots of the crucial phases in the progress of self-assembly between CD and guest molecules as a function of time during the MD simulations.

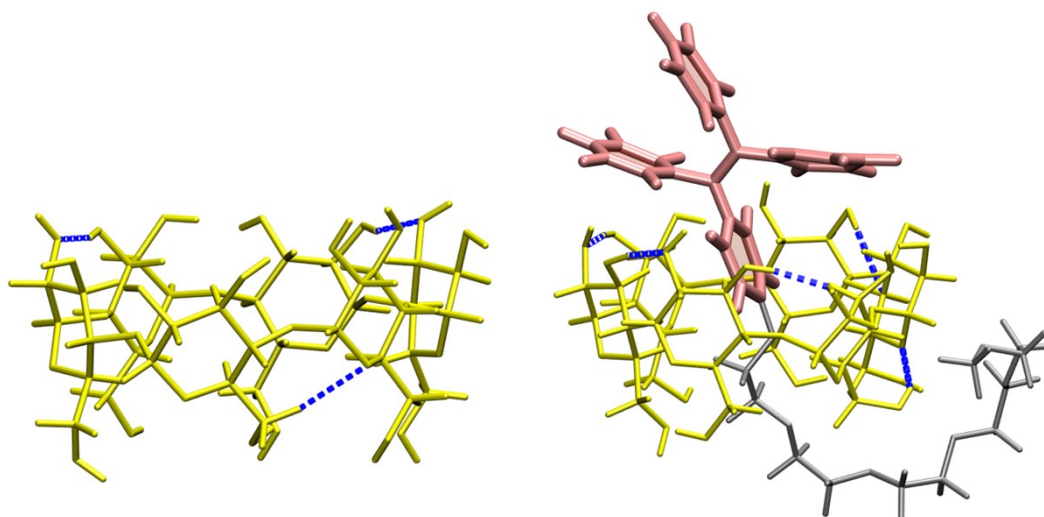


Fig. S3. The intramolecular H-bonds of free CD (left) and CD/G-1 host-guest inclusion (right), remarked by dashed blue line.

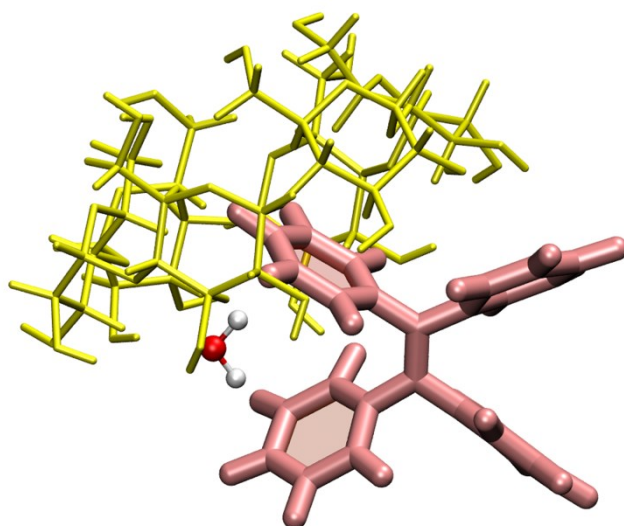


Fig. S4. Representative configuration of CD/TPE and one embedded water is shown in the cavity of CD.

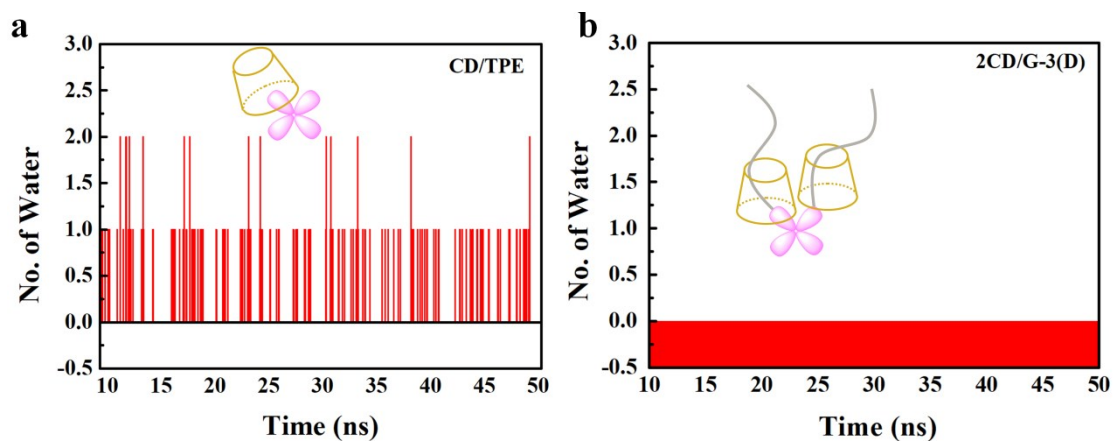


Fig. S5. Evolution of embedded water numbers in the cavity of CD in (a) CD/TPE inclusion and (b) 2CD/G-3(D) as a function of simulation time. CD/TPE is modelled system for comparison.

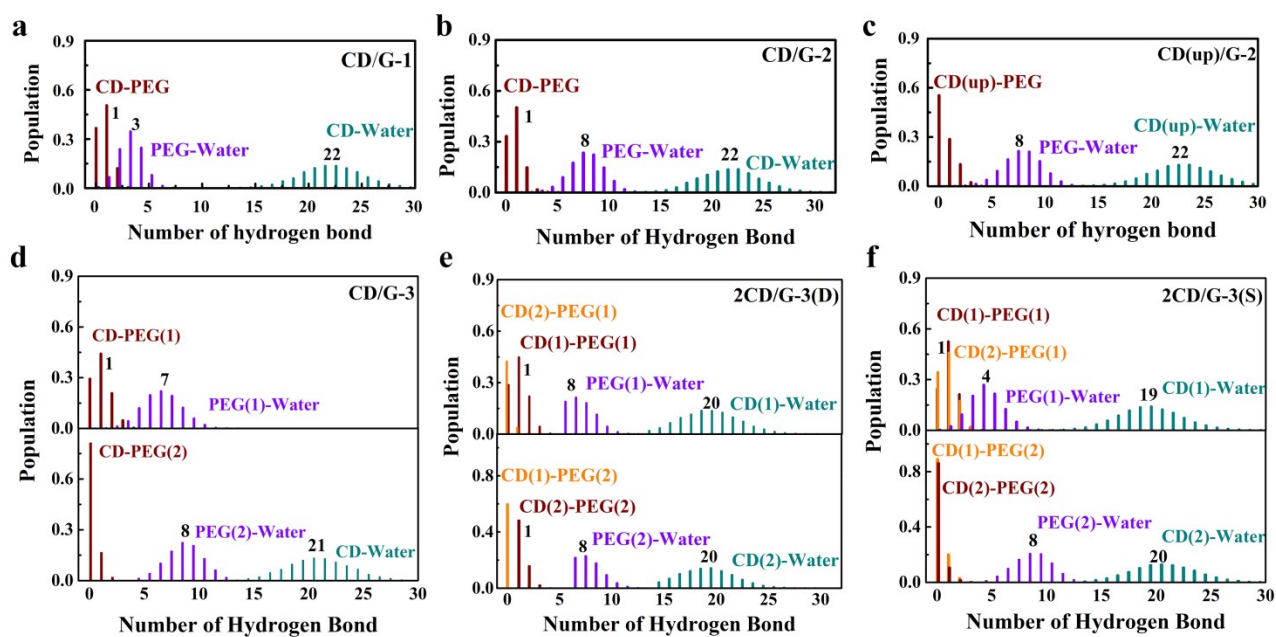


Fig. S6. Distributions of intermolecular H-bonds for six studied host-guest inclusions: (a) CD/G-1, (b) CD/G-2, (c) CD(up)/G-2, (d) CD/G-3, (e) 2CD/G-3(D) and (f) 2CD/G-3(S), respectively.

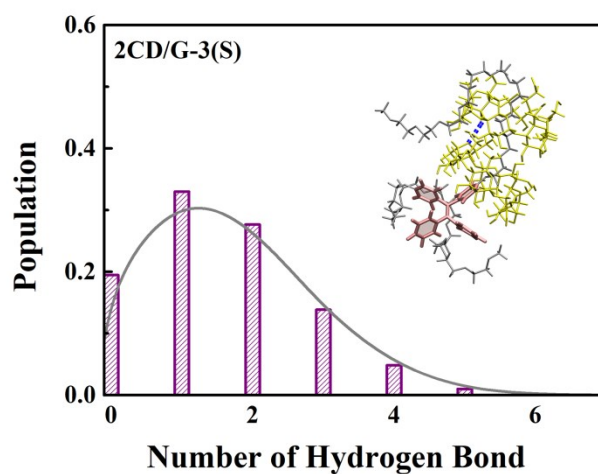
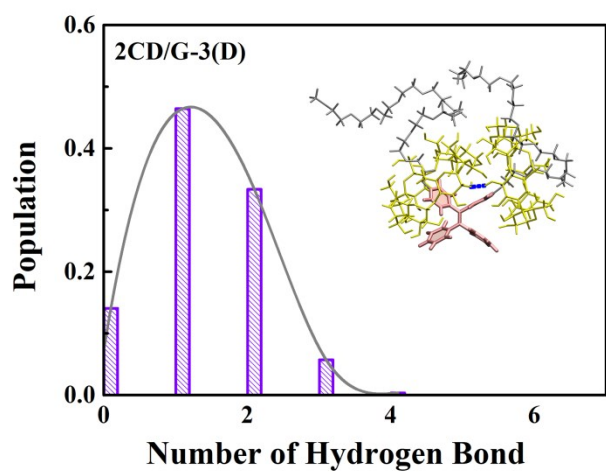


Fig. S7. The population of intermolecular H-bonds of neighboring two CDs in 2CD/G-3(D) (left) and 2CD/G-3(S) (right) inclusions. The representative snapshots of 2CD/G-3(D) and 2CD/G-3(S) were shown in the inset.

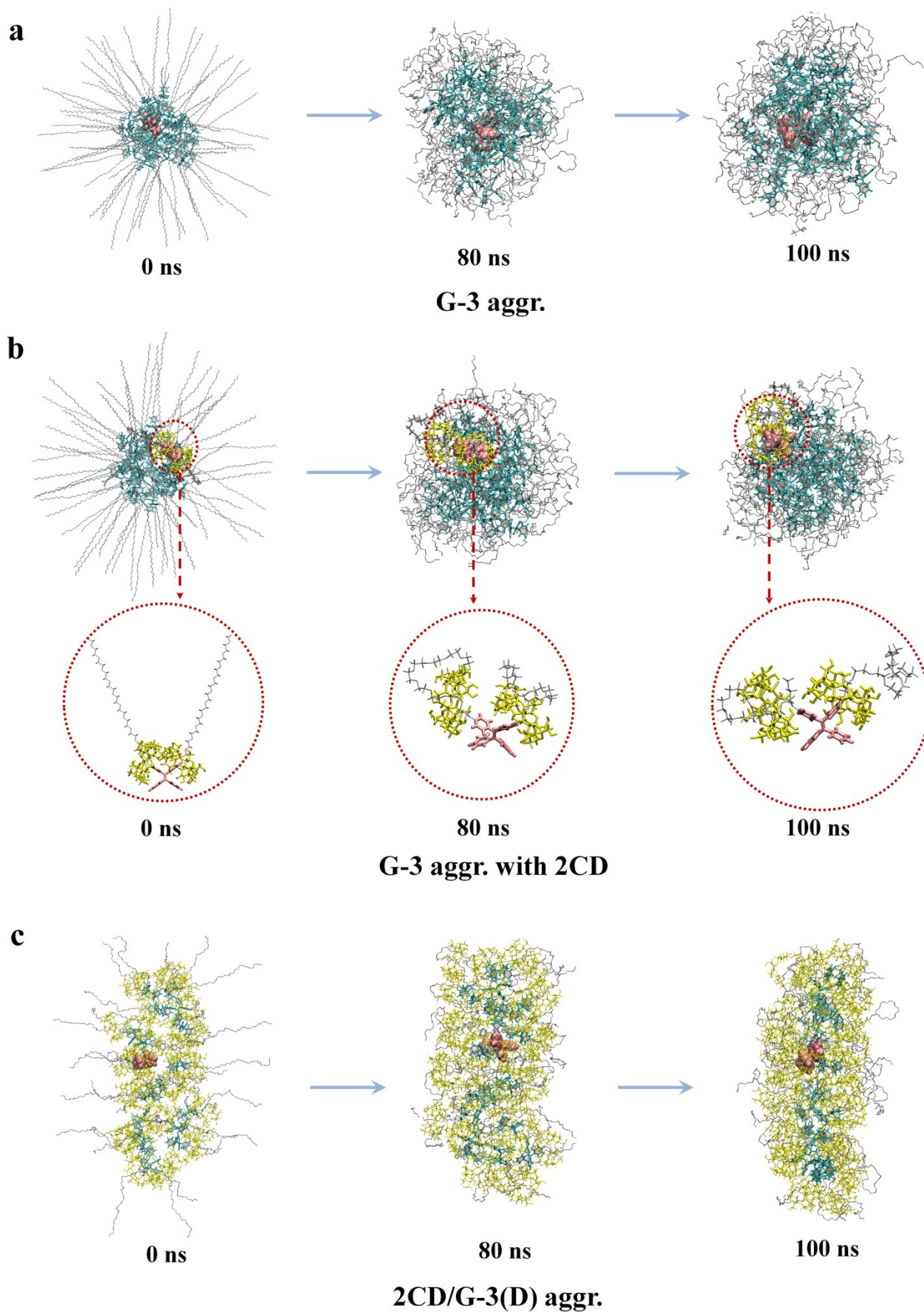


Fig. S8. Snapshots of the crucial phases in the progress of assembly of (a) G-3 aggregate, (b) G-3 aggregate with 2CD and (c) 2CD/G-3(D) aggregate as a function of time during the MD simulations.

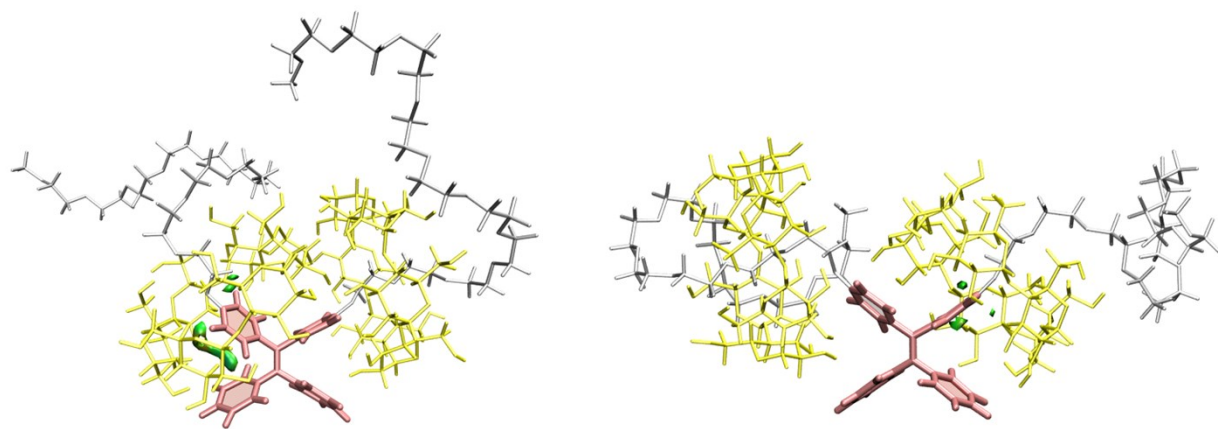


Fig. S9. The Independent Gradient Model (IGM) analysis for 2CD/G-3(D) (left) and one molecule with CD in G-3 aggregate with 2CD (right).

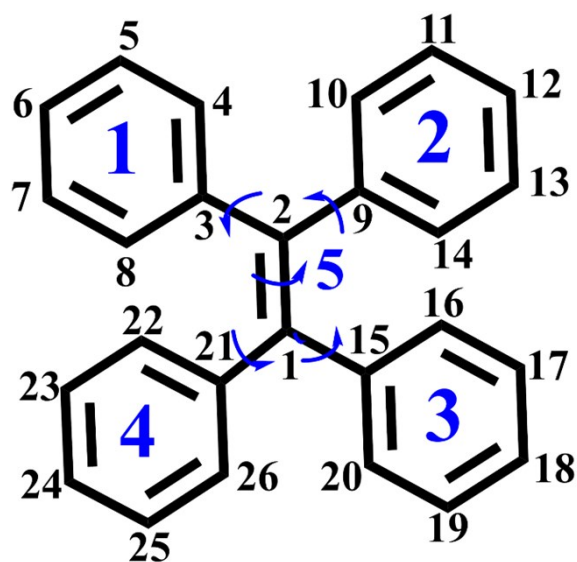


Chart S1. Chemical Structure of QM region of the representative monomers: G-2, G-3, 2CD/G-3(D), and assembled aggregates: G-3 aggregate, G-3 aggregate with 2CD and 2CD/G-3(D) aggregate.

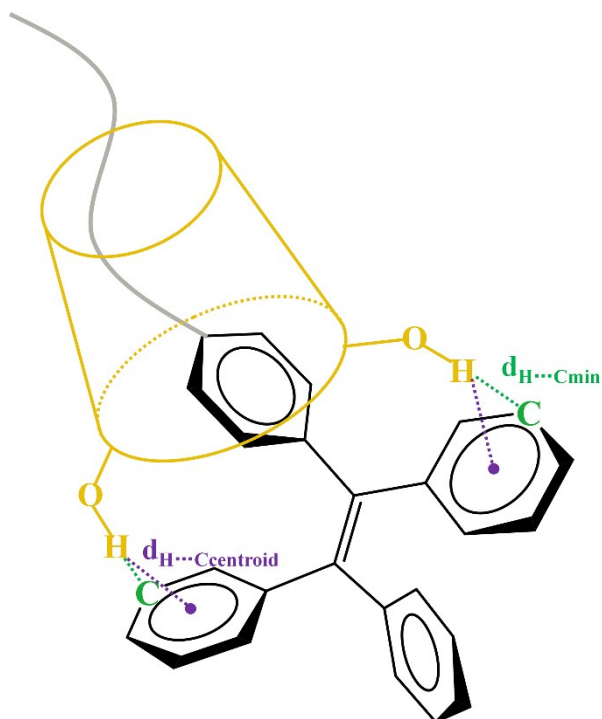


Chart S2. Definitions of distances measured in O-H... π interactions: $d_{H...C_{min}}$ = distance between the hydroxyl group of CD and the nearest aromatic carbon ($H_{hydroxyl}...C_{aromatic}$ distance) at 2- or 4-position of TPE moiety of guests in host-guest inclusions (dashed green line); $d_{H...C_{centroid}}$ = distance between the hydroxyl group of CD and the centroid of the phenyl ring of TPE moiety of guests in host-guest inclusions (dashed purple line).

Table S1. Distances $d_{H\cdots C_{min}}$ and $d_{H\cdots C_{centroid}}$ measured in O–H $\cdots\pi$ interactions at 2- and 4-position of TPE moiety for CD/G-1, CD/G-2, CD(up)/G-2, CD/G-3, 2CD/G-3(D) and 2CD/G-3(S), respectively. The cutoff distance for O–H $\cdots\pi$ interactions are $d_{H\cdots C_{min}} < 2.90$ Å and $d_{H\cdots C_{centroid}} < 3.60$ Å, respectively. Distance larger than cutoff distance is marked in orange.

		$d_{H\cdots C_{min}}(\text{\AA})$	$d_{H\cdots C_{centroid}}(\text{\AA})$
CD/G-1	2-position	2.187	2.760
	4-position	2.851	3.974
CD/G-2	2-position	2.433	2.502
	4-position	2.966	4.056
CD(up)/G-2	2-position	2.986	3.208
	4-position	3.972	4.735
CD/G-3	2-position	2.895	3.524
	4-position	2.080	2.876
2CD/G-3(D)	2-position	2.321	2.985
	4-position	2.548	3.266
2CD/G-3(S)	2-position	2.040	2.865
	4-position	2.511	3.049

Table S2. Calculated vertical excitation energy (VEE), electronic dipole moment (EDM), oscillator strength (f) and the assignment for S_1 of TPE and radiative decay rate constant (k_r).

S_1	VEE	EDM	f	assignment	k_r
G-2	2.02 eV (613 nm)	5.21 D	0.2081	HOMO $\rightarrow$ LUMO (98.0%)	3.70×10^7
G-3	2.06 eV (603 nm)	5.22 D	0.2124	HOMO $\rightarrow$ LUMO (97.8%)	3.90×10^7
2CD/G-3(D)	2.82 eV (439 nm)	4.64 D	0.2312	HOMO $\rightarrow$ LUMO (97.7%)	7.99×10^7
G-3 aggr.	2.73 eV (454 nm)	5.22 D	0.2824	HOMO $\rightarrow$ LUMO (99.4%)	9.14×10^7
G-3 aggr. with 2CD	3.50 eV (354 nm)	5.07 D	0.3409	HOMO $\rightarrow$ LUMO (98.0%)	1.81×10^8
2CD/G-3(D) aggr.	3.31 eV (375 nm)	4.19 D	0.2201	HOMO $\rightarrow$ LUMO (98.1%)	9.56×10^7

Table S3. The extracted key dihedral angles (in degree) of three monomers: G-2, G-3, 2CD/G-3(D) and three nanoaggregates: G-3 aggregate, G-3 aggregate with 2CD and 2CD/G-3(D) aggregate, respectively. S_0/S_1 and $|S_0-S_1|$ represent the geometric parameters extracted from the ground/excited states and the difference between them, respectively.

	G-2			G-3			2CD/G-3(D)			G-3 aggr.			G-3 aggr. with 2CD			2CD/G-3 aggr.		
	S_0	S_1	$ S_0-S_1 $	S_0	S_1	$ S_0-S_1 $	S_0	S_1	$ S_0-S_1 $	S_0	S_1	$ S_0-S_1 $	S_0	S_1	$ S_0-S_1 $	S_0	S_1	$ S_0-S_1 $
1-	134.8	153.2	18.3	136.0	152.8	16.8	102.8	104.0	1.2	125.3	148.1	22.8	77.8	62.5	15.3	82.6	62.7	19.9
2-	133.9	151.1	17.2	134.9	150.5	16.6	134.5	154.0	19.5	60.5	52.3	8.2	102.6	137.0	34.4	90.7	137.5	46.8
3-	47.5	23.2	24.3	47.5	24.0	23.5	60.7	26.1	34.6	127.7	140.9	13.2	78.9	89.7	10.8	69.4	25.5	43.9
4-	47.5	21.5	26.0	47.1	21.1	26.0	46.6	40.8	5.8	129.4	142.2	12.8	136.9	167.9	31.0	75.1	81.3	6.2
5-	170.2	129.0	41.2	168.5	130.0	38.5	3.6	20.5	16.9	175.2	145.5	29.7	178.8	179.9	1.1	2.6	28.2	25.6

Table S4. Selected bond lengths (Å), bond angle (degree), and dihedral angle (degree) of G-2 at S₀ (S₁) minimum and the difference between S₀ and S₁

structural parameters		G-2		
		S ₀	S ₁	S ₀ -S ₁
bond lengths				
	C ₁ -C ₂	1.354	1.465	0.111
	C ₂ -C ₃	1.489	1.440	0.049
	C ₂ -C ₉	1.491	1.447	0.044
	C ₁ -C ₁₅	1.491	1.445	0.046
	C ₁ -C ₂₁	1.491	1.444	0.047
bond angle				
	C ₁ -C ₂ -C ₃	122.7	118.8	3.8
	C ₁ -C ₂ -C ₉	122.4	118.4	4.0
	C ₂ -C ₁ -C ₁₅	122.5	118.4	4.1
	C ₂ -C ₁ -C ₂₁	122.6	118.8	3.8
dihedral angle				
1-	C ₁ -C ₂ -C ₃ -C ₄	134.8	153.2	18.3
2-	C ₁ -C ₂ -C ₉ -C ₁₀	133.9	151.1	17.2
3-	C ₂ -C ₁ -C ₁₅ -C ₁₆	47.5	23.2	24.3
4-	C ₂ -C ₁ -C ₂₁ -C ₂₂	47.5	21.5	26.0
5-	C ₃ -C ₂ -C ₁ -C ₁₅	170.2	129.0	41.2

Table S5. Selected bond lengths (Å), bond angle (degree), and dihedral angle (degree) of G-3 at S₀ (S₁) minimum and the difference between S₀ and S₁

structural parameters		G-3		
		S ₀	S ₁	S ₀ -S ₁
bond lengths				
	C ₁ -C ₂	1.355	1.465	0.110
	C ₂ -C ₃	1.489	1.441	0.048
	C ₂ -C ₉	1.489	1.446	0.043
	C ₁ -C ₁₅	1.491	1.446	0.045
	C ₁ -C ₂₁	1.490	1.443	0.047
bond angle				
	C ₁ -C ₂ -C ₃	122.5	119.1	3.4
	C ₁ -C ₂ -C ₉	122.4	118.4	4.0
	C ₂ -C ₁ -C ₁₅	122.6	118.4	4.2
	C ₂ -C ₁ -C ₂₁	122.5	118.9	3.6
dihedral angle				
1-	C ₁ -C ₂ -C ₃ -C ₄	136.0	152.8	16.8
2-	C ₁ -C ₂ -C ₉ -C ₁₀	134.9	150.5	16.6
3-	C ₂ -C ₁ -C ₁₅ -C ₁₆	47.5	24.0	23.5
4-	C ₂ -C ₁ -C ₂₁ -C ₂₂	47.1	21.1	26.0
5-	C ₃ -C ₂ -C ₁ -C ₁₅	168.5	130.0	38.5

Table S6. Selected bond lengths (Å), bond angle (degree), and dihedral angle (degree) of 2CD/G-3(D) at S₀ (S₁) minimum and the difference between S₀ and S₁

structural parameters		2CD/G-3(D)		
		S ₀	S ₁	S ₀ -S ₁
bond lengths				
	C ₁ -C ₂	1.346	1.437	0.091
	C ₂ -C ₃	1.491	1.433	0.059
	C ₂ -C ₉	1.490	1.467	0.023
	C ₁ -C ₁₅	1.489	1.419	0.070
	C ₁ -C ₂₁	1.489	1.481	0.008
bond angle				
	C ₁ -C ₂ -C ₃	124.2	124.6	0.4
	C ₁ -C ₂ -C ₉	123.3	122.0	1.3
	C ₂ -C ₁ -C ₁₅	121.1	120.4	0.7
	C ₂ -C ₁ -C ₂₁	122.9	120.8	2.1
dihedral angle				
1-	C ₁ -C ₂ -C ₃ -C ₄	102.8	104.0	1.2
2-	C ₁ -C ₂ -C ₉ -C ₁₀	134.5	154.0	19.5
3-	C ₂ -C ₁ -C ₁₅ -C ₁₆	60.7	26.1	34.6
4-	C ₂ -C ₁ -C ₂₁ -C ₂₂	46.6	40.8	5.8
5-	C ₃ -C ₂ -C ₁ -C ₁₅	3.6	20.5	16.9

Table S7. Selected bond lengths (Å), bond angle (degree), and dihedral angle (degree) of G-3 aggregate at S₀ (S₁) minimum and the difference between S₀ and S₁

structural parameters		G-3 aggr.		
		S ₀	S ₁	S ₀ -S ₁
bond lengths				
	C ₁ -C ₂	1.348	1.445	0.097
	C ₂ -C ₃	1.493	1.435	0.058
	C ₂ -C ₉	1.496	1.468	0.028
	C ₁ -C ₁₅	1.494	1.455	0.038
	C ₁ -C ₂₁	1.493	1.440	0.053
bond angle				
	C ₁ -C ₂ -C ₃	122.1	120.8	1.3
	C ₁ -C ₂ -C ₉	121.0	119.4	1.6
	C ₂ -C ₁ -C ₁₅	121.8	119.6	2.2
	C ₂ -C ₁ -C ₂₁	123.0	120.1	2.9
dihedral angle				
1-	C ₁ -C ₂ -C ₃ -C ₄	125.6	148.1	22.5
2-	C ₁ -C ₂ -C ₉ -C ₁₀	60.2	52.3	7.9
3-	C ₂ -C ₁ -C ₁₅ -C ₁₆	127.7	140.9	13.2
4-	C ₂ -C ₁ -C ₂₁ -C ₂₂	129.5	142.2	12.7
5-	C ₃ -C ₂ -C ₁ -C ₁₅	174.9	145.5	29.4

Table S8. Selected bond lengths (Å), bond angle (degree), and dihedral angle (degree) of G-3 aggregate with 2CD at S₀ (S₁) minimum and the difference between S₀ and S₁

structural parameters		G-3 aggr. with 2CD		
		S ₀	S ₁	S ₀ -S ₁
bond lengths				
	C ₁ -C ₂	1.341	1.430	0.089
	C ₂ -C ₃	1.495	1.432	0.063
	C ₂ -C ₉	1.499	1.482	0.017
	C ₁ -C ₁₅	1.496	1.491	0.005
	C ₁ -C ₂₁	1.493	1.425	0.068
bond angle				
	C ₁ -C ₂ -C ₃	124.4	121.7	2.7
	C ₁ -C ₂ -C ₉	122.5	121.1	1.4
	C ₂ -C ₁ -C ₁₅	121.6	119.5	2.1
	C ₂ -C ₁ -C ₂₁	125.7	123.1	2.6
dihedral angle				
1-	C ₁ -C ₂ -C ₃ -C ₄	77.8	62.5	15.3
2-	C ₁ -C ₂ -C ₉ -C ₁₀	102.6	137.0	34.4
3-	C ₂ -C ₁ -C ₁₅ -C ₁₆	78.9	89.7	10.8
4-	C ₂ -C ₁ -C ₂₁ -C ₂₂	136.9	167.9	31.0
5-	C ₃ -C ₂ -C ₁ -C ₁₅	178.8	179.9	1.1

Table S9. Selected bond lengths (Å), bond angle (degree), and dihedral angle (degree) of 2CD/G-3(D) aggregate at S₀ (S₁) minimum and the difference between S₀ and S₁

structural parameters		2CD/G-3(D) aggr.		
		S ₀	S ₀	S ₀ -S ₁
bond lengths				
	C ₁ -C ₂	1.339	1.436	0.097
	C ₂ -C ₃	1.486	1.424	0.062
	C ₂ -C ₉	1.492	1.485	0.007
	C ₁ -C ₁₅	1.500	1.437	0.064
	C ₁ -C ₂₁	1.487	1.479	0.008
bond angle				
	C ₁ -C ₂ -C ₃	121.5	117.3	4.2
	C ₁ -C ₂ -C ₉	122.3	122.0	0.3
	C ₂ -C ₁ -C ₁₅	118.4	116.5	1.9
	C ₂ -C ₁ -C ₂₁	126.0	126.9	0.9
dihedral angle				
1-	C ₁ -C ₂ -C ₃ -C ₄	82.6	62.7	19.9
2-	C ₁ -C ₂ -C ₉ -C ₁₀	90.7	137.5	46.8
3-	C ₂ -C ₁ -C ₁₅ -C ₁₆	69.4	25.5	43.9
4-	C ₂ -C ₁ -C ₂₁ -C ₂₂	75.1	81.3	6.2
5-	C ₃ -C ₂ -C ₁ -C ₁₅	2.6	28.2	25.5

Table S10. Reorganization Energy of each configuration is showed by AP.

	λ_{gs} (eV)	λ_{es} (eV)	λ (eV)
G-2	1.06	0.92	1.98
G-3	1.02	0.89	1.91
2CD/G-3(D)	0.68	0.88	1.56
G-3 aggr.	0.63	0.98	1.61
G-3 aggr. with 2CD	0.48	0.99	1.47
2CD/G-3(D) aggr.	0.54	1.16	1.70