

Supplementary Material

Unveiling the Synergy: A Combined Experimental and Theoretical Study of β -Cyclodextrin with Melatonin

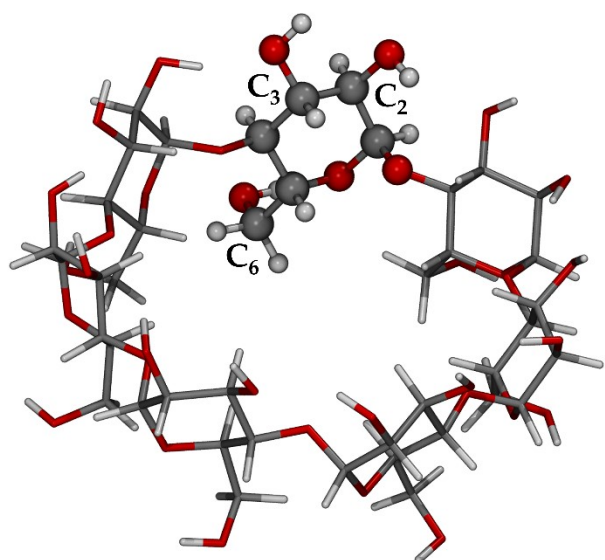
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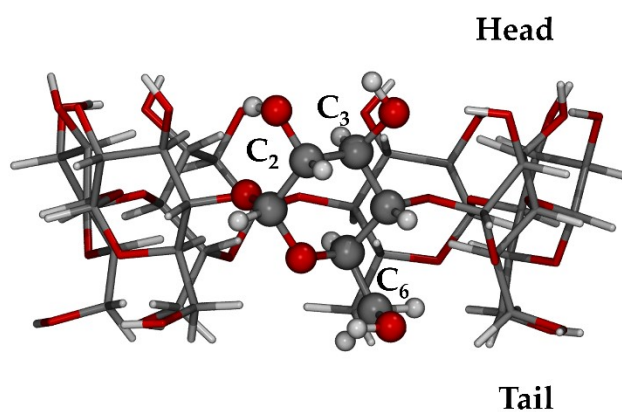
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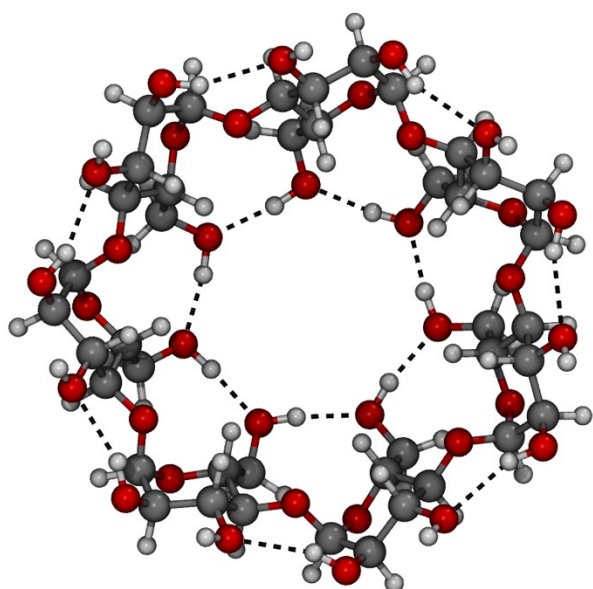
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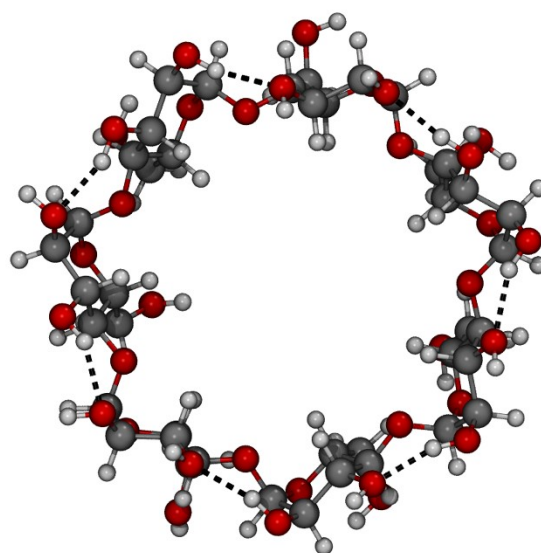
a) β -CD – top view



b) β -CD – side view



c) β -CD1 – top view



d) β -CD2 – top view

Figure S1: Top and side views of β -cyclodextrin (a, b) and open and closed conformations (c, d).

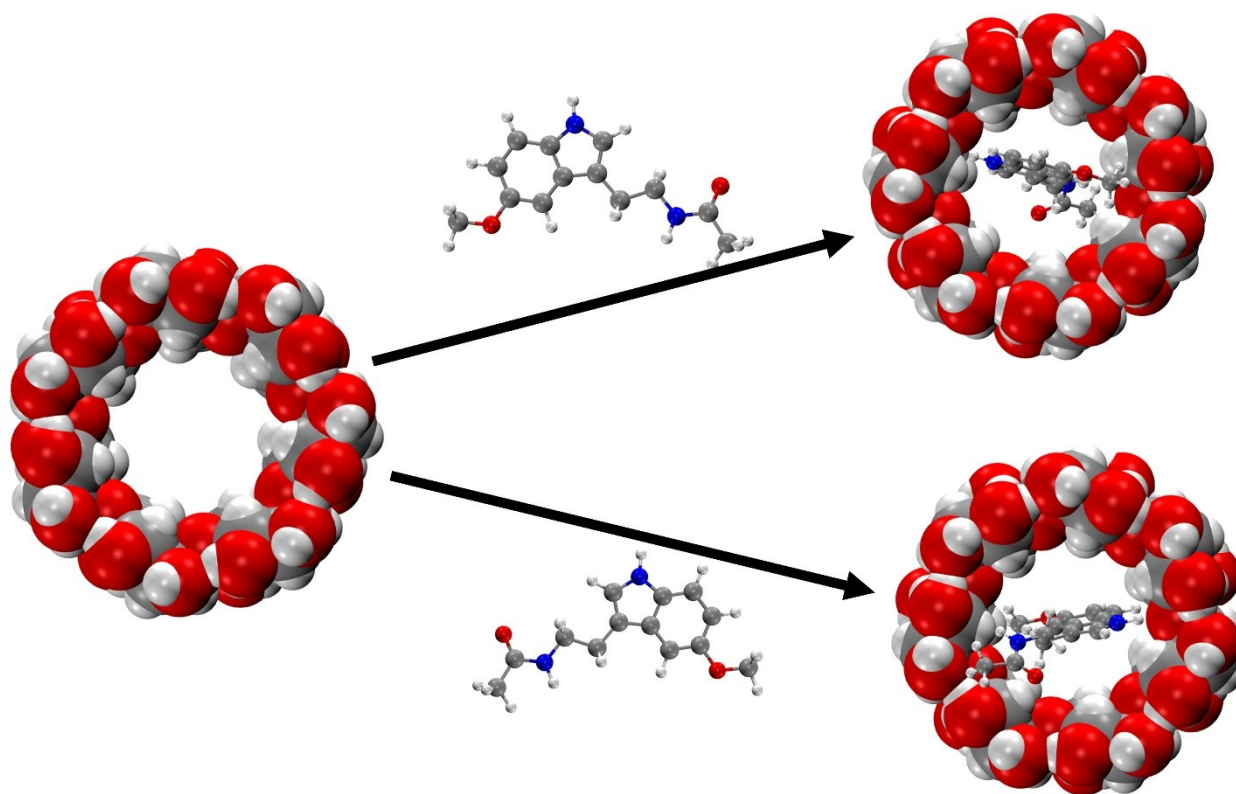


Figure S2: Starting complexes for the conformational search procedure. C in grey, N in blue, O in red, H in white.

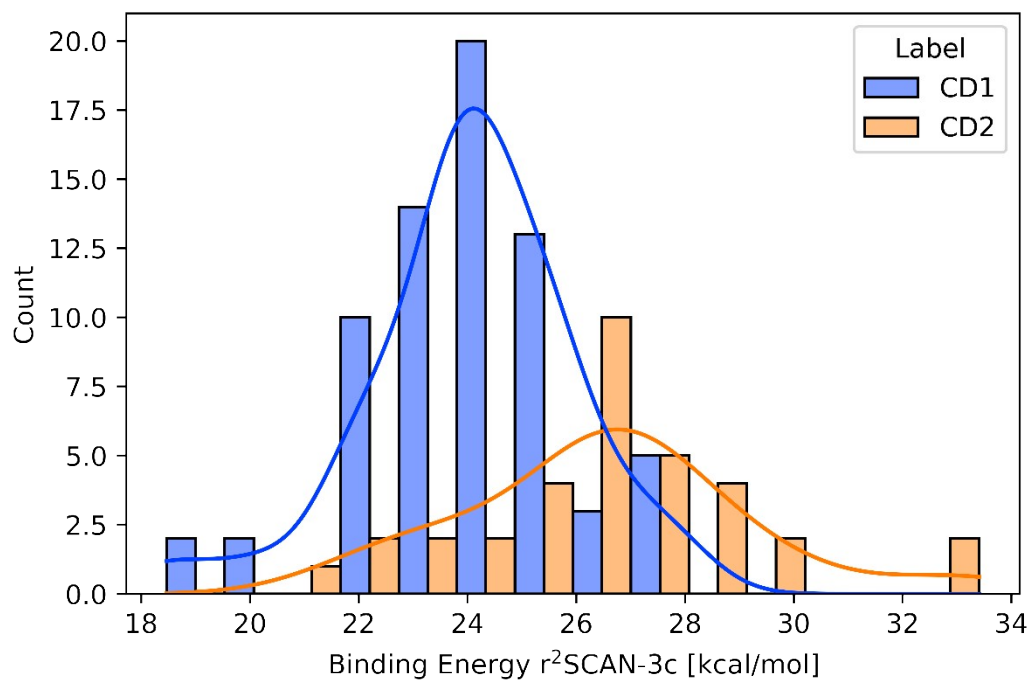


Figure S3: r^2 SCAN-3c binding energy (kcal/mol) distribution of MT/ β -CD complex in gas phase, in which the contributions given by CD1 and CD2 are separated.

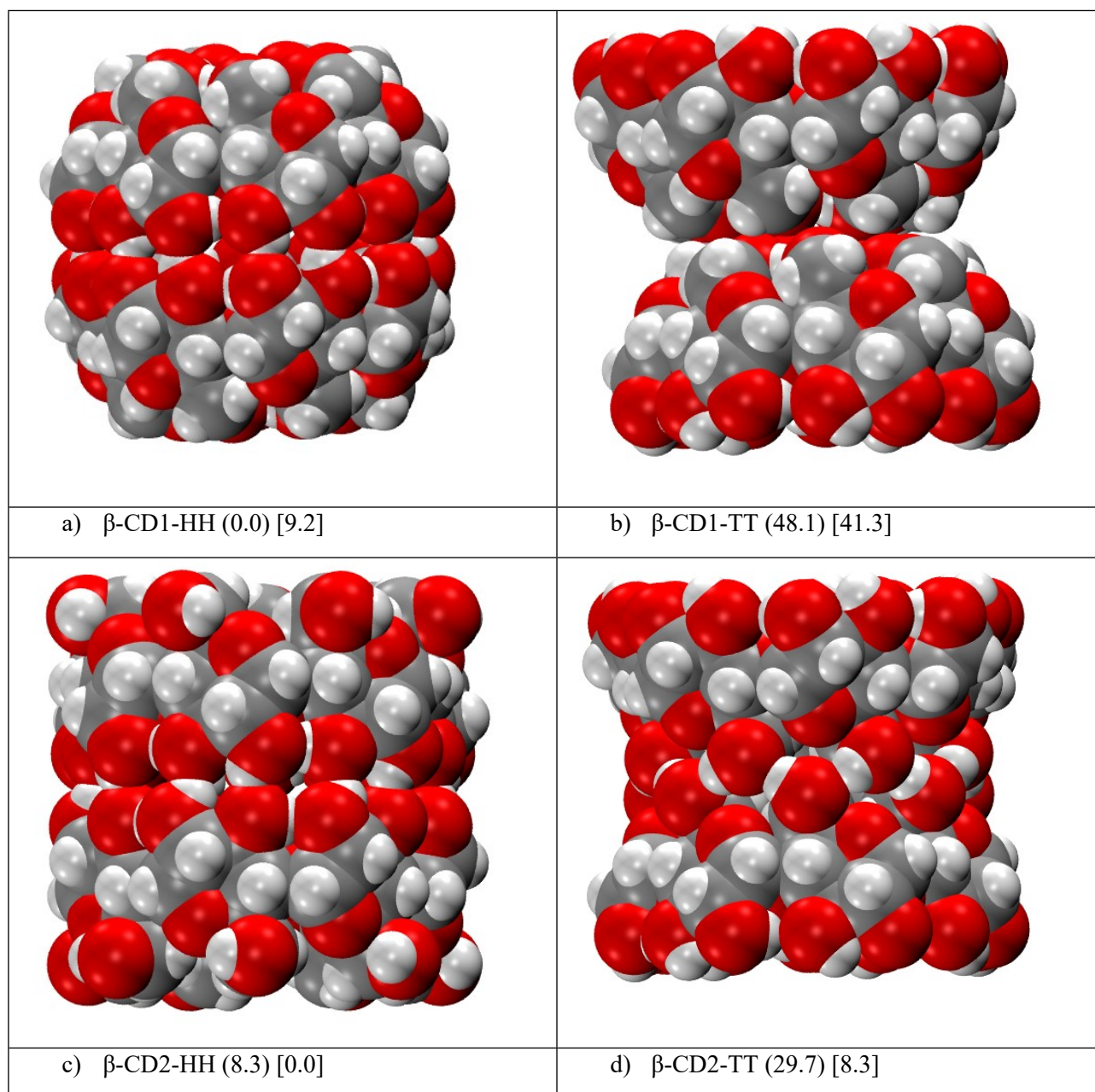


Figure S4: r²SCAN-3c optimized structures of the β -CD dimers in the gas phase. Numbers in round parenthesis correspond to the relative energy in the gas phase, in square parenthesis to the relative energy in water C-PCM (in kcal/mol). H and T stand for Head and Tail sides. Color code: H in white, C in grey, O in red.

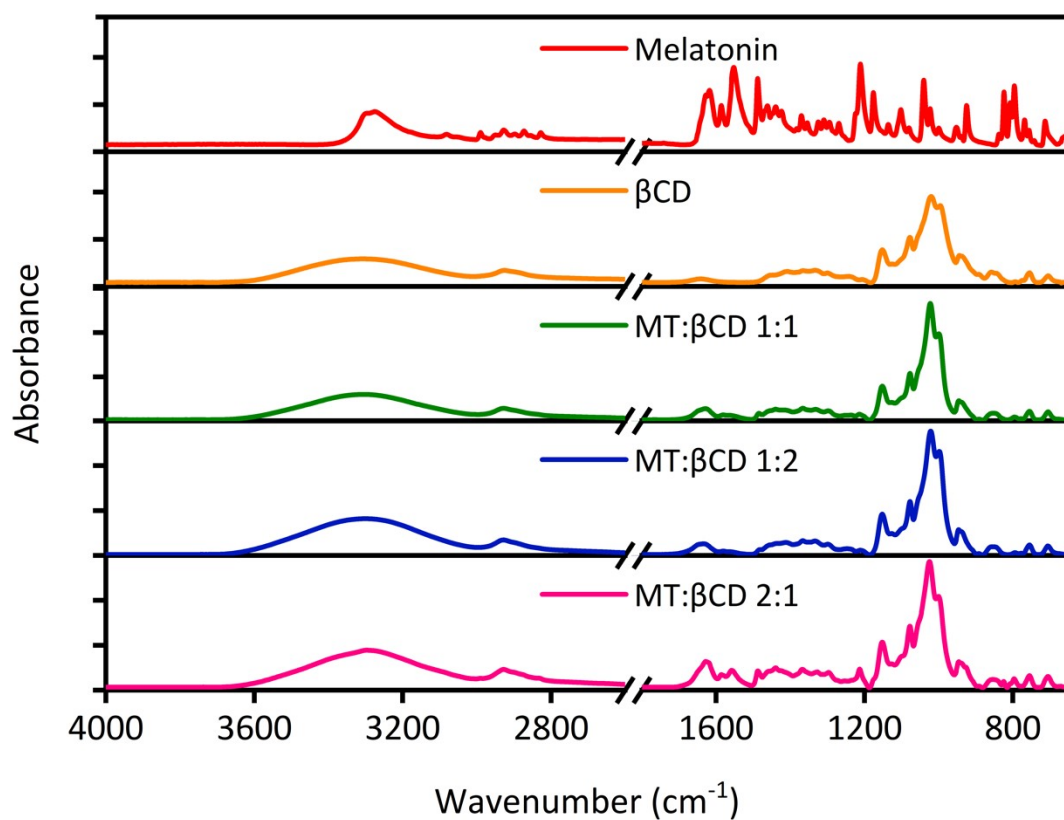


Figure S5: Experimental FTIR spectra of pure molecules (MT in red, β -CD in orange) and of the inclusion complexes in the three molar ratio proposed.

Gas				Water			
GFN2		r ² SCAN-3c		GFN2		r ² SCAN-3c	
Structures	ΔE	Structures	ΔE	Structures	ΔE	Structures	ΔE
x10	0.00	x00	0.00	x06	0.00	x04	0.00
x11	0.00	x01	0.83	x05	0.10	x00	0.34
x07	0.01	x03	1.40	x15	0.10	x03	0.65
x04	0.06	x02	1.73	x00	0.81	x01	0.68
x15	0.07	x04	2.09	x11	0.89	x11	1.06
x05	0.08	x15	2.36	x10	0.90	x07	1.07
x00	0.12	x05	2.37	x07	0.90	x10	1.08
x06	0.17	x10	2.70	x04	0.90	x02	1.27
x09	0.65	x11	2.70	x09	1.04	x09	1.72
x02	0.93	x07	2.71	x01	1.19	x06	1.77
x18	0.98	x09	2.88	x02	1.26	x05	2.38
x21	0.98	x06	2.90	x18	1.50	x15	2.40
x22	1.01	x18	3.61	x22	1.51	x18	2.57
x01	1.15	x21	3.63	x21	1.53	x21	2.61
x03	1.53	x22	3.63	x03	1.80	x22	2.63

Table S1: GFN2 and r²SCAN-3c energetic ranking of melatonin conformers in gas and implicit water phases.