

Supporting Information

Cocrystals of nutraceutical *p*-coumaric acid with caffeine and theophylline: polymorphism and solid-state stability explored in detail using their crystal graphs

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XRPD of cocrystals 1 – 4	3-6
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Figure 1. XRPD of *p*-coumaric acid:caffeine cocrystal, **1**

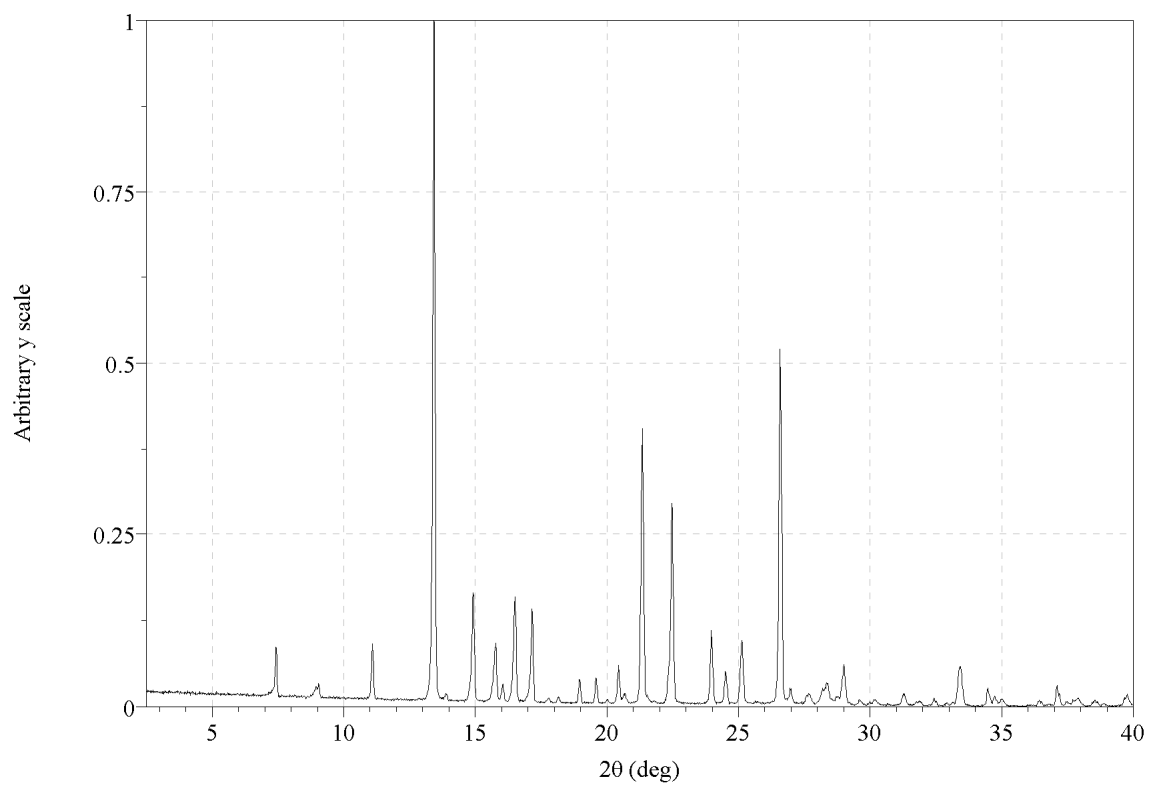


Figure 2. XRPD of *p*-coumaric acid:caffeine cocrystal, **2**

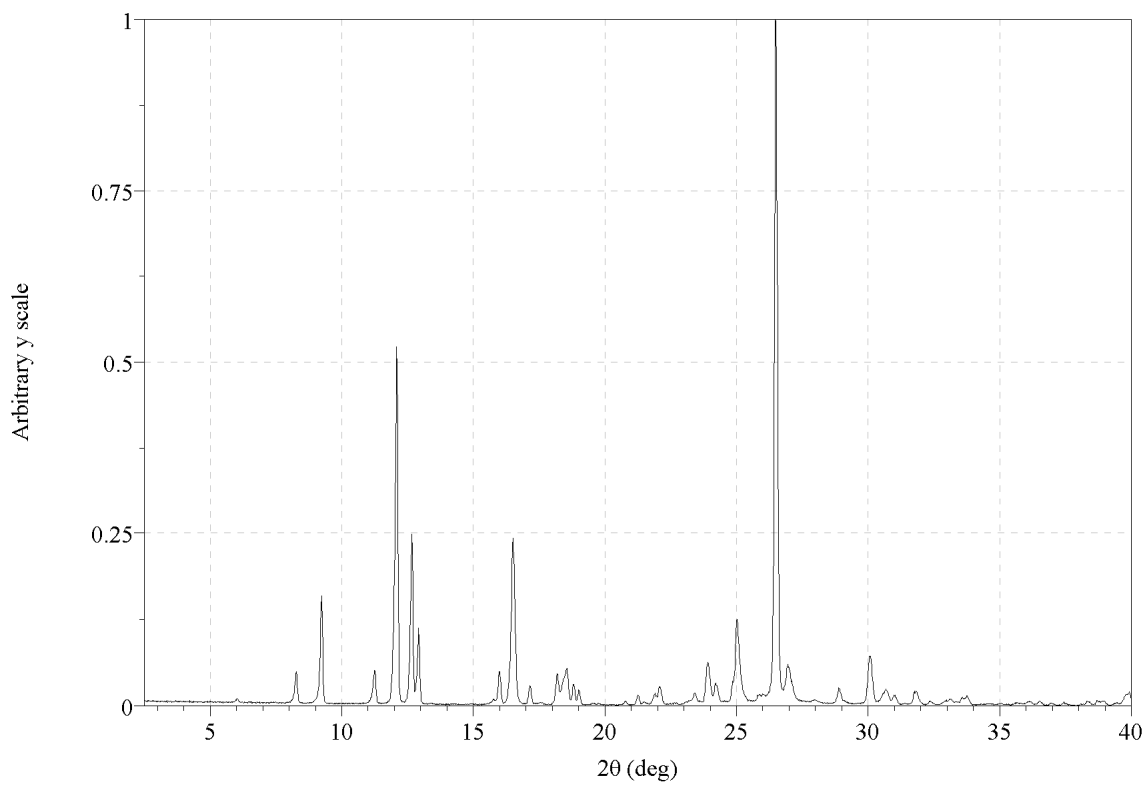


Figure 3. XRPD of *p*-coumaric acid:theophylline cocrystal, **3** (Form I)

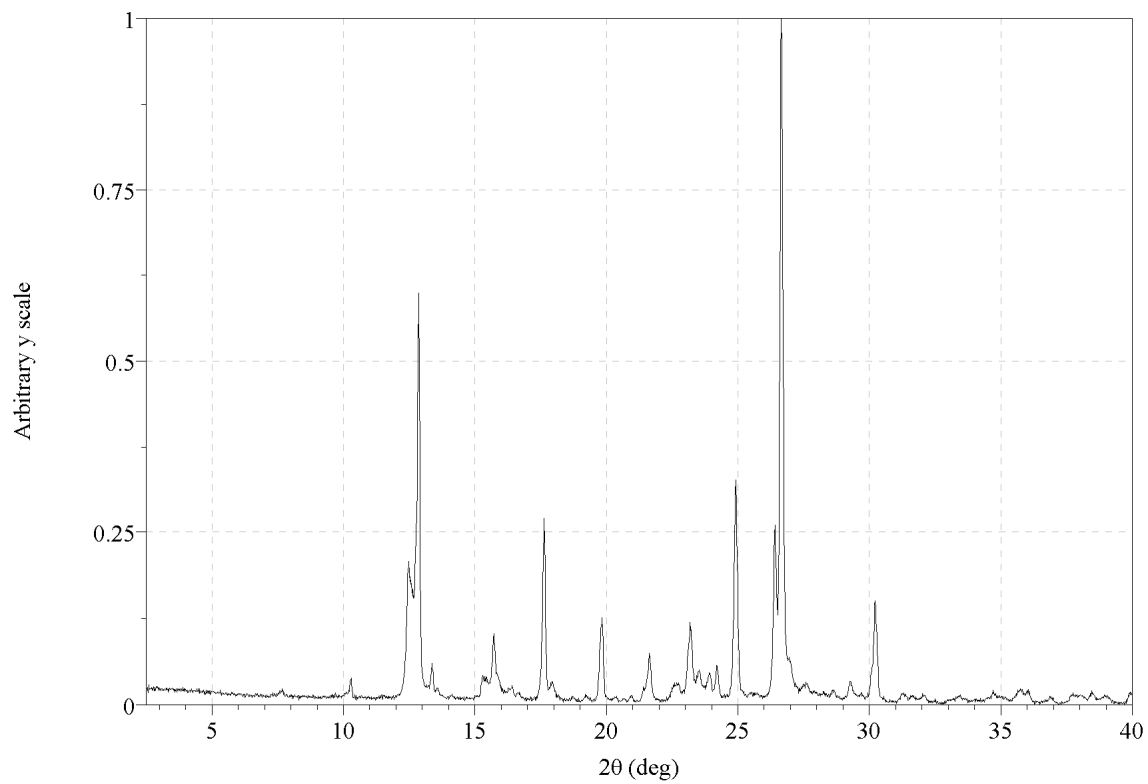


Figure 4. XRPD of *p*-coumaric acid:theophylline cocrystal, **4** (Form II)

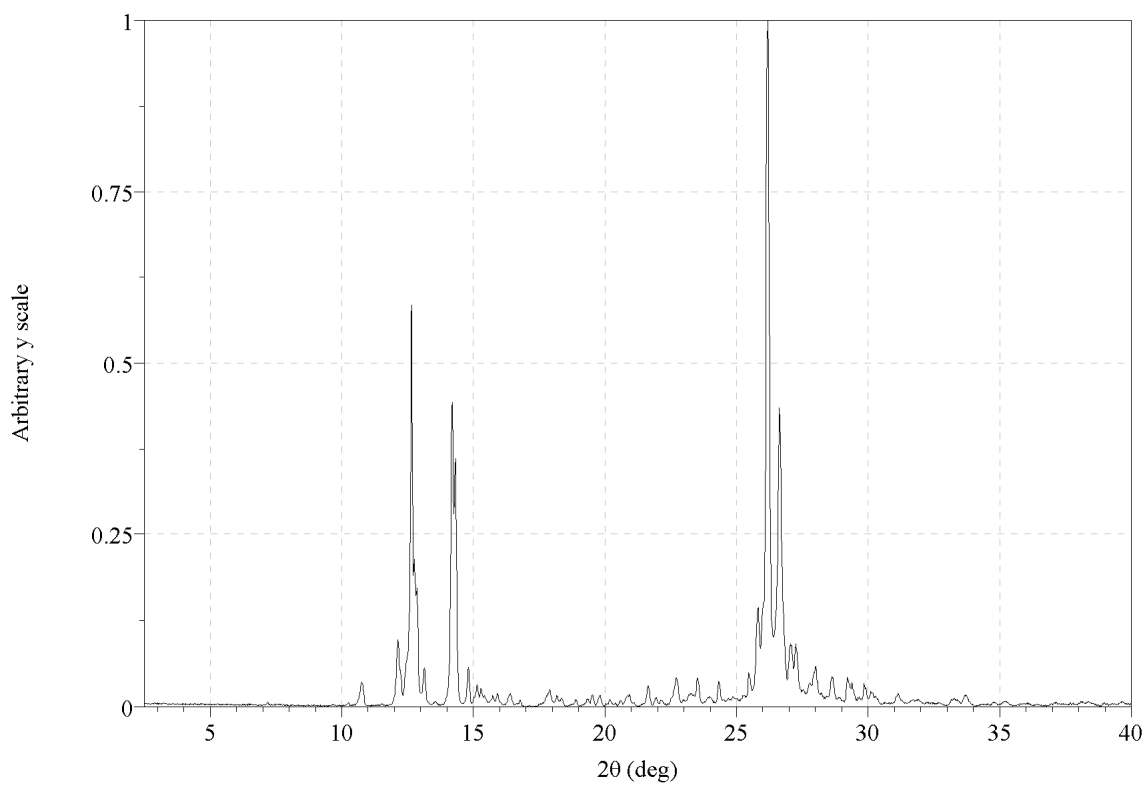


Figure 5. DSC and TGA overlay of 1

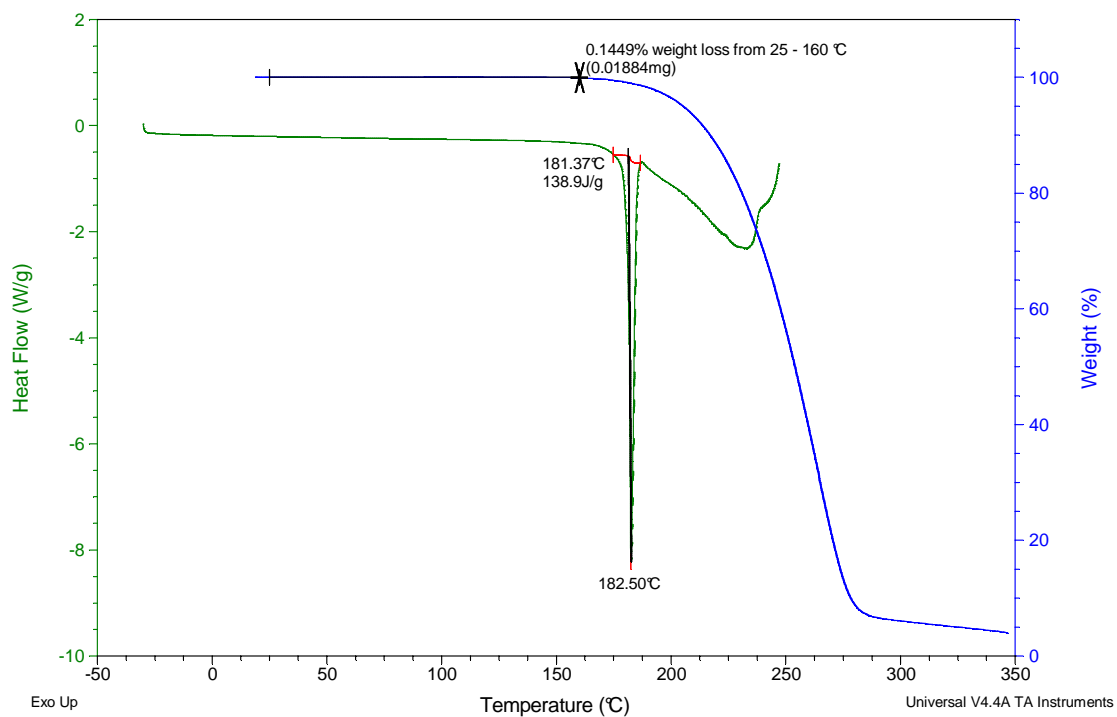


Figure 6. DSC and TGA overlay of **2**

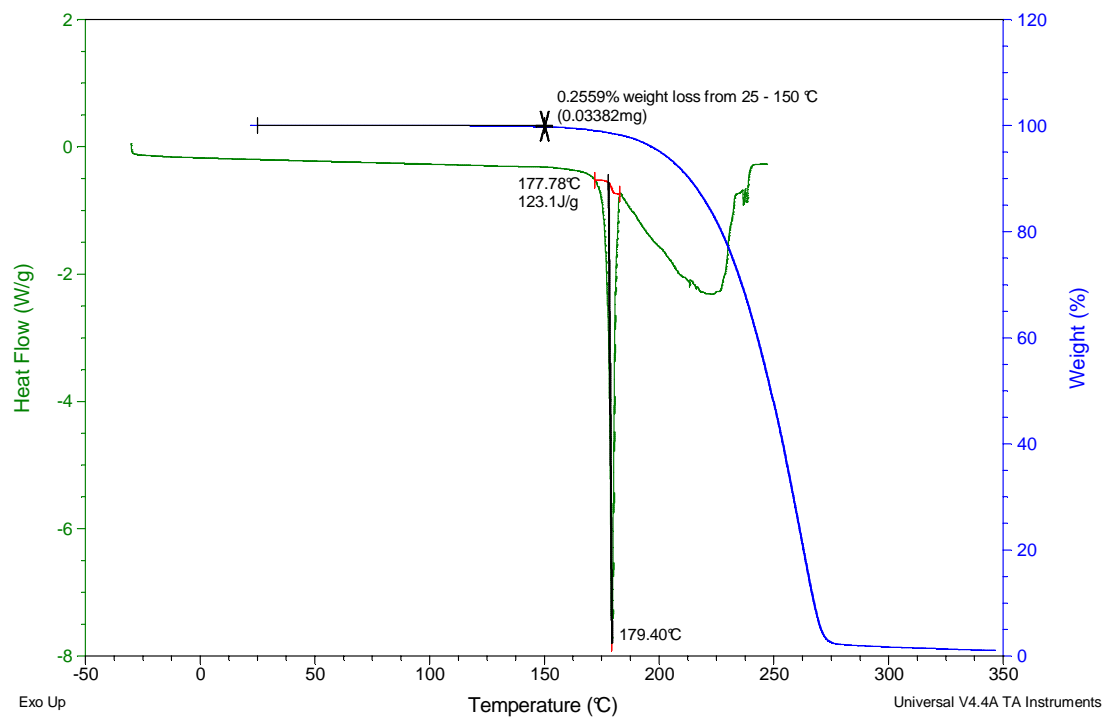


Figure 7. DSC and TGA overlay of **3** (Form I)

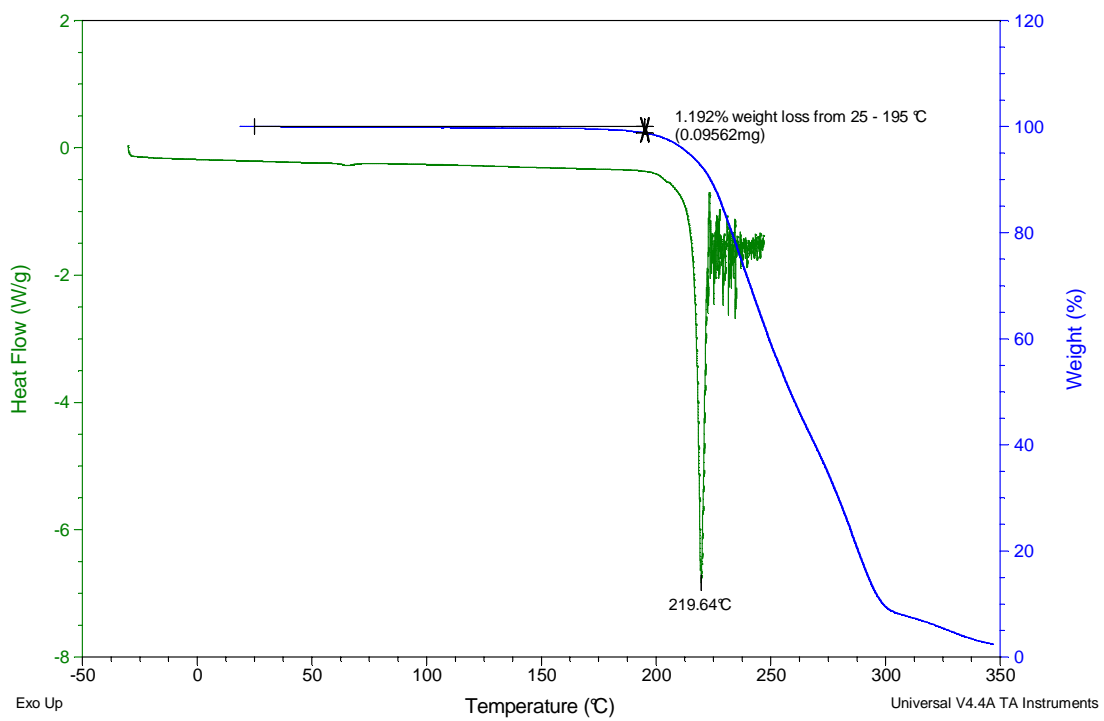


Figure 8. DSC and TGA overlay of **4** (Form II)

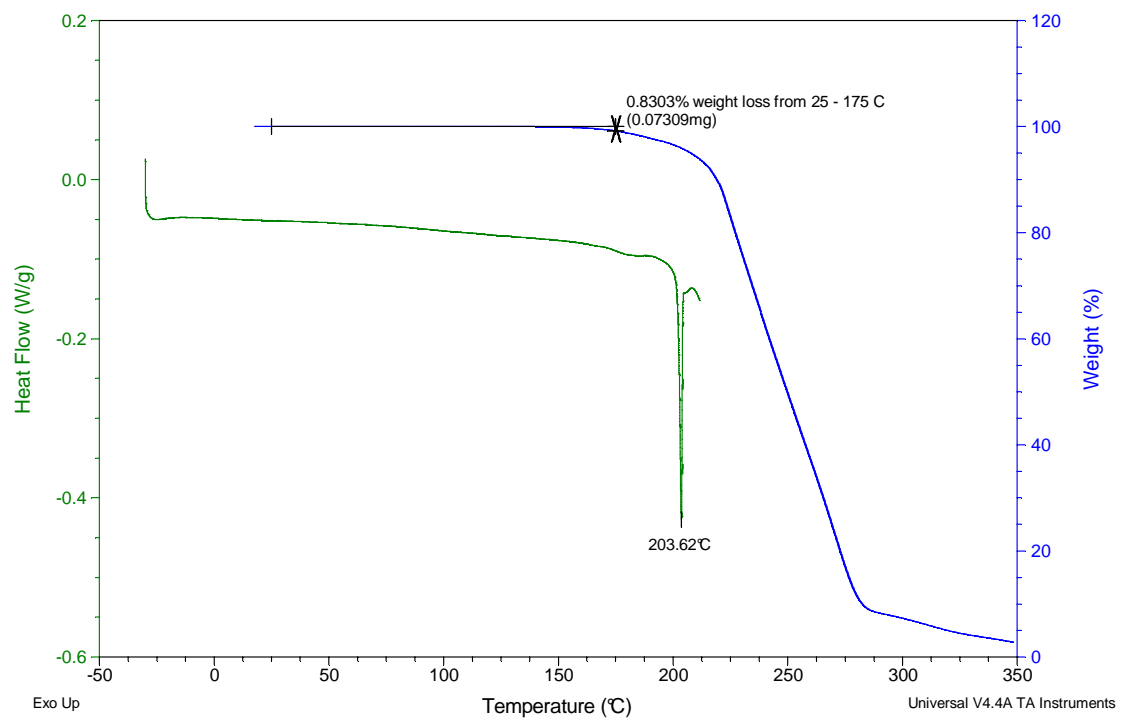


Table 1. Crystal graph for Form I at the default bond cutoff of -0.596 kcal/mol.**key**

Number	Molecule	Reference Coordinate ¹
1	C7 H8 N4 O2	(0.0221497, 0.574097, 0.729905)
2	C7 H8 N4 O2	(0.589332, 0.0899427, 0.867573)
3	C7 H8 N4 O2	(0.410668, 0.9100573, 0.13243)
4	C7 H8 N4 O2	(0.9778503, 0.425903, 0.270095)
5	C9 H8 O3	(0.499478, 0.420611, 0.293522)
6	C9 H8 O3	(0.164907, 0.941541, 0.398635)
7	C9 H8 O3	(0.500522, 0.579389, 0.706478)
8	C9 H8 O3	(0.835093, 0.0584591, 0.601365)

interactions

Molecule 1	Molecule 2	Translation ²	Multiplicity ³	Length (Å) ⁴	Energy (kcal/mol)
1	5	[-1 0 0]	2	8.75699	-13.0476
2	6	[0 -1 0]	2	8.75689	-12.7897
2	3	[0 -1 1]	1	4.23979	-10.5289
1	7	[-1 0 0]	2	4.05232	-9.37448
1	5	[0 0 1]	2	9.00814	-8.54775
2	6	[1 -1 1]	2	8.97188	-8.45311
2	8	[0 0 0]	2	4.34486	-7.63309
1	7	[0 0 0]	2	3.67814	-7.27969
6	8	[0 1 0]	1	6.74735	-4.95618
6	1	[0 0 0]	2	6.18964	-4.43241
5	7	[0 0 0]	1	7.08201	-4.18584
2	7	[0 0 0]	2	6.65223	-4.13371
6	4	[-1 1 0]	2	6.62399	-4.10297
2	8	[-1 0 0]	2	7.40349	-4.09185
5	7	[0 0 -1]	1	8.34279	-3.81668
2	1	[0 0 0]	2	7.59937	-3.50016
2	3	[1 -1 1]	1	7.55594	-3.24017

¹ The reference coordinate represents the position of the molecule in the unit cell, in fractional coordinates, by its center of mass.

² The translation to a neighbouring unit cell in which molecule 2 is located.

³ Space group symmetry applies to the interactions. Multiplicity is the number of occurrences of the interaction in the unit cell.

⁴ Length is the distance between reference points.

1	4	[-1 0 0]	1	7.70752	-2.92737
6	4	[-1 0 0]	2	8.33562	-2.18623
2	7	[0 -1 0]	2	8.33297	-2.13929
6	8	[-1 1 0]	1	4.34183	-2.09630
6	7	[0 0 0]	2	6.31560	-1.93940
2	5	[0 0 1]	2	8.79523	-1.69596
2	5	[0 0 0]	2	8.52143	-1.57281
5	7	[1 0 0]	1	11.0010	-1.47991
6	5	[0 1 0]	2	7.22027	-1.44556
2	1	[1 0 0]	2	7.61871	-1.25770
1	4	[-1 0 1]	1	7.66426	-1.18557
2	4	[0 0 1]	2	9.43033	-0.988586
6	5	[-1 1 0]	2	7.93649	-0.861979
6	1	[1 0 0]	2	8.87328	-0.853893
2	1	[1 -1 0]	2	8.41860	-0.840349
2	1	[0 -1 0]	2	9.64668	-0.786347
1	5	[0 0 0]	2	7.88932	-0.747859
2	4	[0 0 0]	2	9.20499	-0.711493
6	5	[0 0 0]	2	8.08576	-0.694136
6	6	[-1 0 0]	2	7.70166	-0.662726
6	8	[-1 1 -1]	1	11.9018	-0.648548
2	4	[0 -1 1]	2	9.93399	-0.637214
1	1	[-1 0 0]	2	7.70166	-0.615454
2	2	[-1 0 0]	2	7.70166	-0.613198

Table 2. Crystal graph for Form II at the default bond cutoff of -0.596 kcal/mol.**key**

Number	Molecule	Reference Coordinate
1	C7 H8 N4 O2	(0.195864, 0.665954, 0.907673)
2	C7 H8 N4 O2	(0.695864, 0.834046, 0.407673)
3	C7 H8 N4 O2	(0.804136, 0.334046, 0.09233)
4	C7 H8 N4 O2	(0.304136, 0.165954, 0.59233)
5	C9 H8 O3	(0.246106, 0.424849, 0.216406)
6	C9 H8 O3	(0.253894, 0.924849, 0.283594)
7	C9 H8 O3	(0.753894, 0.575151, 0.783594)
8	C9 H8 O3	(0.746106, 0.075151, 0.716406)

interactions

Molecule 1	Molecule 2	Translation	Multiplicity	Length (Å)	Energy (kcal/mol)
1	5	[0 0 0]	4	8.77147	-13.1050
1	6	[0 0 0]	4	8.74081	-7.11590
1	7	[-1 0 0]	4	3.82655	-7.04750
1	7	[0 0 0]	4	4.82726	-6.51352
5	7	[0 0 0]	2	6.68224	-6.12720
5	7	[-1 0 0]	2	7.64219	-4.23363
1	2	[-1 0 0]	4	6.60999	-4.13485
1	5	[0 0 1]	4	6.83216	-2.75957
1	1	[0 0 -1]	4	8.70347	-2.21550
5	7	[-1 0 -1]	2	5.97089	-2.15436
1	7	[-1 0 -1]	4	9.89452	-2.07529
1	6	[0 0 1]	4	7.49143	-2.03350
1	7	[0 0 1]	4	8.12411	-1.75621
1	2	[-1 0 1]	4	7.51421	-1.70961
5	5	[0 0 -1]	4	8.70347	-1.01598
1	3	[0 0 1]	2	9.61863	-0.785475
5	7	[0 0 -1]	2	6.88794	-0.707682
1	1	[-1 0 0]	4	6.87812	-0.677421