

Experimental and Computational Approaches to Produce and Characterise Isostructural Solvates

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1. Pawley Fitting

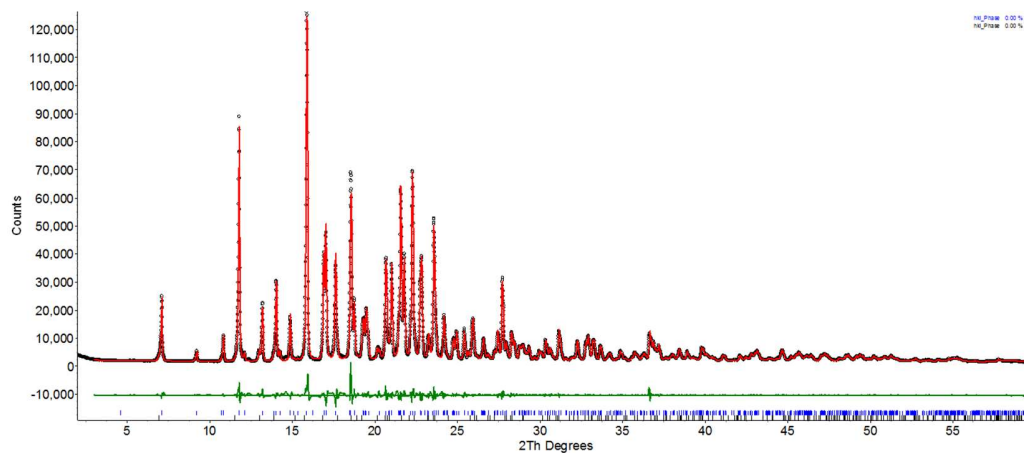


Figure S1. Pawley fit between the PXRD data of $S_{0.5ACN}$ with a model consisting of the cell parameters derived from indexing the PXRD pattern (note that form III impurities are present). Black dots indicate the raw data, while the red line indicates the calculated model. Tick marks (blue – solvate and black – form III) are the 2θ positions for the hkl reflections. The difference pattern is shown in green.

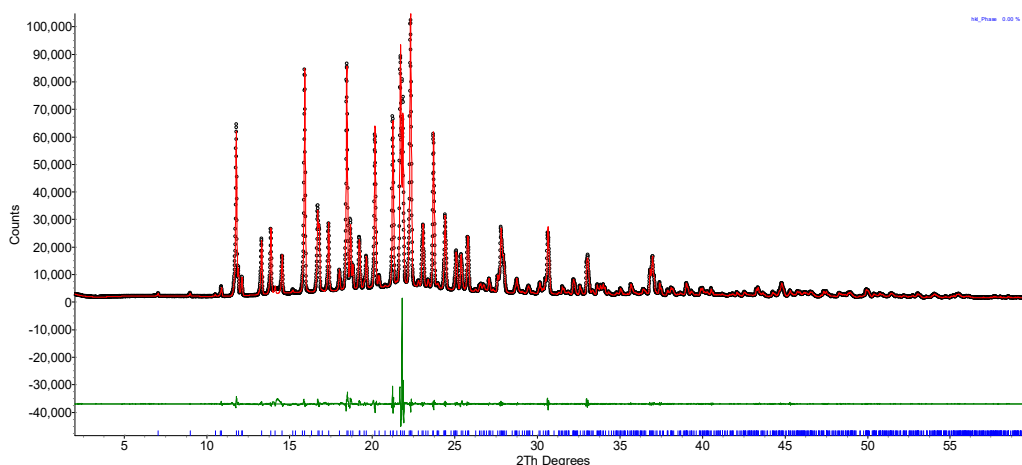


Figure S2. Pawley fit between the PXRD data of $S_{0.5NM}$ with a model consisting of the cell parameters derived from indexing the PXRD pattern. Black dots indicate the raw data, while the red line indicates the calculated model. Tick marks are the 2θ positions for the hkl reflections. The difference pattern is shown in green.

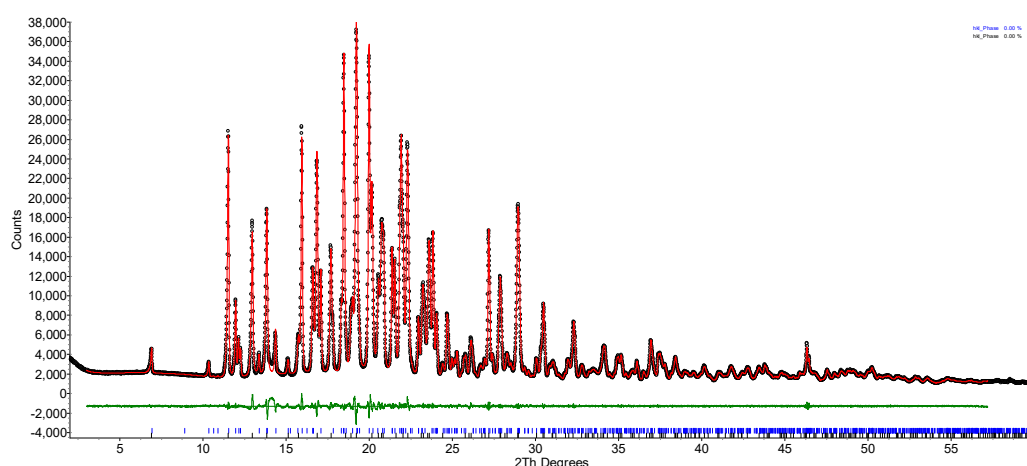


Figure S3. Pawley fit between the PXRD data of **S_{0.5DCM}** with a model consisting of the cell parameters derived from indexing the PXRD pattern (note that form **III** impurities are present). Black dots indicate the raw data, while the red line indicates the calculated model. Tick marks (blue – solvate and black – form **III**) are the 2θ positions for the hkl reflections. The difference pattern is shown in green.

Table S1. Unit cell parameters and space group symmetry derived from indexation of DDS hemisolvates and Form **III** PXRD patterns recorded at room temperature.

Solid Form	SG	a / Å	b / Å	c / Å	β / °	Volume / Å ³	R_{exp}	R_{wp}	R_p
<i>Anhydrate</i>									
Form III	$P2_12_12_1$	5.7501(<1)	8.0472(<1)	25.4957(3)	90	1179.73(2)	3.65	3.95	3.04
<i>Hemi-Solvates</i>									
S_{0.5ACN}	$P2_1/c$	8.5352(2)	16.3078(4)	22.2536(7)	120.500(1)	2668.87(14)	1.49	6.04	3.69
S_{0.5NM}	$P2_1/c$	8.4638(3)	16.2489(7)	22.8563(9)	120.578(<1)	2705.71(19)	1.50	3.21	4.26
S_{0.5DCM}	$P2_1/c$	8.3560(2)	16.6089(4)	20.5183(7)	104.250(1)	2759.99(15)	1.65	3.10	1.91

2. Single Crystal X-ray Diffraction

Table S2. Crystallographic data.

Compound	S_{0.5ACN}	S_{0.5NM}
Chemical formula	C ₂₆ H ₂₇ N ₅ O ₄ S ₂	C ₂₅ H ₂₇ N ₅ O ₆ S ₂
Chemical formula	2(C ₁₂ H ₁₂ N ₂ O ₂ S) · C ₂ H ₃ N	2(C ₁₂ H ₁₂ N ₂ O ₂ S) · CH ₃ NO ₂
Formula Mass	537.64	557.63
Crystal system	Monoclinic	Monoclinic
Space group	<i>P</i> 2 ₁ / <i>n</i>	<i>P</i> 2 ₁ / <i>n</i>
No. of formula units per unit cell, <i>Z</i>	4	4
<i>a</i> / Å	8.5132(2)	8.4167(2)
<i>b</i> / Å	16.1830(3)	16.3415(3)
<i>c</i> / Å	18.9865(4)	19.1383(4)
<i>β</i> / °	97.818(2)	95.832(2)
Unit cell volume / Å ³	2591.44(10)	2618.69(10)
Temperature / K	100	100
No. of reflections measured	23182	48993
No. of independent reflections	5290	4977
No. of parameters	386	395
<i>R</i> _{int}	0.0376	0.0866
Final <i>R</i> 1 value [<i>I</i> > 2σ(<i>I</i>)]	0.0393	0.0491
Final <i>wR</i> (<i>F</i> ²) value (all data)	0.1056	0.1276
CCDC no.	1915951	1915952

3. Crystal Explorer Calculations

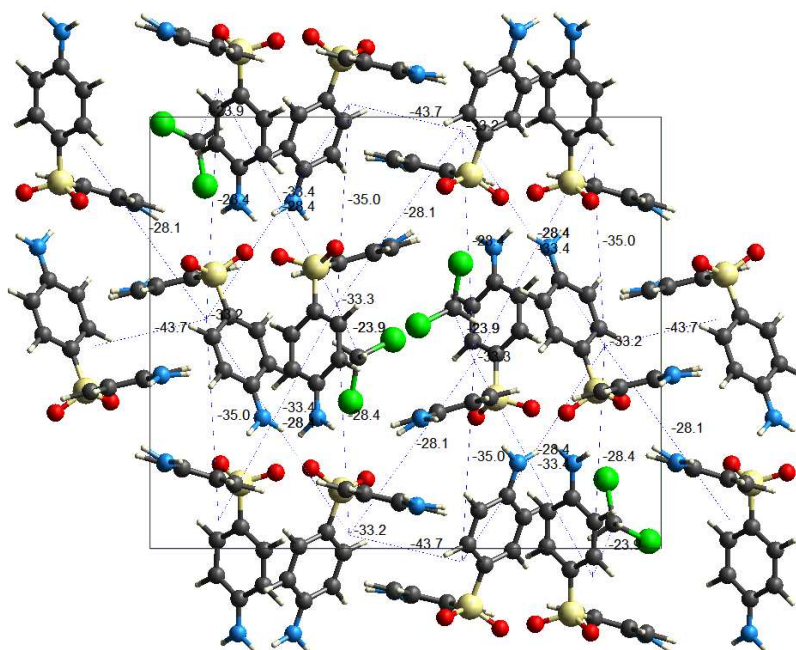


Figure S4. Energy frameworks (total energy, annotated) for $S_{0.5DCM}$, viewed along the crystallographic *a* axis. Pair-wise interaction energies with magnitudes smaller than 20 kJ mol⁻¹ have been omitted.

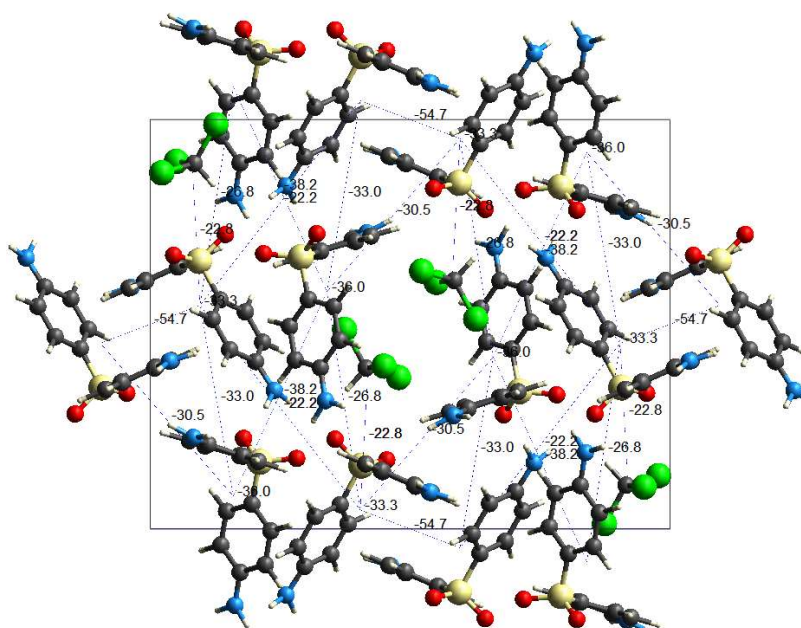


Figure S5. Energy frameworks (total energy, annotated) for $S_{0.5SCHLF}$, viewed along the crystallographic *a* axis. Pair-wise interaction energies with magnitudes smaller than 20 kJ mol⁻¹ have been omitted.

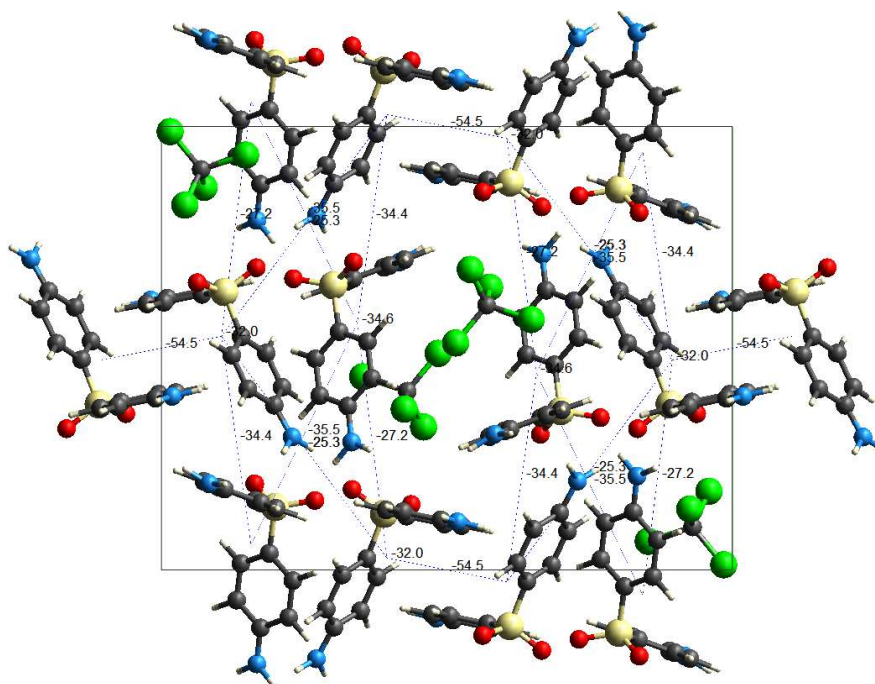


Figure S6. Energy frameworks (total energy, annotated) for $S_{0.5CTC}$, viewed along the crystallographic a axis. Pair-wise interaction energies with magnitudes smaller than 20 kJ mol^{-1} have been omitted.

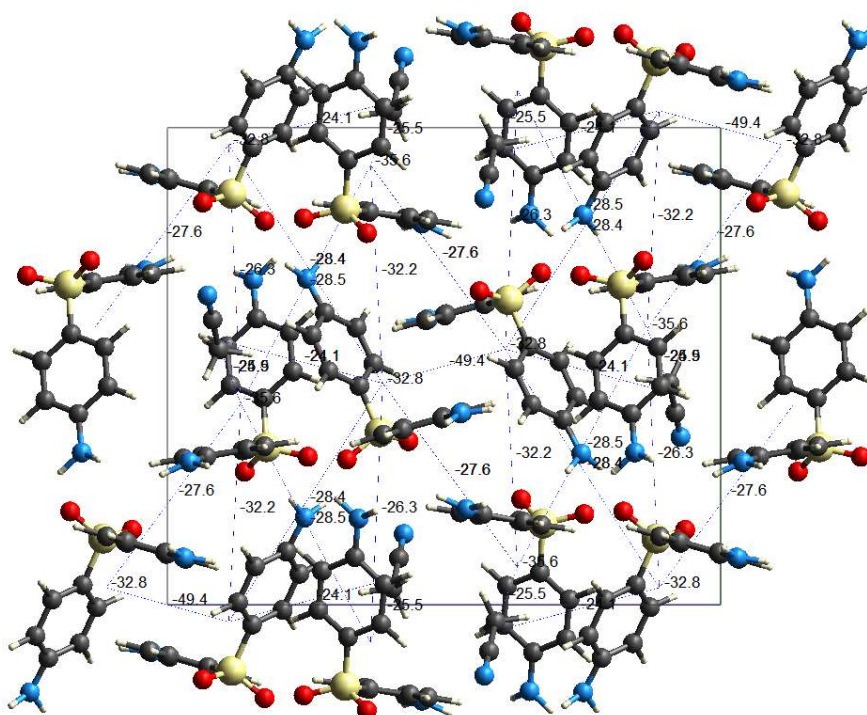


Figure S7. Energy frameworks (total energy, annotated) for $S_{0.5ACN}$, viewed along the crystallographic a axis. Pair-wise interaction energies with magnitudes smaller than 20 kJ mol^{-1} have been omitted.

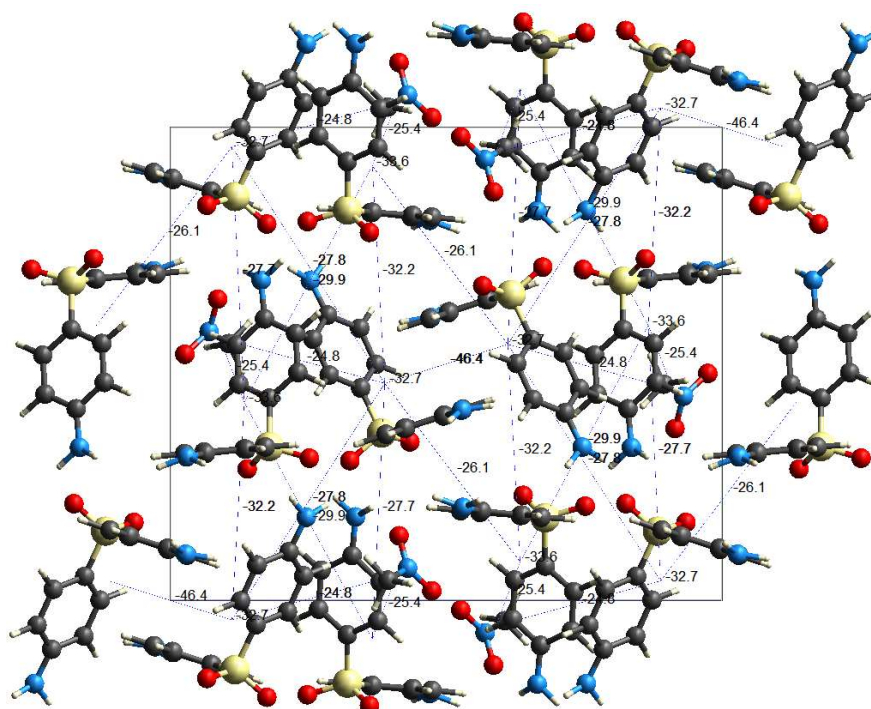


Figure S8. Energy frameworks (total energy, annotated) for $S_{0.5NM}$, viewed along the crystallographic a axis. Pair-wise interaction energies with magnitudes smaller than 20 kJ mol^{-1} have been omitted.

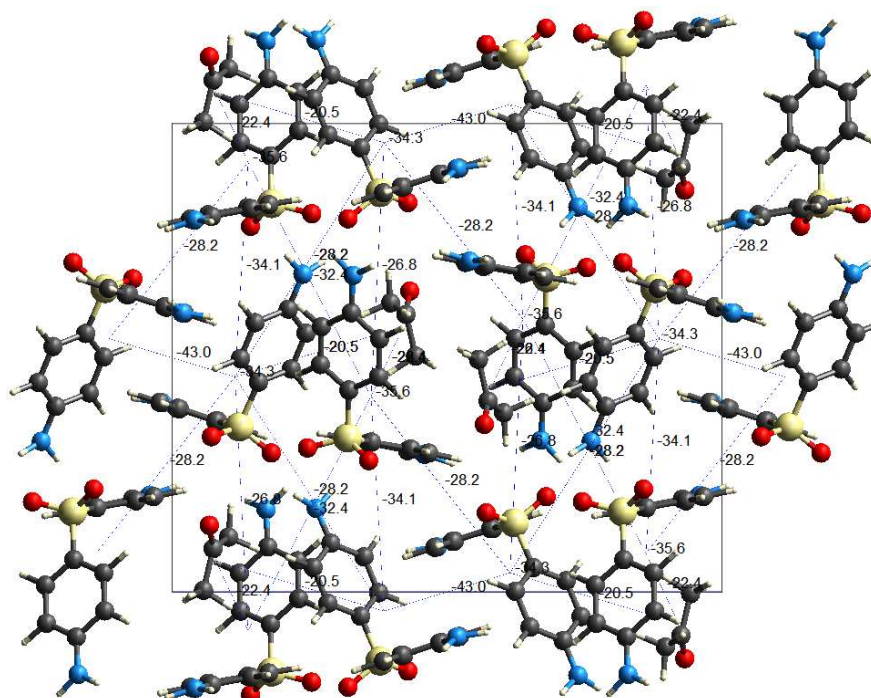


Figure S9. Energy frameworks (total energy, annotated) for $S_{0.5ACO}$, viewed along the crystallographic a axis. Pair-wise interaction energies with magnitudes smaller than 20 kJ mol^{-1} have been omitted.

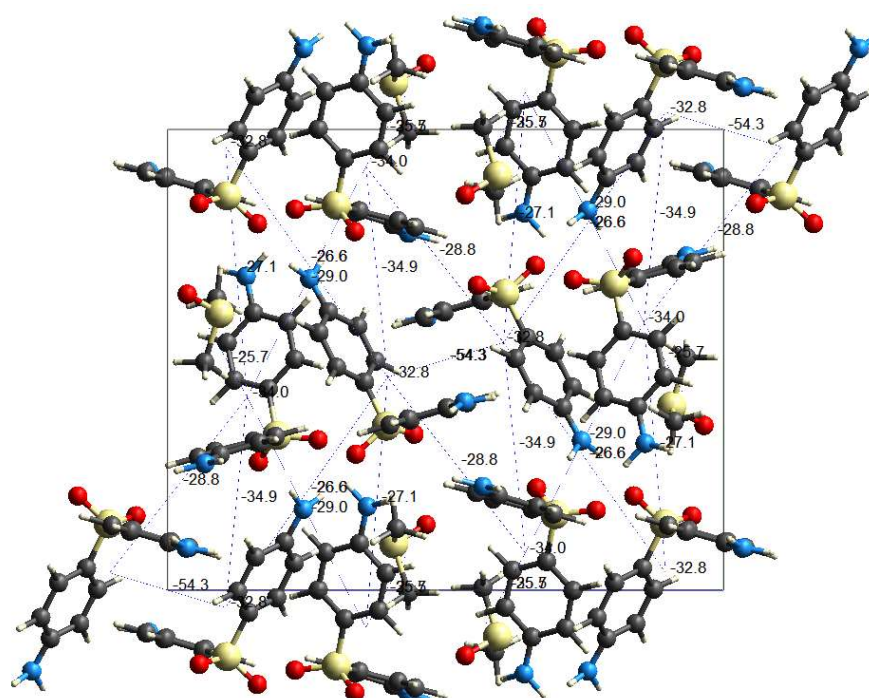


Figure S10. Energy frameworks (total energy, annotated) for $S_{0.5DMSO}$, viewed along the crystallographic *a* axis. Pair-wise interaction energies with magnitudes smaller than 20 kJ mol⁻¹ have been omitted.

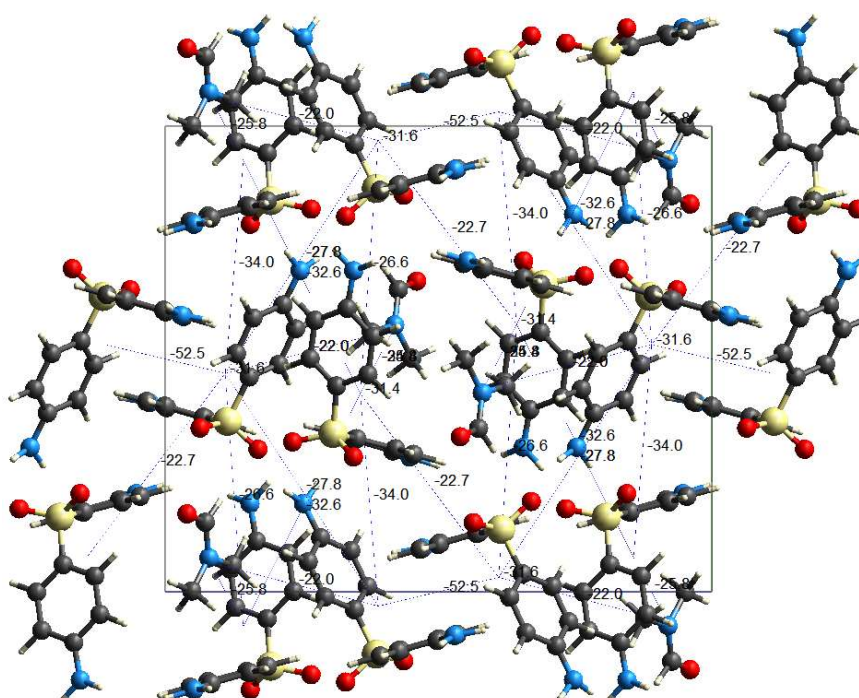


Figure S11. Energy frameworks (total energy, annotated) for $S_{0.5DMF}$, viewed along the crystallographic *a* axis. Pair-wise interaction energies with magnitudes smaller than 20 kJ mol⁻¹ have been omitted.

Table S3. Pair-wise intermolecular interaction energies^a of DDS S_{0.5DCM} (PBE-TS structure).

Inter-action	Molecules Involved	Distance (Å)	E_E (kJ mol ⁻¹)	E_P (kJ mol ⁻¹)	E_D (kJ mol ⁻¹)	E_R (kJ mol ⁻¹)	E_{tot}^b (kJ mol ⁻¹)
S-0.5DCM, $E_{cluster} = -184.9$ kJ mol⁻¹							
1 (1)	DDS...DDS	6.53	-15.5	-5.9	-48.9	31.8	-43.7
2 (2)	DDS...DDS	9.53	-38.5	-9.4	-9.9	34.3	-35
3 (2)	DDS...DDS	9.36	-33.4	-8.7	-9.2	26.5	-33.4
4 (2)	DDS...DDS	8.28	-29.5	-8.8	-17.7	32.2	-33.3
5 (2)	DDS...DDS	8.28	-28	-8.5	-18.3	30.1	-33.2
6 (2)	DDS...DDS	10.36	-21.1	-4.9	-8.7	8.2	-28.4
7 (2)	DDS...DDS	9.93	-27.4	-6.7	-6.9	18.6	-28.4
8 (2)	DDS...DDS	10.29	-23.1	-5.3	-6.5	9.5	-28.1
9 (2)	DDS...S	3.8	-9.8	-2.8	-29.8	23.4	-23.9
10 (1)	DDS...DDS	6.24	-7.2	-2	-25.6	18.9	-19.7
11 (2)	DDS...DDS	6.33	1.4	-6.3	-27.8	21.4	-14.3
12 (2)	DDS...S	6.89	-8.5	-1.3	-11	10.5	-13.1
13 (2)	DDS...DDS	6.29	7.2	-5.4	-30.4	18.5	-11.4
14 (2)	DDS...S	6.91	-6	-0.7	-7.8	4.6	-10.8
15 (1)	S...S	4.93	-7.7	-0.7	-7.1	7.9	-10
16 (2)	DDS...DDS	7.95	3.7	-4.1	-22.6	14.4	-9.9
17 (2)	DDS...S	4.32	-6.9	-0.7	-19.8	24.6	-9.9
18 (1)	DDS...DDS	9.48	-3.2	-0.3	-2.2	0	-5.5
19 (2)	DDS...DDS	9.53	-2.9	-0.5	-2.4	0.1	-5.4
20 (1)	DDS...DDS	6.47	9.5	-2.3	-18.4	5.3	-4.3
21 (2)	DDS...S	8.51	-1.7	-0.3	-2.9	0.5	-4.1
22 (2)	DDS...DDS	8.29	4.2	-2.3	-13.3	7.8	-4
23 (2)	DDS...S	6.08	2.4	-1.3	-10.1	9.7	-1.3
24 (1)	S...S	6.56	0.2	0	-0.9	0.1	-0.5
25 (2)	DDS...DDS	9.23	4.2	-1.1	-5.5	1.2	-0.4
26 (2)	DDS...DDS	9.11	0.2	0	-0.4	0	-0.1
27 (2)	DDS...DDS	9.56	0.6	0	-0.3	0	0.4

^a(1) – interaction present once, (2) – interaction present twice. ^bElectrostatic (E_E), polarization (E_P), dispersion (E_D) and exchange-repulsion energy (E_R) contributions. ^b $E_{tot} = k_E E_E + k_P E_P + k_D E_D + k_R E_R$, with k being scale factors.

Table S4. Pair-wise intermolecular interaction energies^a of DDS S_{0.5CHLF} (PBE-TS structure).

Inter-action	Molecules Involved	Distance (Å)	E_E (kJ mol ⁻¹)	E_P (kJ mol ⁻¹)	E_D (kJ mol ⁻¹)	E_R (kJ mol ⁻¹)	E_{tot}^b (kJ mol ⁻¹)
S-0.5CHLF, $E_{cluster} = -189.4$ kJ mol⁻¹							
1 (1)	DDS...DDS	6.30	-27.2	-9	-65.6	61.2	-54.7
2 (2)	DDS...DDS	8.67	-37	-10.5	-16.2	36.9	-38.2
3 (2)	DDS...DDS	8.43	-34.6	-9.6	-19.4	39.9	-36
4 (2)	DDS...DDS	8.43	-29.1	-8.6	-17.8	31.4	-33.3
5 (2)	DDS...DDS	9.63	-31.7	-7.7	-7.7	21	-33
6 (2)	DDS...DDS	10.01	-26.6	-6.6	-7.8	15.2	-30.5
7 (2)	DDS...DDS	10.32	-20.8	-5.2	-9	11.2	-26.8
8 (2)	DDS...S	5.52	-13.4	-4.5	-17.7	16.3	-22.8
9 (2)	DDS...DDS	10.24	-17	-4.1	-5.6	5.9	-22.2
10 (1)	DDS...DDS	6.05	-8.1	-2.4	-29.7	27.1	-19.5
11 (2)	DDS...DDS	6.22	-4.9	-7.5	-30.1	31.6	-17.4
12 (2)	DDS...S	4.06	-7	-1	-28.8	34.4	-12
13 (2)	DDS...DDS	6.4	6.9	-5	-33.7	25.2	-10.1
14 (2)	DDS...DDS	8.1	0.7	-3.2	-19.8	14.8	-9.8
15 (2)	DDS...DDS	7.71	0.3	-4.3	-26.3	26	-9.7
16 (2)	DDS...DDS	8.98	-4.7	-0.8	-3.4	0.2	-8.4
17 (2)	DDS...S	4.98	-4.3	-0.3	-15.7	17.3	-7.8
18 (2)	DDS...S	7.96	-4.8	-0.7	-1.2	0	-6.6
19 (2)	DDS...S	7.52	-4.1	-1	-11.3	15.3	-5.5
20 (2)	DDS...S	7.65	0.2	-1.2	-11.1	9	-4.8
21 (2)	DDS...S	8.86	-6.1	-0.4	-6	12.4	-4.3
22 (1)	DDS...DDS	10.80	-2.3	-0.2	-1.1	0	-3.6
23 (1)	S...S	4.81	-0.3	-0.1	-5.6	3.5	-3.1
24 (1)	S...S	6.77	0.2	0	-1	0	-0.6
25 (2)	DDS...DDS	8.91	4.7	-1.2	-5.4	1.1	0.1
26 (1)	DDS...DDS	7.70	6.6	-0.6	-5.3	0.1	2

^a(1) – interaction present once, (2) – interaction present twice. ^bElectrostatic (E_E), polarization (E_P), dispersion (E_D) and exchange-repulsion energy (E_R) contributions. ^b $E_{tot} = k_E E_E + k_P E_P + k_D E_D + k_R E_R$, with k being scale factors.

Table S5. Pair-wise intermolecular interaction energies^a of DDS S_{0.5CTC} (PBE-TS structure).

Inter-action	Molecules Involved	Distance (Å)	E_E (kJ mol ⁻¹)	E_P (kJ mol ⁻¹)	E_D (kJ mol ⁻¹)	E_R (kJ mol ⁻¹)	E_{tot}^b (kJ mol ⁻¹)
S-0.5CTC, $E_{cluster} = -170.9$ kJ mol⁻¹							
1 (1)	DDS...DDS	6.32	-23.6	-8.2	-56.7	41.9	-54.5
2 (2)	DDS...DDS	8.82	-33.3	-9.9	-14.4	31.7	-35.5
3 (2)	DDS...DDS	8.72	-34	-9	-14.9	33.9	-34.6
4 (2)	DDS...DDS	9.7	-36	-8.9	-9.2	29.5	-34.4
5 (2)	DDS...DDS	8.72	-29	-7.9	-13.6	26.5	-32
6 (2)	DDS...DDS	10.36	-19.7	-4.7	-8.5	7.3	-27.2
7 (2)	DDS...DDS	10.08	-20.9	-5.1	-6.1	9.5	-25.3
8 (2)	DDS...DDS	10.96	-11.5	-1.8	-2.9	0.3	-15.8
9 (2)	DDS...DDS	6.35	-0.9	-7.4	-24.8	19.9	-15.6
10 (2)	DDS...S	4.14	-4.7	-0.8	-28.6	27.3	-13.5
11 (1)	DDS...DDS	6.48	-0.2	-1.8	-17	6.2	-12.5
12 (2)	DDS...DDS	6.45	7.3	-5.9	-28.7	17.1	-11.1
13 (2)	DDS...S	5.68	-8.4	-2.3	-18.3	25.6	-10.8
14 (2)	DDS...S	4.98	-5.3	-0.3	-20.6	21.4	-10.6
15 (2)	DDS...DDS	8.18	0.5	-2.9	-20.2	15	-10
16 (2)	DDS...DDS	8.31	1.2	-2.8	-18.6	12.9	-9.1
17 (2)	DDS...S	7.3	-6	-1.1	-14.1	19.6	-7.4
18 (2)	DDS...S	7.4	-8.5	-1	-12.9	23.1	-6.7
19 (2)	DDS...DDS	9.28	-2.6	-0.5	-1.2	0	-4.2
20 (1)	DDS...DDS	10.38	-2.6	-0.2	-1.2	0	-4
21 (1)	S...S	6.95	0	0	-1.4	0	-1.2
22 (1)	S...S	5.33	-8.2	-0.1	-10.2	26.7	-1.1
23 (2)	DDS...DDS	8.41	0.4	-0.5	-1	0	-0.9
24 (2)	DDS...S	9.2	0.8	-0.5	-5.5	6.3	-0.4
25 (2)	DDS...S	10.83	0.5	-0.1	-0.8	0	-0.3
26 (1)	DDS...DDS	7.45	5.7	-0.6	-6	0.1	0.5

^a(1) – interaction present once, (2) – interaction present twice. ^bElectrostatic (E_E), polarization (E_P), dispersion (E_D) and exchange-repulsion energy (E_R) contributions. ^b $E_{tot} = k_E E_E + k_P E_P + k_D E_D + k_R E_R$, with k being scale factors.

Table S6. Pair-wise intermolecular interaction energies^a of **DDS S_{0.5ACN}** (PBE-TS structure).

Inter-action	Molecules Involved	Distance (Å)	E_E (kJ mol ⁻¹)	E_P (kJ mol ⁻¹)	E_D (kJ mol ⁻¹)	E_R (kJ mol ⁻¹)	E_{tot}^b (kJ mol ⁻¹)
S-0.5ACN, $E_{cluster} = -194.8$ kJ mol⁻¹							
1 (1)	DDS...DDS	6.38	-19.7	-7	-55.7	40.6	-49.4
2 (2)	DDS...DDS	8.51	-33.2	-9.2	-16.7	33.8	-35.6
3 (2)	DDS...DDS	8.51	-28	-7.8	-15.6	26.2	-32.8
4 (2)	DDS...DDS	9.35	-29	-8	-11.4	23.4	-32.2
5 (2)	DDS...DDS	9.16	-23	-7.3	-10.2	16.3	-28.5
6 (2)	DDS...DDS	9.69	-28.9	-7.8	-8.8	25.1	-28.4
7 (2)	DDS...DDS	10.17	-22.9	-5.4	-6.9	10.7	-27.6
8 (2)	DDS...DDS	10.4	-18.8	-4.4	-8.2	6.4	-26.3
9 (2)	DDS...S	4.07	-13.1	-3.1	-26.2	21.8	-25.5
10 (2)	DDS...S	6.97	-22.3	-4	-8.9	15	-24.9
11 (2)	DDS...S	6.47	-15.7	-2.4	-12.8	8.8	-24.1
12 (1)	DDS...DDS	6.26	-4.8	-2	-24.1	16.1	-17.6
13 (2)	DDS...DDS	6.39	1.8	-5.9	-27.9	18.1	-15.6
14 (2)	DDS...DDS	6.19	5.3	-6.2	-31.8	23.7	-12.2
15 (2)	DDS...DDS	7.44	3.2	-5	-28	20.2	-12.2
16 (2)	DDS...S	8.72	-9.5	-1.9	-4	5.4	-11.7
17 (1)	DDS...DDS	6.48	6	-3.7	-26	14.4	-10.2
18 (2)	DDS...S	4.36	-1	-2.1	-14	9.8	-8.8
19 (1)	DDS...DDS	9.46	-2.4	-0.5	-3.9	0.3	-6.2
20 (2)	DDS...DDS	8.03	3.2	-2	-11.8	4.2	-5.8
21 (2)	DDS...DDS	9.2	-2.1	-0.6	-3.4	0.3	-5.4
22 (2)	DDS...S	10.68	-3.3	-0.2	-0.3	0	-3.9
23 (1)	S...S	5.40	2.2	-0.8	-3.7	1.5	-0.5
24 (1)	S...S	13.69	0.7	-0.2	-0.7	0	0
25 (2)	DDS...DDS	9.24	4.4	-0.8	-4.5	0.8	0.6
26 (2)	DDS...S	5.7	17.1	-3.1	-6.9	5.4	13.2

^a(1) – interaction present once, (2) – interaction present twice. ^bElectrostatic (E_E), polarization (E_P), dispersion (E_D) and exchange-repulsion energy (E_R) contributions. ^b $E_{tot} = k_E E_E + k_P E_P + k_D E_D + k_R E_R$, with k being scale factors.

Table S7. Pair-wise intermolecular interaction energies^a of **DDS S_{0.5NM}** (PBE-TS structure).

Inter-action	Molecules Involved	Distance (Å)	E_E (kJ mol ⁻¹)	E_P (kJ mol ⁻¹)	E_D (kJ mol ⁻¹)	E_R (kJ mol ⁻¹)	E_{tot}^b (kJ mol ⁻¹)
S-0.5NM, $E_{cluster} = -194.0$ kJ mol⁻¹							
1 (1)	DDS...DDS	6.38	-19.6	-6.9	-54	42.7	-46.4
2 (2)	DDS...DDS	8.42	-26.8	-7.8	-17.2	25	-33.6
3 (2)	DDS...DDS	8.42	-27.2	-7.9	-16.3	26.1	-32.7
4 (2)	DDS...DDS	9.26	-31.1	-8.5	-12	28.3	-32.2
5 (2)	DDS...DDS	9.17	-27.2	-8.2	-10.6	23	-29.9
6 (2)	DDS...DDS	9.74	-25.9	-6.9	-8.2	19.2	-27.8
7 (2)	DDS...DDS	10.27	-21.2	-5	-9.1	10.2	-27.7
8 (2)	DDS...DDS	10.37	-20.7	-4.6	-6.3	7.5	-26.1
9 (2)	DDS...S	4.03	-11.6	-4.3	-28.5	24	-25.4
10 (2)	DDS...S	6.57	-16.4	-2.6	-14.5	11.5	-24.8
11 (2)	DDS...S	6.64	-14.3	-2	-10	9.7	-19.3
12 (1)	DDS...DDS	6.46	-5	-1.8	-19.5	9.5	-17.6
13 (1)	S...S	4.65	-13	-1.9	-6.5	6.6	-16.7
14 (2)	DDS...DDS	6.33	2.3	-5.8	-29.4	20.4	-14.8
15 (2)	DDS...DDS	7.36	3.8	-5.1	-28.8	19.5	-12.9
16 (2)	DDS...DDS	6.2	5.2	-5.5	-32.1	22.7	-12.5
17 (2)	DDS...S	4.2	-2.9	-2.9	-18.4	17.1	-10.8
18 (1)	DDS...DDS	6.80	4.4	-3	-20	8.9	-9.5
19 (2)	DDS...DDS	9.36	-2.6	-0.5	-3.1	0.2	-5.7
20 (1)	DDS...DDS	9.08	-1.9	-0.5	-3.9	0.2	-5.7
21 (2)	DDS...S	8.96	-2.9	-0.6	-1.7	0.2	-4.9
22 (2)	DDS...S	8.49	-3.5	-0.5	-0.6	0	-4.6
23 (2)	DDS...DDS	8.2	3.3	-1.7	-9.7	2.9	-4.4
24 (2)	DDS...DDS	9.32	4	-0.7	-4	0.5	0.5
25 (1)	S...S	6.41	3.2	-0.2	-0.4	0	2.9
26 (2)	DDS...S	6.05	12	-1.7	-5.3	2.3	8.2

^a(1) – interaction present once, (2) – interaction present twice. ^bElectrostatic (E_E), polarization (E_P), dispersion (E_D) and exchange-repulsion energy (E_R) contributions. ^b $E_{tot} = k_E E_E + k_P E_P + k_D E_D + k_R E_R$, with k being scale factors.

Table S8. Pair-wise intermolecular interaction energies^a of DDS S_{0.5ACO} (PBE-TS structure).

Inter-action	Molecules Involved	Distance (Å)	E_E (kJ mol ⁻¹)	E_P (kJ mol ⁻¹)	E_D (kJ mol ⁻¹)	E_R (kJ mol ⁻¹)	E_{tot}^b (kJ mol ⁻¹)
S-0.5ACO, $E_{cluster} = -197.9$ kJ mol⁻¹							
1 (1)	DDS...DDS	6.55	-18.9	-6.1	-49.5	39.8	-43
2 (2)	DDS...DDS	8.37	-35.3	-10.6	-18.4	41.4	-35.6
3 (2)	DDS...DDS	8.37	-32.2	-9.1	-18.7	36.8	-34.3
4 (2)	DDS...DDS	9.42	-33.6	-8.8	-10.3	27.3	-34.1
5 (2)	DDS...DDS	9.19	-30.5	-8.8	-11	25.7	-32.4
6 (2)	DDS...DDS	10.15	-24.6	-5.9	-7.1	13.6	-28.2
7 (2)	DDS...DDS	9.74	-30.2	-7.6	-8.1	26.6	-28.2
8 (2)	DDS...DDS	10.42	-19.5	-4.6	-8.6	7.6	-26.8
9 (2)	DDS...S	6.93	-23.2	-4.4	-14.8	23.6	-26.1
10 (2)	DDS...S	4.18	-14.2	-2.1	-33.8	38.1	-22.4
11 (2)	DDS...S	6.67	-12	-2.1	-16.5	13.2	-20.5
12 (1)	DDS...DDS	6.49	-5.6	-1.8	-19.6	10.5	-17.9
13 (2)	DDS...DDS	6.27	0.5	-6.6	-30.6	23	-16.8
14 (2)	DDS...DDS	6.11	4.6	-6.1	-35.7	29.4	-12.6
15 (2)	DDS...DDS	7.56	0.7	-4.8	-27.6	23.1	-12.5
16 (2)	DDS...S	8.55	-9.2	-2.1	-4.9	5.7	-12.1
17 (2)	DDS...S	4.44	-1.3	-1.3	-23.3	18.4	-11.2
18 (1)	DDS...DDS	9.67	-3.1	-0.4	-2.6	0	-5.8
19 (2)	DDS...DDS	9.43	-2.9	-0.5	-2.7	0.1	-5.7
20 (1)	S...S	6.48	-2.4	-0.2	-1.5	0.2	-4
21 (2)	DDS...DDS	8.09	4.6	-2.3	-13.6	8.3	-3.6
22 (2)	DDS...S	8.26	-1.8	-0.8	-1.1	0	-3.5
23 (1)	S...S	5.31	1.1	-0.8	-8.1	7	-2.3
24 (1)	DDS...DDS	6.80	12.1	-2.3	-15.4	3.3	-0.3
25 (2)	DDS...DDS	9.24	4.4	-0.9	-4.3	0.6	0.5
26 (2)	DDS...S	5.71	13.6	-3.2	-11.6	8	6.9

^a(1) – interaction present once, (2) – interaction present twice. ^b $E_{tot} = k_E E_E + k_P E_P + k_D E_D + k_R E_R$, with k being scale factors.

Table S9. Pair-wise intermolecular interaction energies^a of DDS S_{0.5DMSO} (PBE-TS structure).

Inter-action	Molecules Involved	Distance (Å)	E_E (kJ mol ⁻¹)	E_P (kJ mol ⁻¹)	E_D (kJ mol ⁻¹)	E_R (kJ mol ⁻¹)	E_{tot}^b (kJ mol ⁻¹)
S-0.5DMSO, $E_{cluster} = -184.9$ kJ mol⁻¹							
1 (1)	DDS...DDS	6.19	-27.2	-8.9	-68.6	65.9	-54.3
2 (2)	DDS...S	6.89	-30.8	-6.6	-18.1	28.7	-35.5
3 (2)	DDS...DDS	9.4	-36.2	-9.3	-9.6	30.2	-34.9
4 (2)	DDS...DDS	8.4	-33.1	-10.7	-17.6	39.1	-34
5 (2)	DDS...DDS	8.4	-29.3	-8.5	-18.8	33.5	-32.8
6 (2)	DDS...DDS	8.96	-20.3	-6.5	-11.2	11.4	-29
7 (2)	DDS...DDS	10.14	-25	-5.9	-7.3	13.4	-28.8
8 (2)	DDS...DDS	10.38	-19.7	-4.9	-9.6	9.3	-27.1
9 (2)	DDS...DDS	9.88	-24.7	-6.2	-6.8	16.2	-26.6
10 (2)	DDS...S	4.3	-18.6	-3.1	-37	46.1	-25.7
11 (1)	DDS...DDS	6.22	-4.7	-2	-24.5	17.2	-17.2
12 (2)	DDS...S	8.48	-14.2	-3.9	-5.2	8.6	-17.2
13 (2)	DDS...DDS	6.39	-0.4	-6.5	-26.8	19.8	-16.4
14 (2)	DDS...S	4.66	-6.6	-4.7	-24.8	27.4	-15.2
15 (2)	DDS...S	6.87	-11.9	-1.9	-14.6	18.7	-15.1
16 (2)	DDS...DDS	7.53	-2.8	-4.7	-29.1	28.5	-14.2
17 (2)	DDS...DDS	6.16	6.6	-6.2	-35.7	30.6	-9.8
18 (1)	S...S	6.32	-6.3	-0.7	-1.5	0	-8.5
19 (2)	DDS...DDS	7.97	2.3	-2.7	-18.2	13.3	-7.2
20 (2)	DDS...S	8.24	-4.8	-0.9	-1.2	0	-6.8
21 (2)	DDS...DDS	9.1	-2.7	-0.6	-3.3	0.2	-6
22 (1)	DDS...DDS	9.92	-2.7	-0.5	-3.1	0.1	-5.8
23 (1)	S...S	5.60	0.4	-0.9	-7.6	5	-3.7
24 (2)	DDS...DDS	9.11	5	-0.9	-4.7	0.8	1
25 (1)	DDS...DDS	7.00	13	-2.4	-14.2	3	1.5
26 (2)	DDS...S	5.52	10.1	-4.9	-16.2	15.7	2.7

^a(1) – interaction present once, (2) – interaction present twice. ^bElectrostatic (E_E), polarization (E_P), dispersion (E_D) and exchange-repulsion energy (E_R) contributions. ^b $E_{tot} = k_E E_E + k_P E_P + k_D E_D + k_R E_R$, with k being scale factors.

Table S10. Pair-wise intermolecular interaction energies^a of DDS S_{0.5DMF} (PBE-TS structure).

Inter-action	Molecules Involved	Distance (Å)	E_E (kJ mol ⁻¹)	E_P (kJ mol ⁻¹)	E_D (kJ mol ⁻¹)	E_R (kJ mol ⁻¹)	E_{tot}^b (kJ mol ⁻¹)
S-0.5DMF, $E_{cluster} = -209.0$ kJ mol⁻¹							
1 (1)	DDS...DDS	6.28	-26.6	-8.4	-65	62.2	-52.5
2 (2)	DDS...S	6.68	-30.8	-6.6	-20.8	34.2	-34.4
3 (2)	DDS...DDS	9.26	-36.8	-9.8	-11.4	35.8	-34
4 (2)	DDS...DDS	9.00	-27.9	-8.3	-11.6	21.3	-32.6
5 (2)	DDS...DDS	8.51	-28.1	-7.8	-14.4	26.5	-31.6
6 (2)	DDS...DDS	8.51	-25.2	-8.2	-15.3	23.7	-31.4
7 (2)	DDS...DDS	9.76	-27	-7	-8.1	21	-27.8
8 (2)	DDS...DDS	10.32	-19.7	-4.9	-9.5	9.8	-26.6
9 (2)	DDS...S	4.38	-20	-2.9	-45.1	59.5	-25.8
10 (2)	DDS...DDS	10.66	-16.8	-3.3	-5	2.9	-22.7
11 (2)	DDS...S	6.62	-12.7	-2.4	-21.5	19.2	-22
12 (2)	DDS...S	8.98	-16.3	-3.7	-5.4	8.3	-19.5
13 (2)	DDS...DDS	6.3	-1.5	-6.9	-30	24.2	-17.8
14 (1)	DDS...DDS	6.31	-4.5	-2	-21.7	12.3	-17.7
15 (2)	DDS...DDS	7.7	0.7	-4.2	-26.7	22.9	-11.5
16 (2)	DDS...DDS	6.09	5.9	-6.4	-36.4	33	-9.7
17 (2)	DDS...S	4.42	0.5	-2.8	-24.9	23.1	-9
18 (2)	DDS...DDS	9.06	-3.6	-0.7	-3.6	0.3	-7.2
19 (2)	DDS...S	8.33	-5.1	-0.9	-1.1	0	-7
20 (1)	S...S	5.29	-1.7	-0.9	-9.8	6.7	-6.8
21 (1)	S...S	6.48	-4.5	-0.5	-2	0.2	-6.8
22 (1)	DDS...DDS	9.29	-3.2	-0.5	-3.4	0.1	-6.5
23 (2)	DDS...DDS	8.19	2.6	-2.2	-14	7.6	-6.4
24 (1)	DDS...DDS	7.02	12.1	-2.6	-14.4	3.5	0.5
25 (2)	DDS...DDS	9.37	4.5	-0.8	-4.2	0.7	0.8
26 (2)	DDS...S	5.83	10.8	-4.5	-14.5	12.3	3.1

^a(1) – interaction present once, (2) – interaction present twice. ^bElectrostatic (E_E), polarization (E_P), dispersion (E_D) and exchange-repulsion energy (E_R) contributions. ^b $E_{tot} = k_E E_E + k_P E_P + k_D E_D + k_R E_R$, with k being scale factors.

4. Packing Similarity

Table S11. CrystalCMP similarity matrix.

	S _{0.5} NM	S _{0.5} ACN	S _{0.5} ACO	S _{0.5} CTC	S _{0.5} DCM	S _{0.5} DMF	S _{0.5} DMSO	S _{0.5} DMSO
S _{0.5} NM								
S _{0.5} ACN	0.3831							
S _{0.5} ACO	0.2535	0.3567						
S _{0.5} CTC	1.3776	1.4667	1.2727					
S _{0.5} DCM	1.6801	1.0122	1.5993	0.9824				
S _{0.5} DMF	0.7888	0.7011	0.7542	1.4982	1.3345			
S _{0.5} DMSO	0.4909	0.3476	0.5631	0.9840	0.9204	0.6795		
S _{0.5} DMSO	0.8164	0.5106	0.6978	0.6611	1.0431	1.0406	0.4599	
Form III	27.6893	28.0526	27.8906	27.0483	27.0544	27.1303	27.2600	27.0526

5. DMSO Hemisolvate

The .res file of the computationally generated $S_{0.5DMSO}$ is given below.

```
TITL DMSO_HemiSolvate_PBE-TS
CELL 1.54180 8.4035 16.0552 19.5606 90.000 97.567 90.000
ZERR 4 0.0000 0.0000 0.0000 0.000 0.000 0.000
LATT 1
SYMM 0.50000 - X , 0.50000 + Y , 0.50000 - Z
SFAC C H N O S
C1 1 0.15112 0.04821 0.14579 11.00000 0.0500
C2 1 0.24623 -0.05913 0.22767 11.00000 0.0500
C3 1 0.26963 0.01615 0.19621 11.00000 0.0500
C4 1 0.10167 -0.10431 0.21055 11.00000 0.0500
C5 1 -0.01411 -0.07237 0.15802 11.00000 0.0500
C6 1 0.01054 0.00258 0.12609 11.00000 0.0500
C7 1 0.35141 0.13047 0.06345 11.00000 0.0500
C8 1 0.33396 0.10563 -0.00580 11.00000 0.0500
C9 1 0.46830 0.09270 -0.03825 11.00000 0.0500
C10 1 0.62397 0.10404 -0.00229 11.00000 0.0500
C11 1 0.63946 0.12823 0.06771 11.00000 0.0500
C12 1 0.50506 0.14085 0.10011 11.00000 0.0500
C13 1 0.71410 0.06246 0.31876 11.00000 0.0500
C14 1 0.65154 0.04362 0.38005 11.00000 0.0500
C15 1 0.60671 -0.03718 0.39278 11.00000 0.0500
C16 1 0.62313 -0.10180 0.34501 11.00000 0.0500
C17 1 0.68791 -0.08169 0.28386 11.00000 0.0500
C18 1 0.73319 -0.00084 0.27116 11.00000 0.0500
C19 1 0.97359 0.17650 0.34446 11.00000 0.0500
C20 1 1.10139 0.16589 0.30588 11.00000 0.0500
C21 1 1.25645 0.18043 0.33599 11.00000 0.0500
C22 1 1.28895 0.20625 0.40559 11.00000 0.0500
C23 1 1.15915 0.21181 0.44473 11.00000 0.0500
C24 1 1.00329 0.19835 0.41445 11.00000 0.0500
C25 1 0.30579 0.50540 0.06690 11.00000 0.0500
C26 1 0.48234 0.36639 0.08849 11.00000 0.0500
H1 2 0.16267 -0.20374 0.27656 11.00000 -1.20000
H2 2 0.74638 0.08203 -0.08563 11.00000 -1.20000
H3 2 -0.02086 -0.21270 0.22376 11.00000 -1.20000
H4 2 0.86671 0.11147 -0.00998 11.00000 -1.20000
H5 2 0.33834 -0.08389 0.26673 11.00000 -1.20000
H6 2 0.38143 0.05002 0.21076 11.00000 -1.20000
H7 2 -0.12392 -0.10795 0.14341 11.00000 -1.20000
H8 2 -0.07943 0.02607 0.08551 11.00000 -1.20000
H9 2 0.21410 0.09737 -0.03372 11.00000 -1.20000
H10 2 0.45551 0.07354 -0.09219 11.00000 -1.20000
H11 2 0.75840 0.13815 0.09609 11.00000 -1.20000
H12 2 0.52041 0.16003 0.15398 11.00000 -1.20000
H13 2 0.51106 -0.19278 0.39656 11.00000 -1.20000
H14 2 0.59373 -0.22789 0.32401 11.00000 -1.20000
H15 2 1.46977 0.24145 0.48388 11.00000 -1.20000
H16 2 1.53326 0.21969 0.40431 11.00000 -1.20000
H17 2 0.63926 0.09154 0.41860 11.00000 -1.20000
```

H18	2	0.55959	-0.05272	0.44052	11.00000	-1.20000
H19	2	0.70315	-0.13102	0.24694	11.00000	-1.20000
H20	2	0.78385	0.01381	0.22413	11.00000	-1.20000
H21	2	1.07862	0.14801	0.25185	11.00000	-1.20000
H22	2	1.35554	0.17337	0.30583	11.00000	-1.20000
H23	2	1.18304	0.22949	0.49890	11.00000	-1.20000
H24	2	0.90395	0.20589	0.44462	11.00000	-1.20000
H25	2	0.34067	0.50542	0.01474	11.00000	-1.50000
H26	2	0.39704	0.53589	0.10391	11.00000	-1.50000
H27	2	0.19061	0.53724	0.06721	11.00000	-1.50000
H28	2	0.51807	0.38324	0.03831	11.00000	-1.50000
H29	2	0.48580	0.29867	0.09555	11.00000	-1.50000
H30	2	0.56016	0.39626	0.13097	11.00000	-1.50000
N1	3	0.07285	-0.17549	0.24424	11.00000	0.0500
N2	3	0.75729	0.09145	-0.03360	11.00000	0.0500
N3	3	0.58158	-0.18134	0.35868	11.00000	0.0500
N4	3	1.44068	0.22518	0.43331	11.00000	0.0500
O1	4	0.04265	0.16136	0.05675	11.00000	0.0500
O2	4	0.22822	0.20464	0.16153	11.00000	0.0500
O3	4	0.78156	0.17354	0.22926	11.00000	0.0500
O4	4	0.67637	0.22171	0.33651	11.00000	0.0500
O5	4	0.17190	0.35768	0.03540	11.00000	0.0500
S1	5	0.18442	0.14393	0.10669	11.00000	0.0500
S2	5	0.77843	0.16401	0.30332	11.00000	0.0500
S3	5	0.27995	0.39944	0.09342	11.00000	0.0500
END						