

Supporting information

Polymorphism of R-encencline hydrochloride: access to the highest number of structurally characterized polymorphs using desolvation of various solvates

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Table of Contents SI

1. Solvate screening SI
2. Thermal Characterization and VT-PXRD. SI
3. Monohydrate MH-V and MH-VI characterization
4. Crystal structure determination SI
5. Conformation analysis SI
6. Packing analysis SI
7. Thermal characterization SI

SI 1. Solvate screening

Table S1. Solvates obtained in solid form screening, their lattice parameters and grouping according to identified isomorphism/isostructurality.

type	solvent	space group	a. Å	b. Å	c. Å	β. °	V. Å ³	stoichiomety	desolvation product
1.	isobutanol	P_{212121}	26.53	12.16	6.69	90	2160	1:1	V _D
	chloroform		26.08	12.15	6.69		2119	1:1	V _D
	trifluoroethanol		26.23	12.14	6.70		2136	1:1	V _D
2.	2-methoxyethanol	P_{212121}	7.04	13.08	22.81	90	2100	1:1	IV _D
	ethyl acetate		6.98	12.87	22.66		2040	1:1	IV _D
	dichloromethane		7.01	12.72	22.73		2028	1:1	IV _D
	nitromethane		6.96	12.91	22.38		2006	1:1	IV _D
	n- propanol		7.00	13.09	22.83		2087	1:1	IV _D
	n- butanol		7.07	12.82	23.05		2087	1:1	IV _D
	isopropanol		7.03	12.99	23.03		2095	1:1	IV _D
	n- pentanol		7.02	12.93	22.95		2084	1:1	IV _D
	cyclohexanol		7.04	12.73	22.83		2088	1:1	IV _D
	tetrahydrofuran		7.02	12.98	23.14		2109	1:1	IV _D
	1,3-dioxolane		7.03	13.01	22.82		2084	1:1	IV _D
	ethanol		6.96	12.98	22.63		2047	1:1	IV _D
	methanol		6.98	13.10	22.58		2068	1:1	IV _D
	acetone		6.98	13.00	22.70		2063	1:1	IV _D
	pyridine		7.00	12.81	22.98		2064	1:1	IV _D
	benzene		7.09	13.07	23.88		2214	1:1	IV _D
3.	dimethylacetamide	P_{21}	13.67	7.39	11.34	90.2	1110	1:1	IV _D
	N-methyl-2 pyrrolidone		13.92	7.35	11.39	90.3	1167	1:1	IV _D
	acetic acid		13.96	7.05	11.39	91.0	1119	1:2	VIII _D / IX _D
	propionic acid		14.49	7.02	11.64	90.7	1185	1:2	amorphous
Other	acetonitrile	P_{21}	13.33	7.55	10.45	112.3	972	1:1	IV _D /VI _D
	acetonitrile (5 °C)	P_{212121}	9.33	34.97	6.81	90	2224	1:2	S _{ACN-II}
	acetonitrile ^a	P_{212121}	9.47	29.42	6.99	90	1945	1:1	IV _D /XII _D
	formic acid	P_{21}	14.21	7.05	12.38	114.5	1132	1:2	S _{FA}
	formic acid ^a	P_{212121}	28.50	19.06	6.98	90	3797	1:1	VII _D /XI _D
	p-xylene	P_{21}	17.09	7.28	17.54	90.2	2206	1:0.5	IV _D
	cyclohexanol	P_{21}	13.92	12.07	6.84	90.8	1153	1:1	IV _D
	benzyl alcohol	P_{212121}	18.87	17.76	7.02	90	2352	1:1	IV _D

^a obtained in solid state transition from higher solvate

SI 2. Thermal Characterization and VT-PXRD.

Table S2. Thermal data (DSC, TGA) for the Enc-HCl solvates

	guest (T _{peak}), °C	loss loss, %	stoichiometry	thermal desolvation product
S _{IIBA}	138	16.5	1:1	V _D
DS _{ACN}	79	18.5	2:1	IV _D
S _{ACN-I}	84	9.8	1:1	IV _D
S _{ACN-II}	72	10.6	1:1	IV _D
S _{FA}	120	11.5	1:1	VII _D
DS _{FA}	60; 120	24.0	2:1	S _{FA} / VII _D
DS _{AA}	115	21.6	2:1	IXD
S _{BnOH}	123	24.9	1:1	IV _D
S _{EtOH}	99	11.2	1:1	IV _D

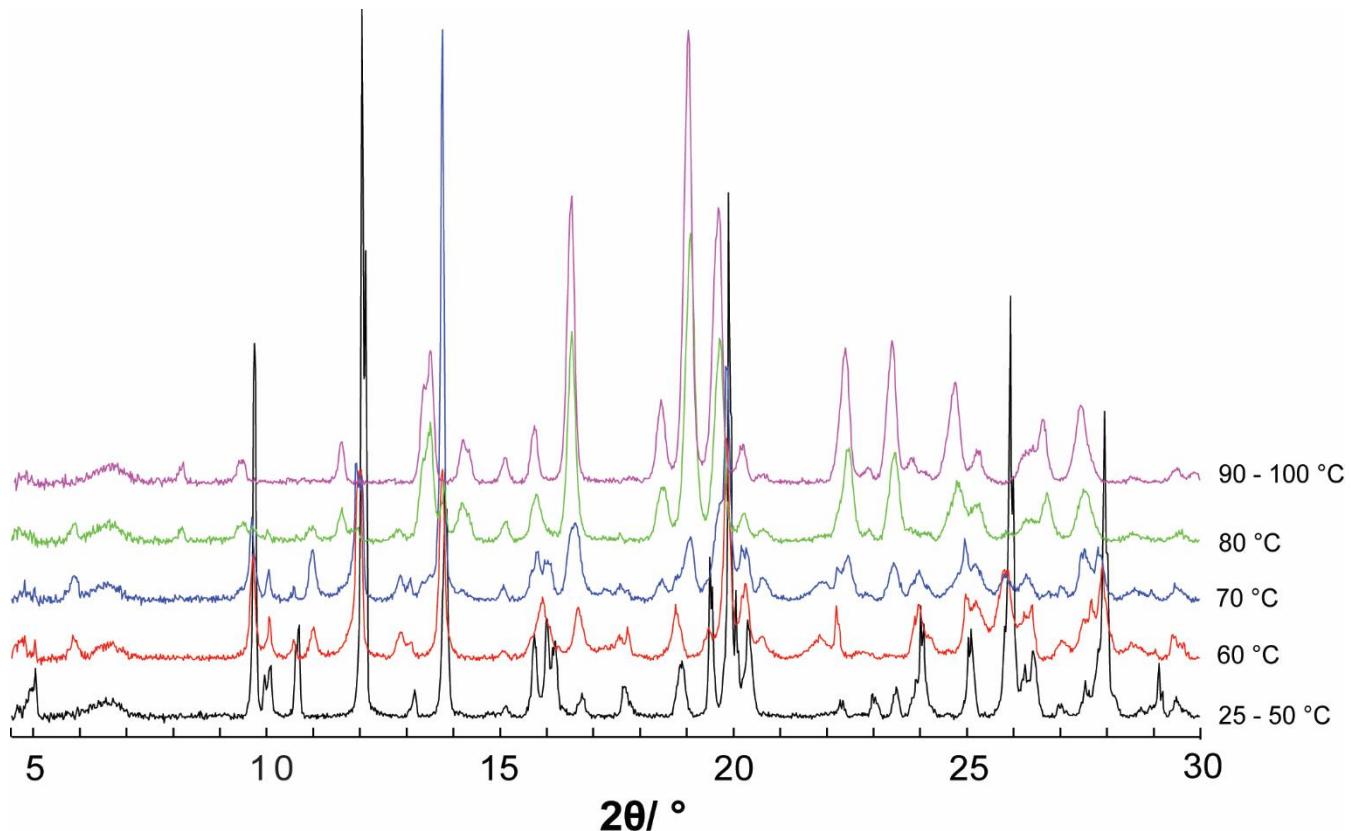


Figure S1. VT-PXRD patterns of Enc-HCl acetonitrile disolvate- DS_{ACN}.

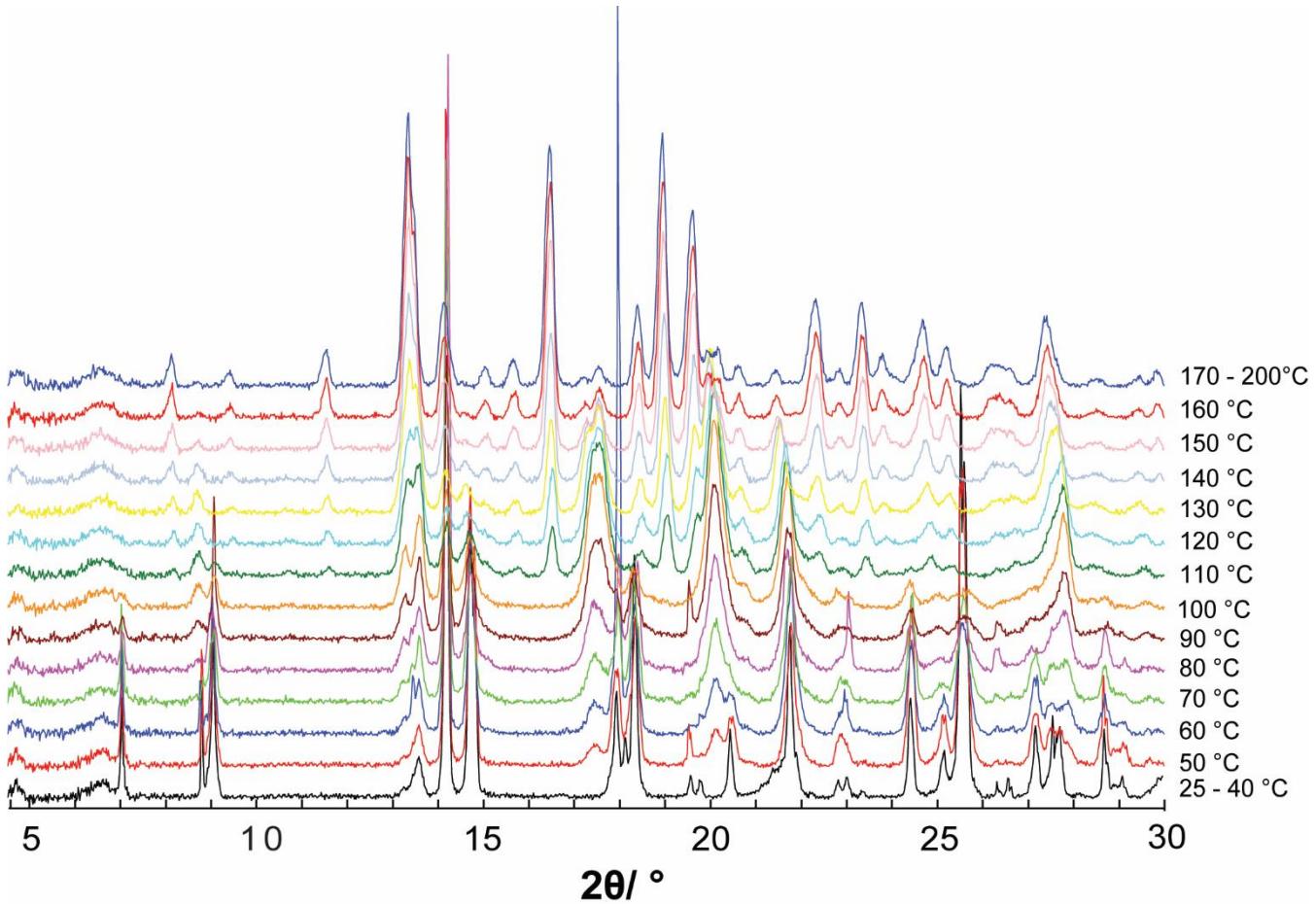


Figure S2. VT-PXRD patterns of Enc-HCl acetonitrile monosolvate- $S_{\text{ACN-I}}$.

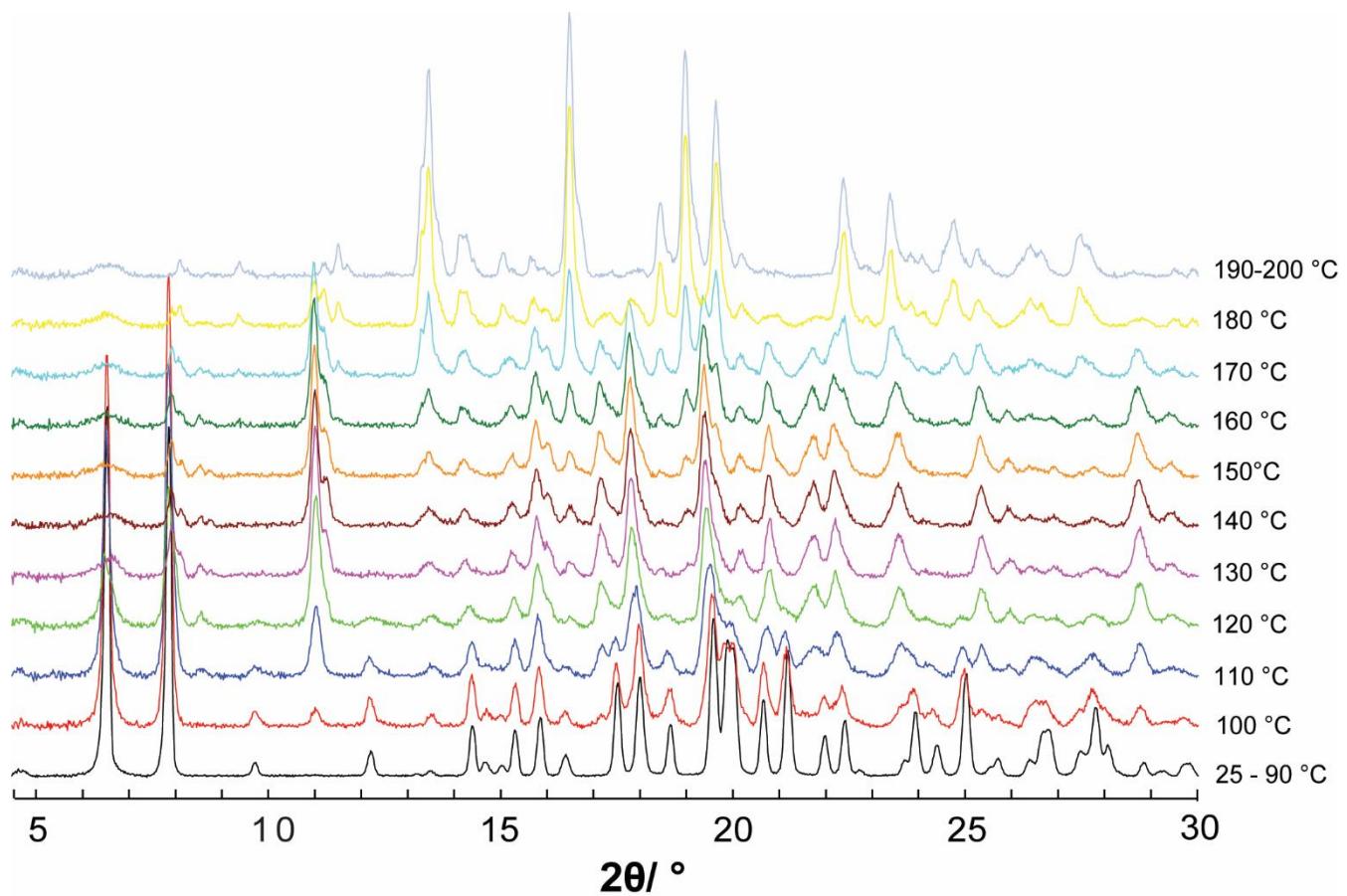


Figure S3. VT-PXRD patterns of Enc-HCl isobutanol monosolvate- S_{IBA} .

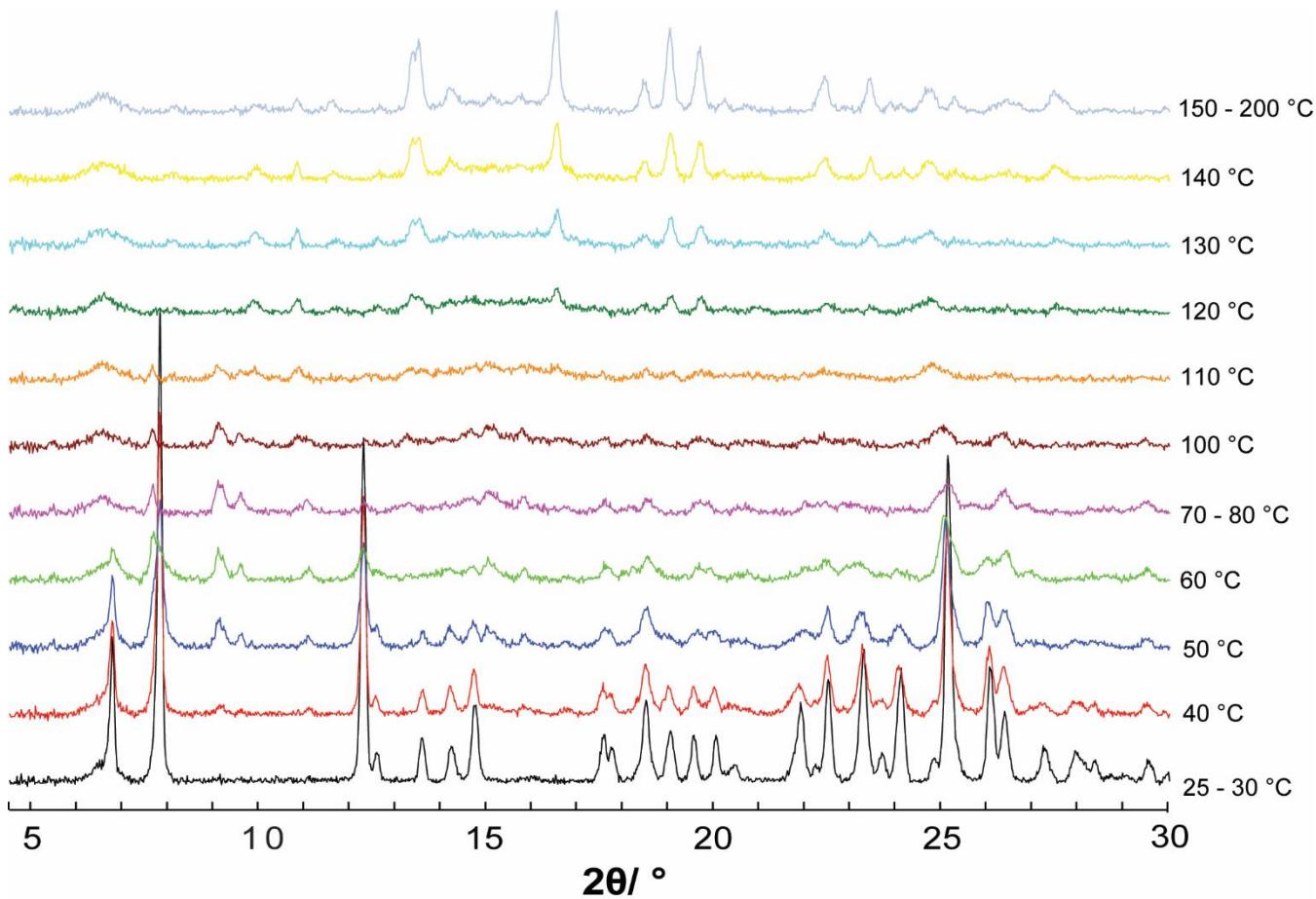


Figure S4. VT-PXRD patterns of Enc-HCl formic acid disolvate- DS_{FA}.

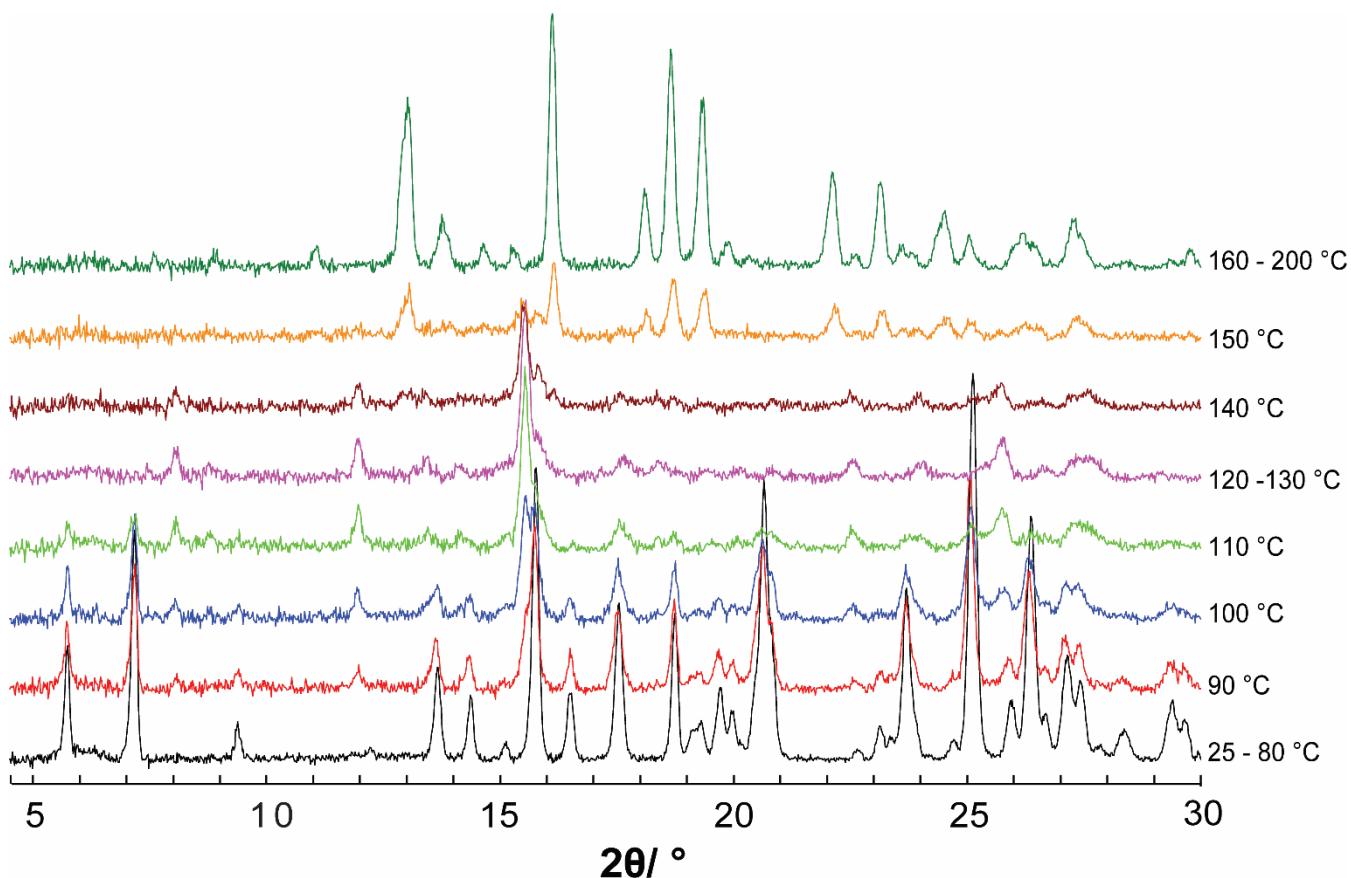


Figure S5. VT-PXRD patterns of Enc-HCl acetic acid disolvate- DS_{AA}.

3. Monohydrate MH-V and MH-VI characterization

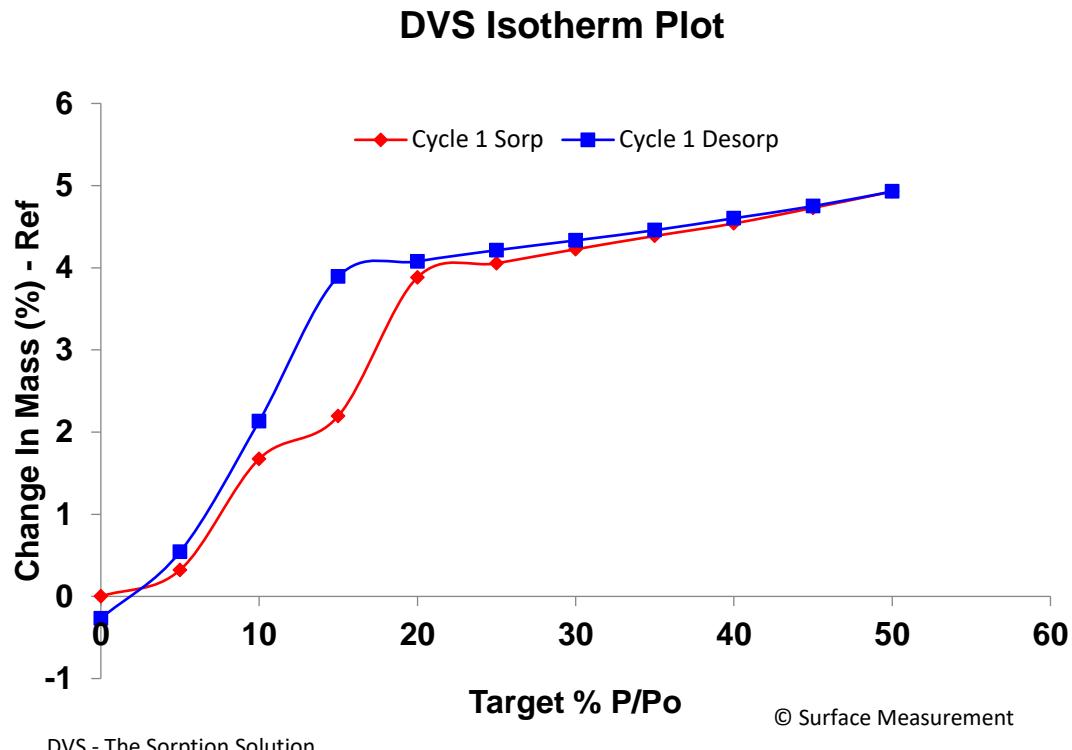


Figure S6. DVS isotherms of Enc-HCl monohydrate MH-V at 25 °C.

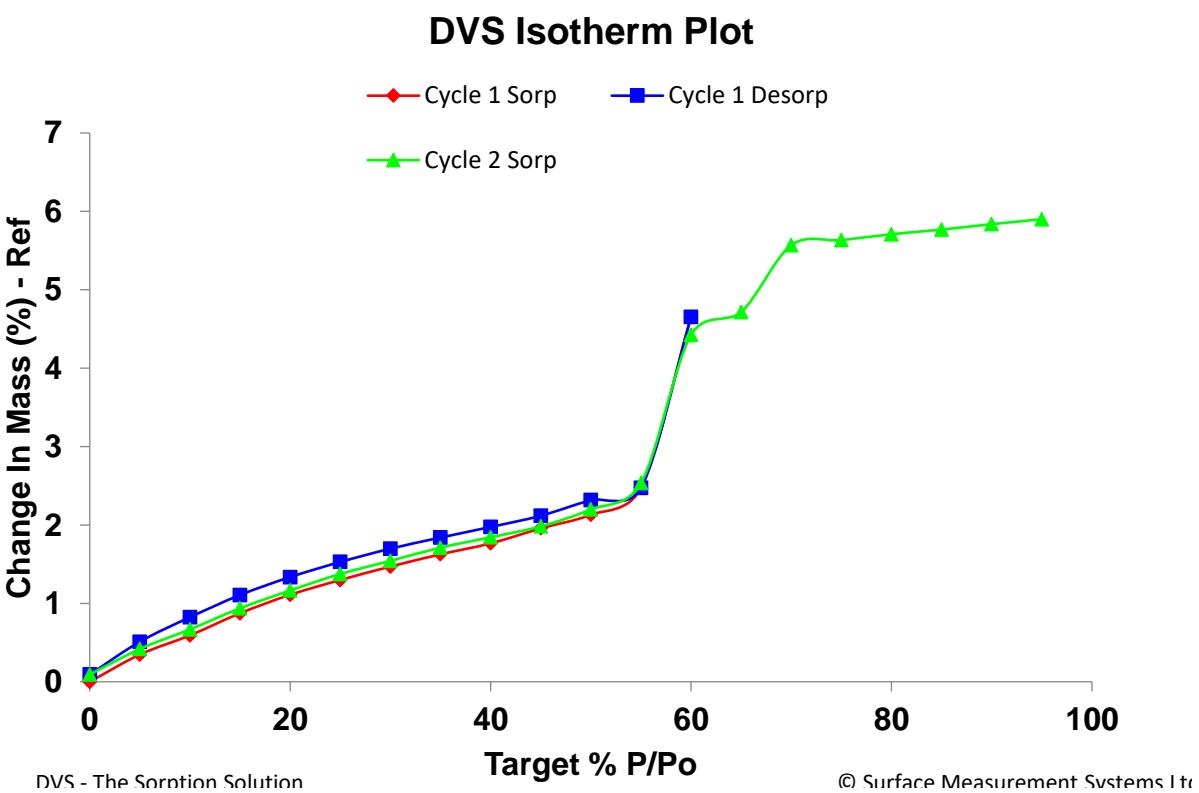


Figure S7. DVS isotherms of Enc-HCl monohydrate MH-VI at 25 °C.

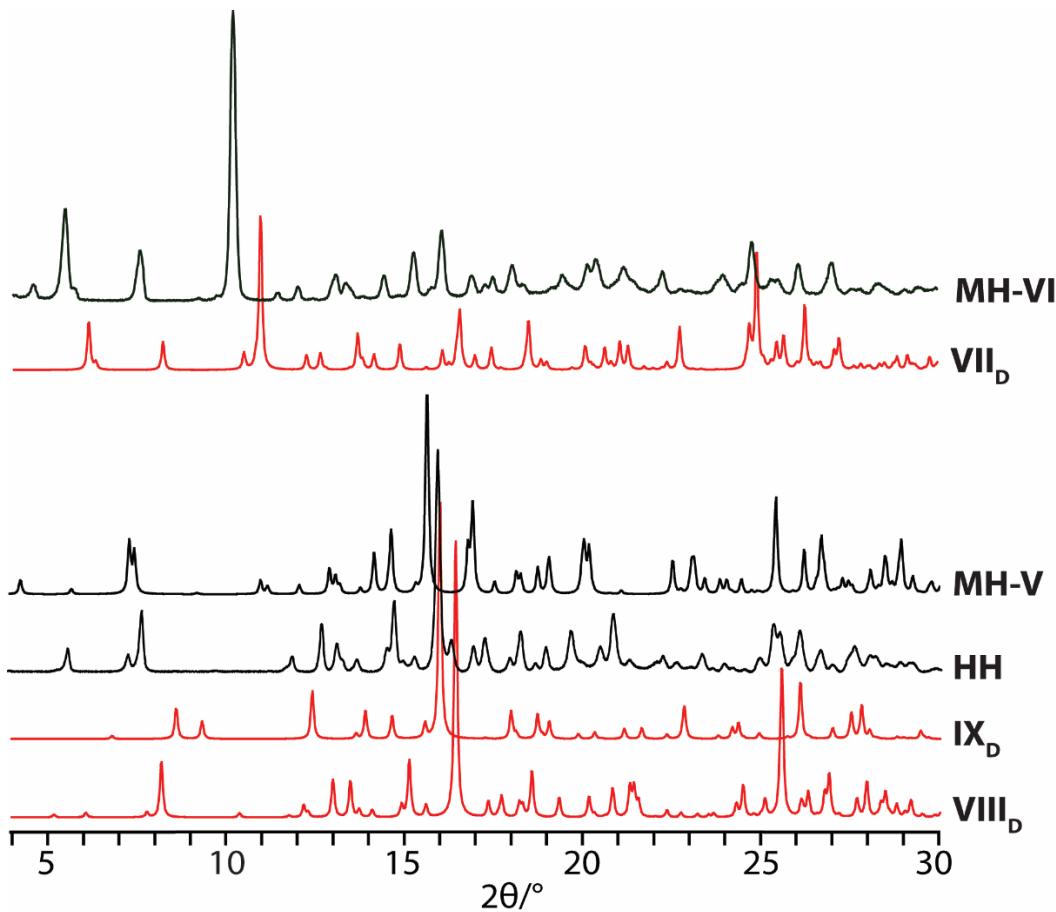


Figure S8. PXRD patterns of Enc-HCl nonsolvated (red) crystalline forms and their respective hydrated phases (black).

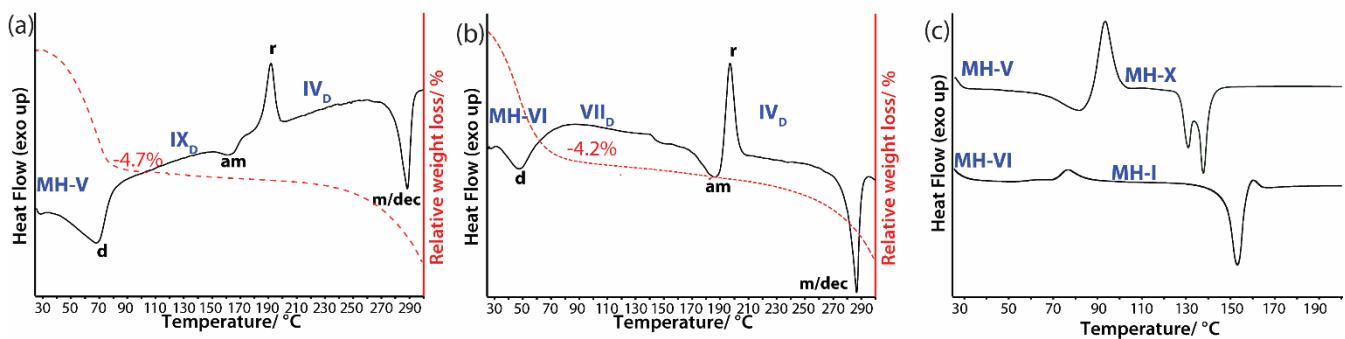


Figure S9. (a-b) DSC/TG traces of Enc-HCl monohydrates. d – desolvation, am – amorphization, r – recrystallization, m/dec – melting with decomposition (10 $^\circ\text{C}/\text{min}$ heating rate) in open crucibles. (c) Overlay of DSC curves of Enc-HCl monohydrates at 10 $^\circ\text{C}/\text{min}$ heating rate in sealed crucibles.

SI 4. Crystal structure determination

Table S3. Crystal data for Enc-HCl solvated phases

	DS_{AA}	S_{EtOH}	DS_{ACN}	S_{ACN-I}	S_{ACN-II}	S_{IBA}	S_{FA}	S_{BnOH}	MH-V
formula	C ₂₀ H ₂₆ Cl ₂ N ₂ O ₅ S	C ₁₈ H ₂₃ Cl ₂ N ₂ O ₂ S	C ₂₀ H ₂₄ Cl ₂ N ₄ OS	C ₁₈ H ₂₁ Cl ₂ N ₃ OS	C ₁₈ H ₂₁ Cl ₂ N ₃ OS	C ₂₀ H ₂₈ Cl ₂ N ₂ O ₂ S	C ₁₇ H ₂₀ Cl ₂ N ₂ O ₃ S	C ₂₃ H ₂₆ Cl ₂ N ₂ O ₂ S	C ₁₆ H ₂₀ Cl ₂ N ₂ O ₂ S
formula weight, g·mol ⁻¹	477.39	403.35	439.39	398.34	398.34	431.40	403.31	465.42	375.30
sample type	crystal	crystal	crystal	crystal	powder	powder	powder	powder	powder
crystal system	monoclinic	orthorhombic	orthorhombic	monoclinic	orthorhombic	orthorhombic	orthorhombic	orthorhombic	monoclinic
space group	P ₂ ₁	P ₂ ₁ 2 ₁ 2 ₁	P ₂ ₁ 2 ₁ 2 ₁	P ₂ ₁	P ₂ ₁ 2 ₁ 2 ₁	P ₂ ₁ 2 ₁ 2 ₁	P ₂ ₁ 2 ₁ 2 ₁	P ₂ ₁ 2 ₁ 2 ₁	P ₂ ₁
a, Å	13.8812(4)	6.9170(2)	6.6818(2)	10.25590(10)	29.424(3)	26.533(16)	28.5929(10)	18.8294(7)	18.3333(8)
b, Å	6.8563(2)	13.0268(5)	9.2833(2)	7.50830(10)	9.4777(10)	12.1649(8)	19.1266(8)	17.7219(7)	6.98761(17)
c, Å	11.4582(4)	21.8980(8)	35.0845(5)	13.3761(2)	6.9991(7)	6.6952(4)	7.0043(2)	7.0016(3)	14.2176(5)
α, °	90	90	90	90	90	90	90	90	90
β, °	90.293(3)	90	90	112.478(2)	90	90	90	90	90.052(2)
γ, °	90	90	90	90	90	90	90	90	90
V, Å ³	1090.51(6)	1973.15(12)	2176.26(9)	951.76(2)	1951.9(3)	2161.0(13)	3830.6(2)	2336.38(16)	1821.05(11)
ρ _{calc} , g·cm ⁻³	1.45	1.36	1.34	1.39	1.36	1.33	1.40	1.32	1.37
Z/ Z'	2/1	4/1	4/1	2/1	4/1	4/1	8/2	4/1	4/2
temperature, K	174	174	174	150	293	293	293	293	293
2θ _{min} –2θ _{max} /increment, °	-	-	-	-	4.5–70/0.01	4.5–70/0.01	5.0–70/0.005	4.0–70/0.01	4.0–70/0.01
Rwp	-	-	-	-	0.04911	0.03790	0.05833	0.04829	0.02597
Rp	-	-	-	-	0.07324	0.02917	0.04338	0.03661	0.03695
R _{exp}	-	-	-	-	0.04460	0.01124	0.05465	0.01239	0.00641
R ₁ (wR ₂)	0.0406 (0.0510)	0.0335 (0.0398)	0.0303 (0.0368)	0.0255 (0.0680)	-	-	-	-	-
Packing index, % ^a	73.5	72.1	73.3	73.6	72.2	72.8	72.2	72.0	68.2
CCDC no.	1895195	1895190	1916268	1916267	1916269	1895198	1895189	1895191	1916266

^a after geometry optimization by relaxing positions of all atoms and unit cell parameters

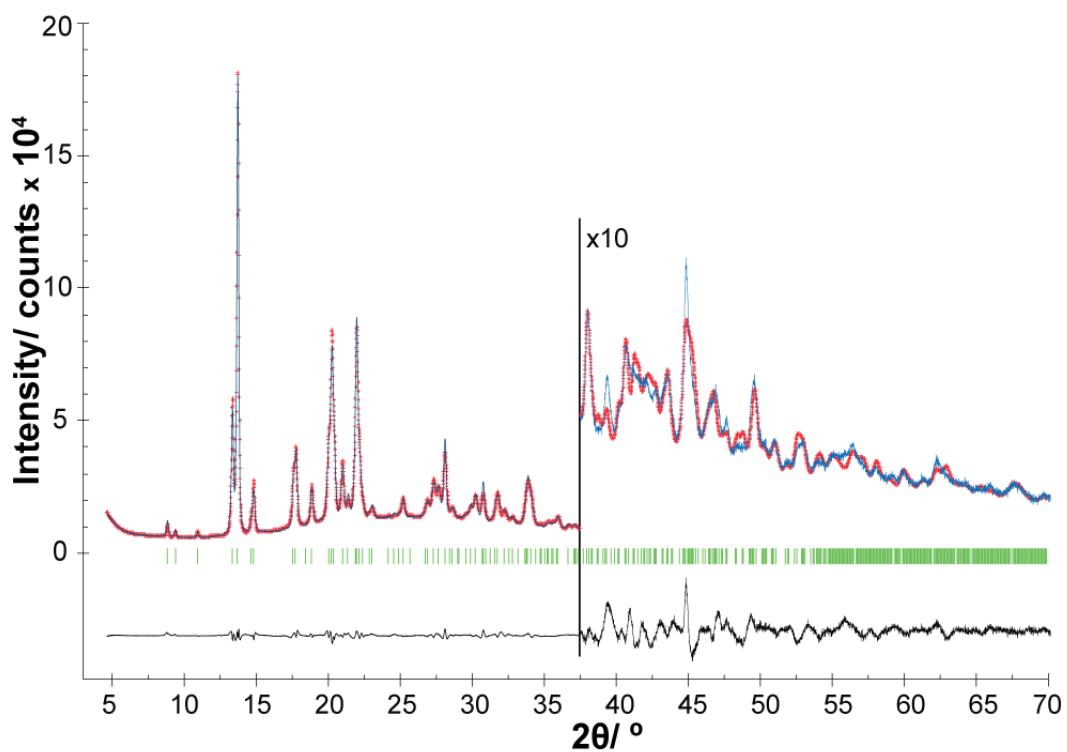


Figure S10. Final Rietveld fit for Enc-HCl polymorph VI_D: red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

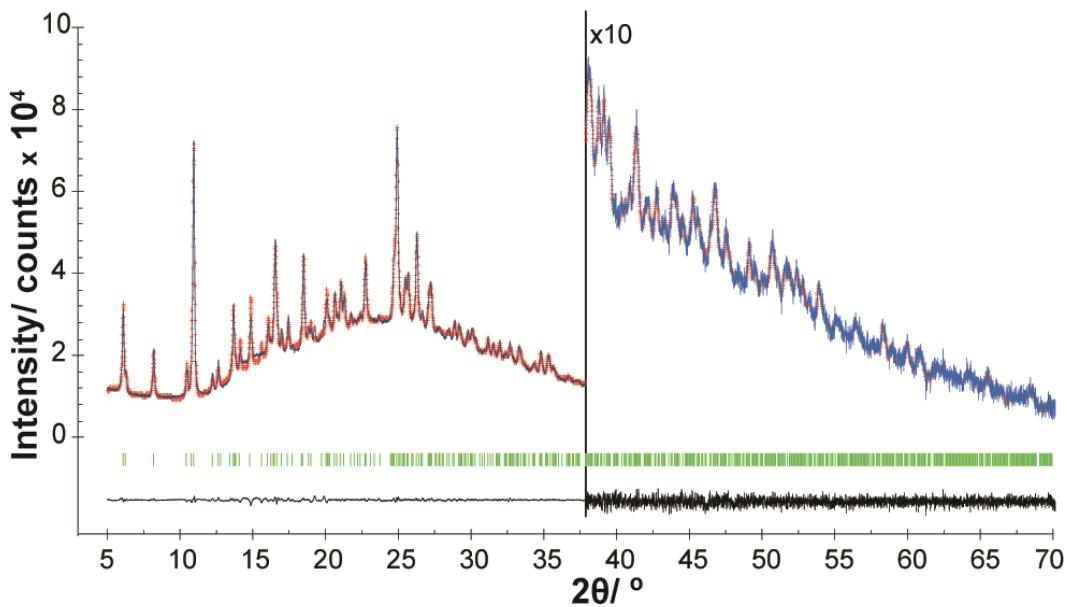


Figure S11. Final Rietveld fit for Enc-HCl polymorph VII_D: red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

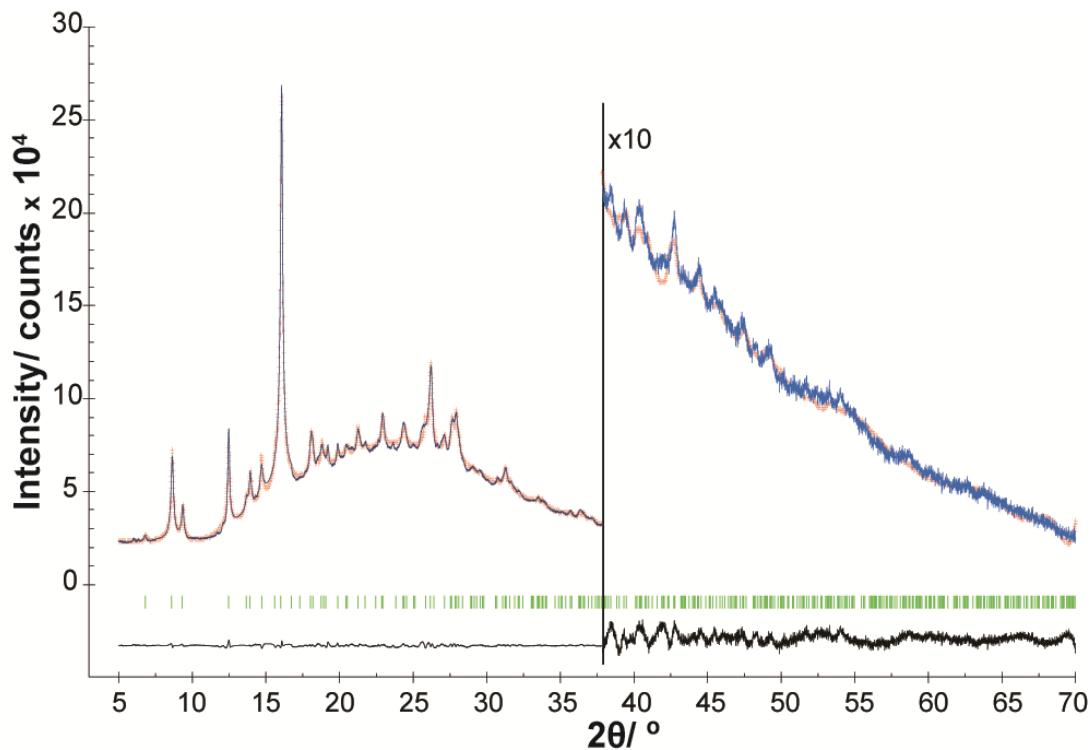


Figure S12. Final Rietveld fit for Enc-HCl polymorph VIII_D : red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

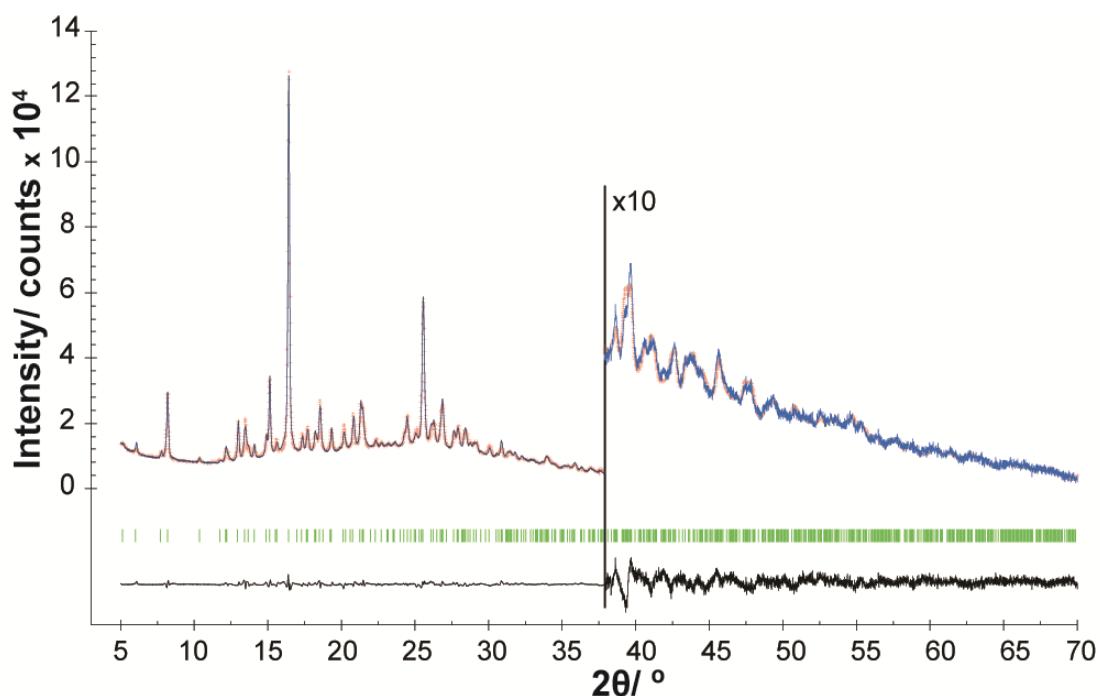


Figure S13. Final Rietveld fit for Enc-HCl polymorph IX_D : red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

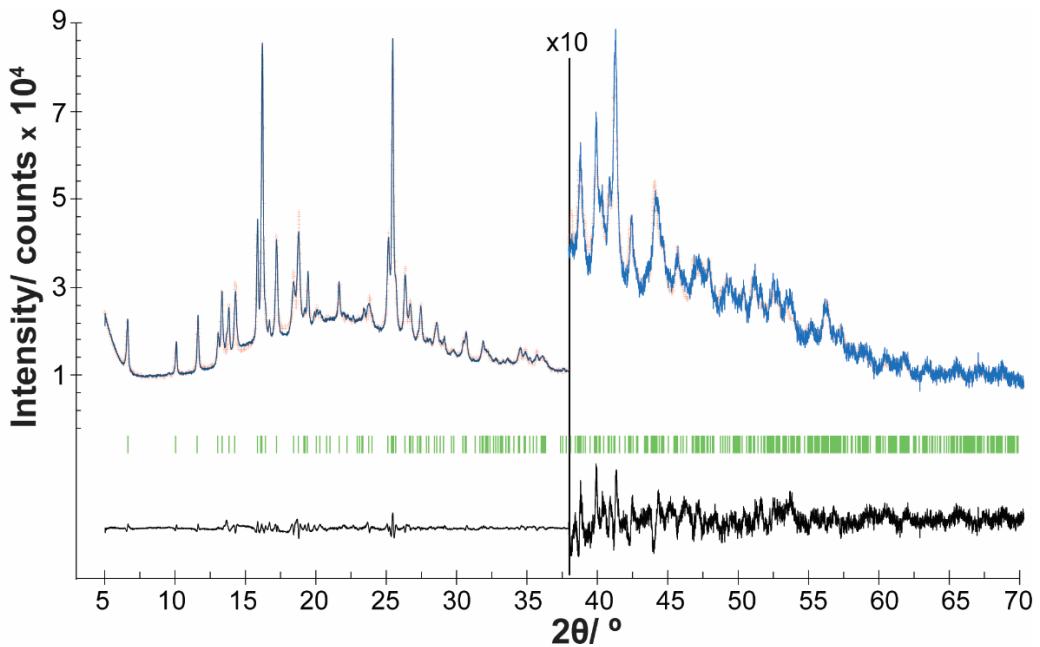


Figure S14. Final Rietveld fit for Enc-HCl polymorph XII_D: red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

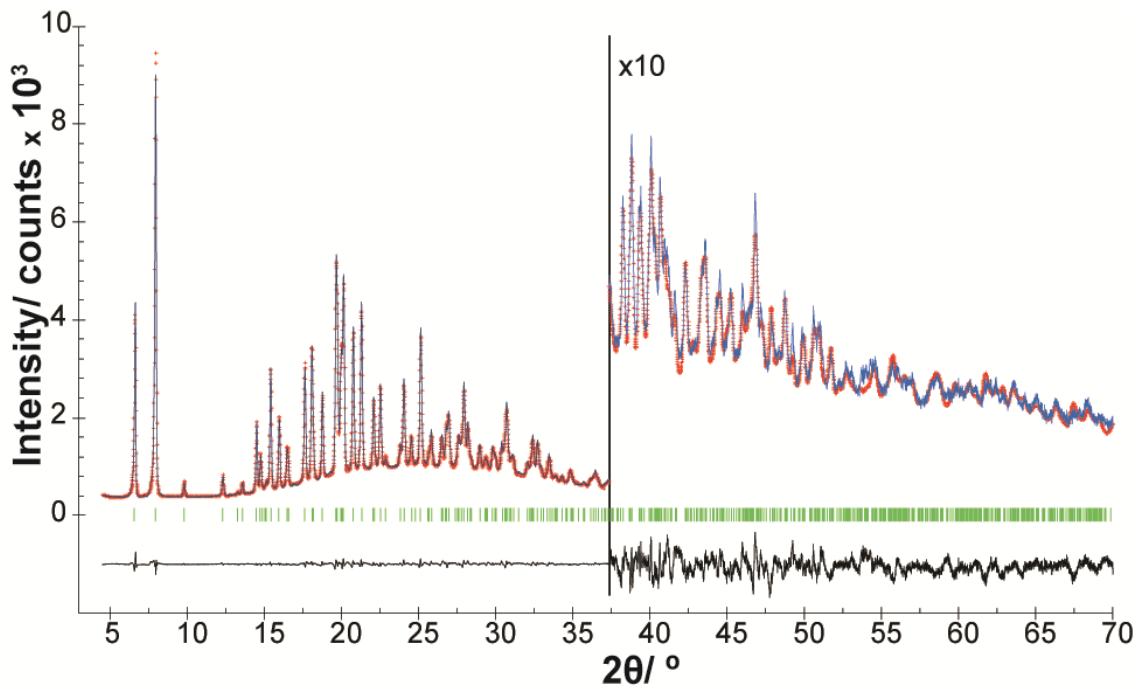


Figure S15. Final Rietveld fit for Enc-HCl i-butanol monosolvate S_{IBA}: red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

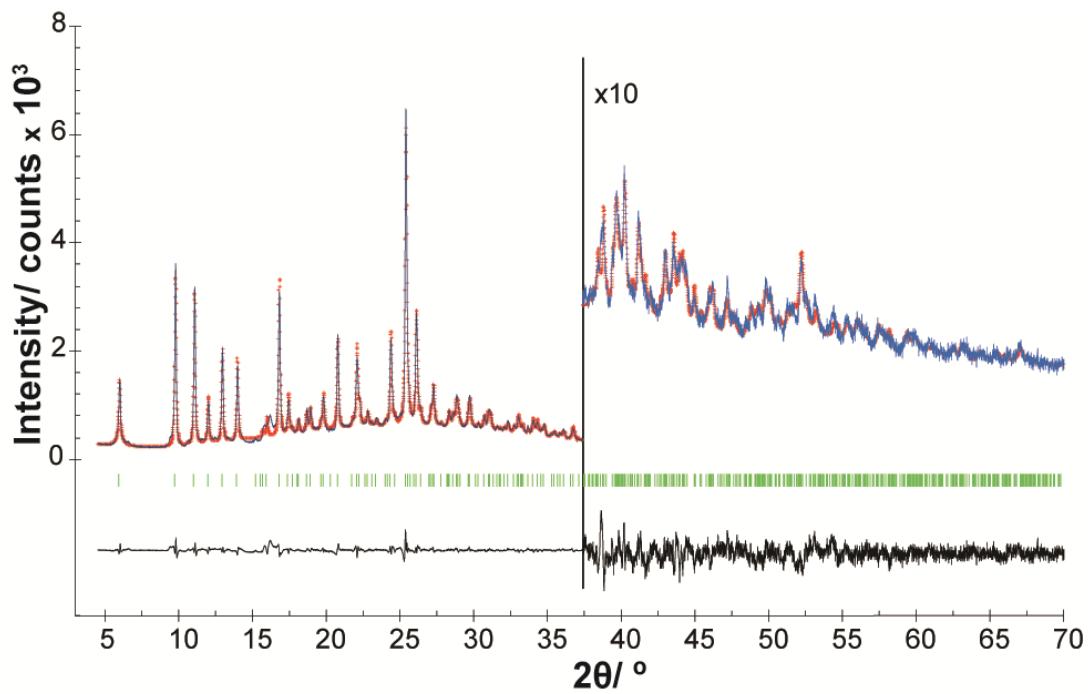


Figure S16. Final Rietveld fit for Enc-HCl acetonitrile monosolvate I $S_{\text{ACN-II}}$: red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

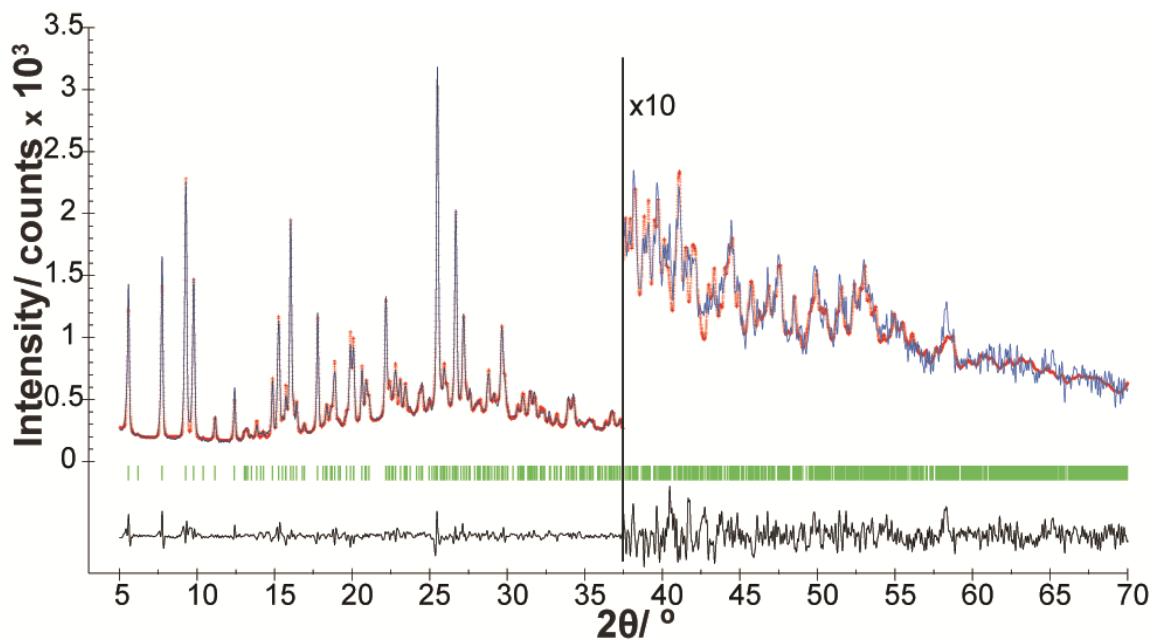


Figure S17. Final Rietveld fit for Enc-HCl formic acid monosolvate S_{FA} : red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

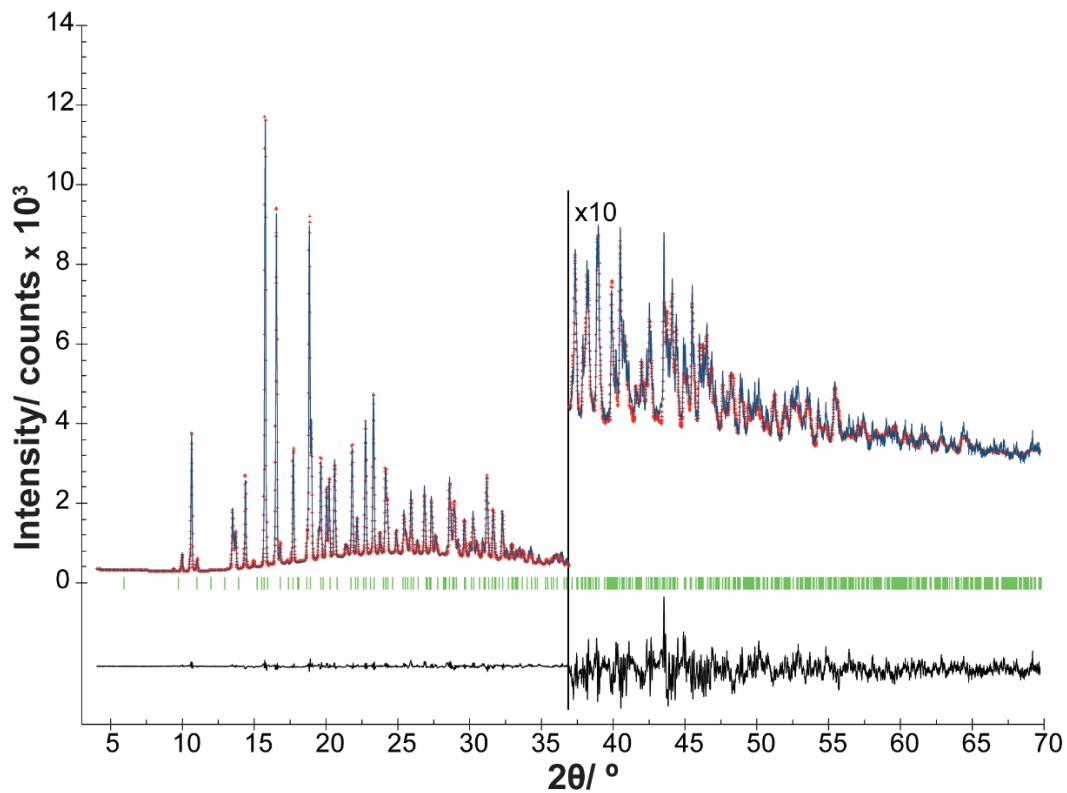


Figure S18. Final Rietveld fit for Enc-HCl benzyl alcohol monosolvate S_{BnOH} : red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

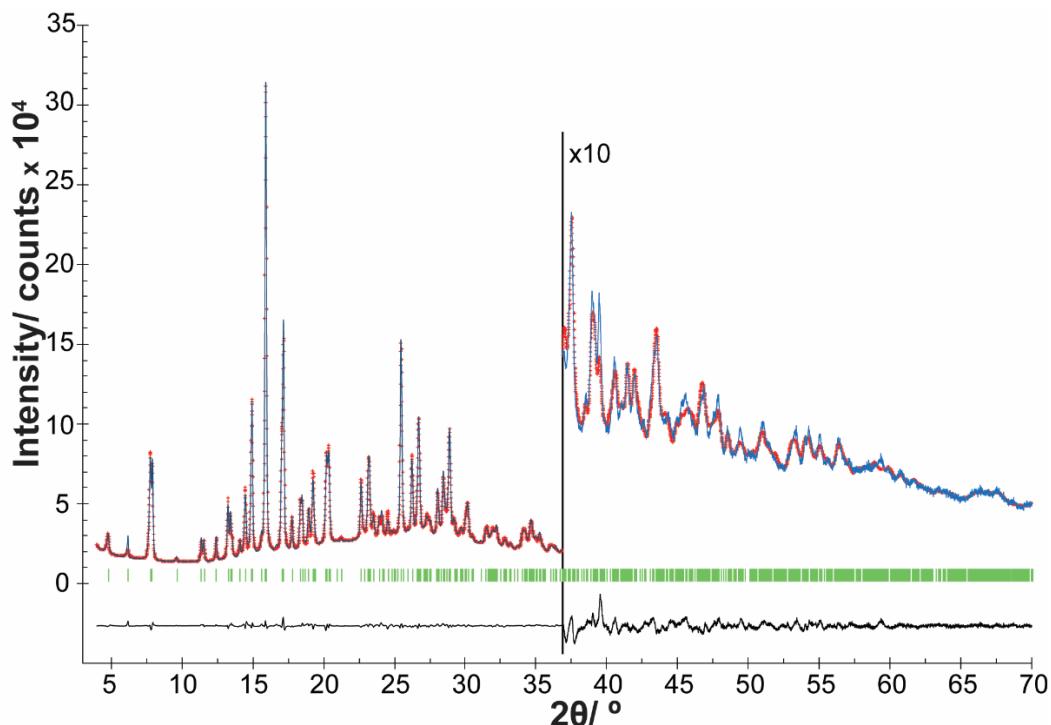


Figure S19. Final Rietveld fit for Enc-HCl monohydrate MH-V: red crosses – measured data points; blue line – calculated profile; black line – difference curve; green tick marks – calculated peak positions.

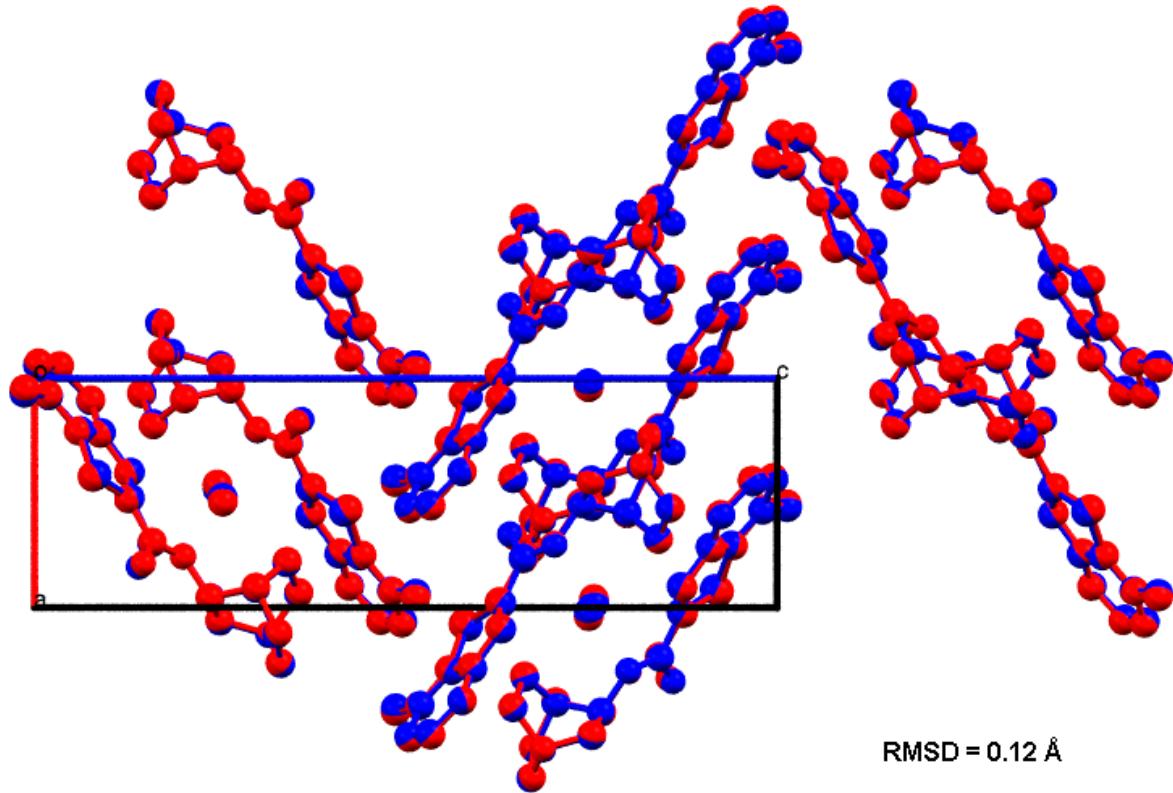


Figure S20. Overlay of form V_D crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD_{15} value for non-hydrogen atoms.

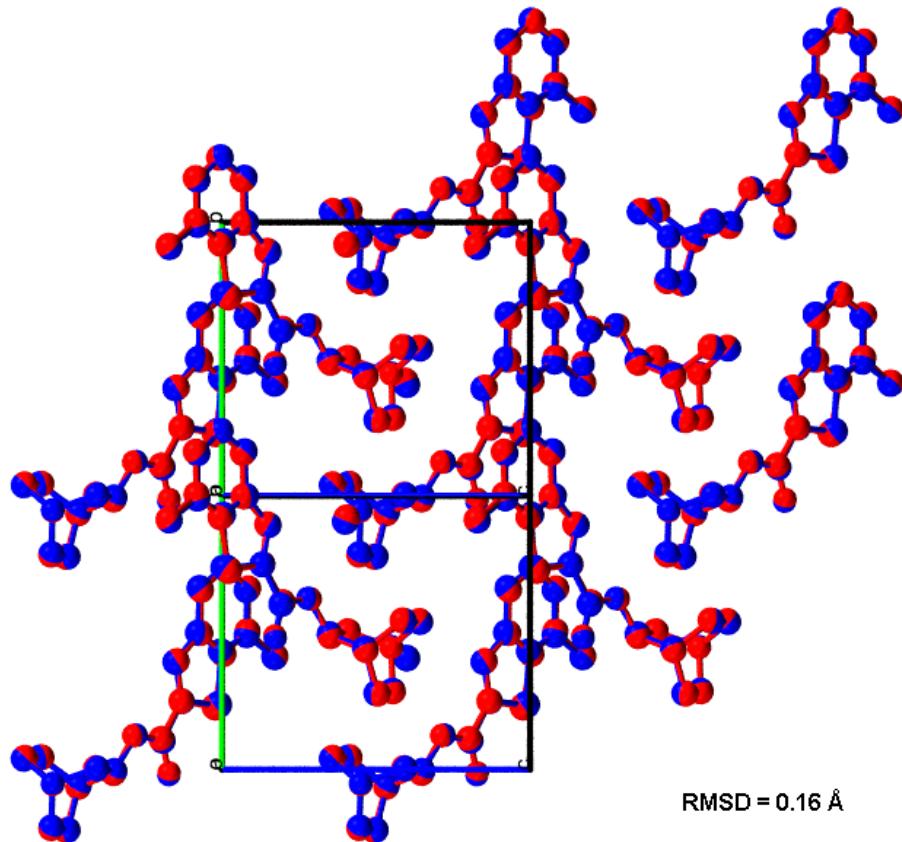


Figure S21. Overlay of form VI_D crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD_{15} value for non-hydrogen atoms.

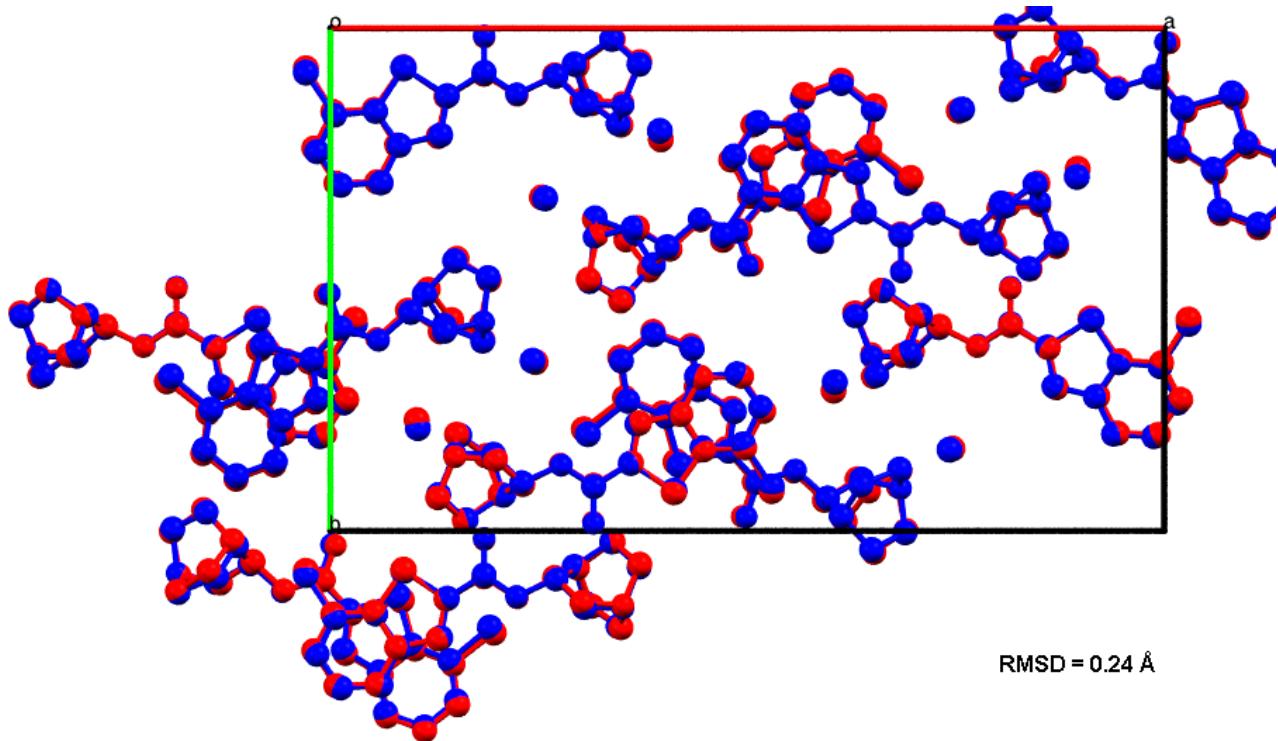


Figure S22. Overlay of form VII_D crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD_{15} value for non-hydrogen atoms.

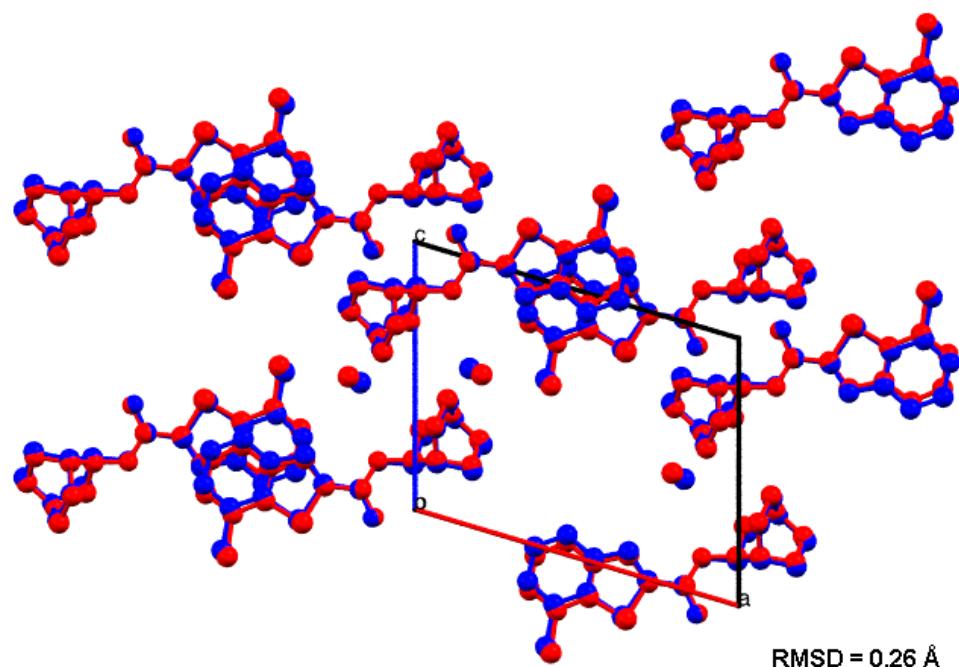


Figure S23. Overlay of form VIII_D crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD_{15} value for non-hydrogen atoms.

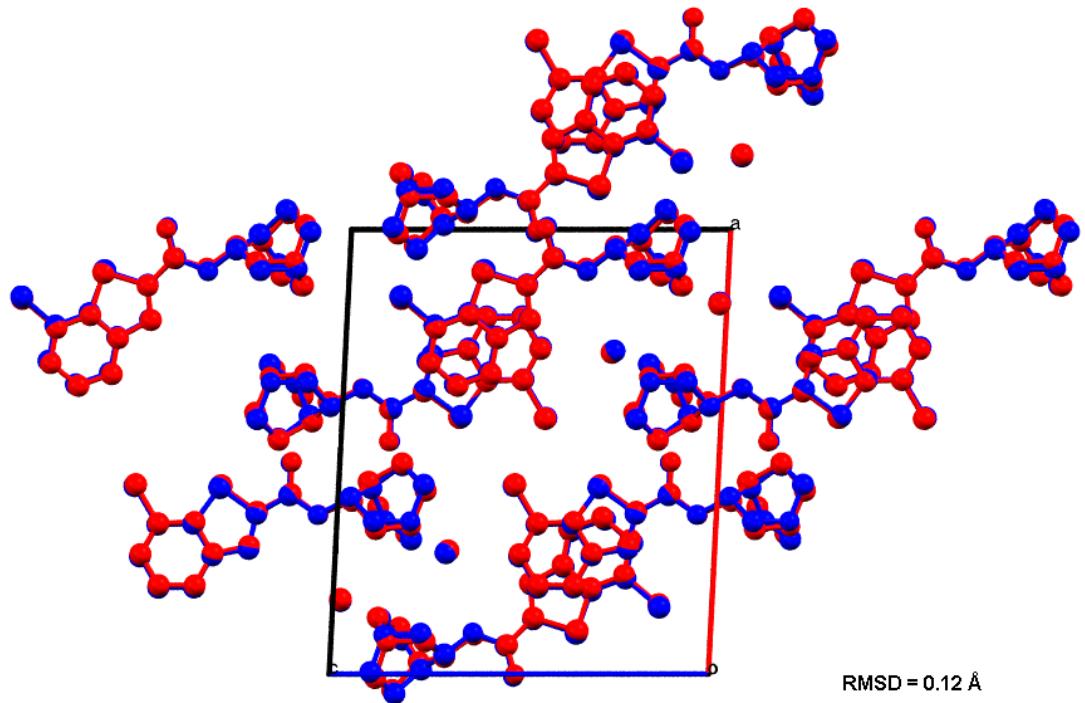


Figure S24. Overlay of form IX_D crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD_{15} value for non-hydrogen atoms.

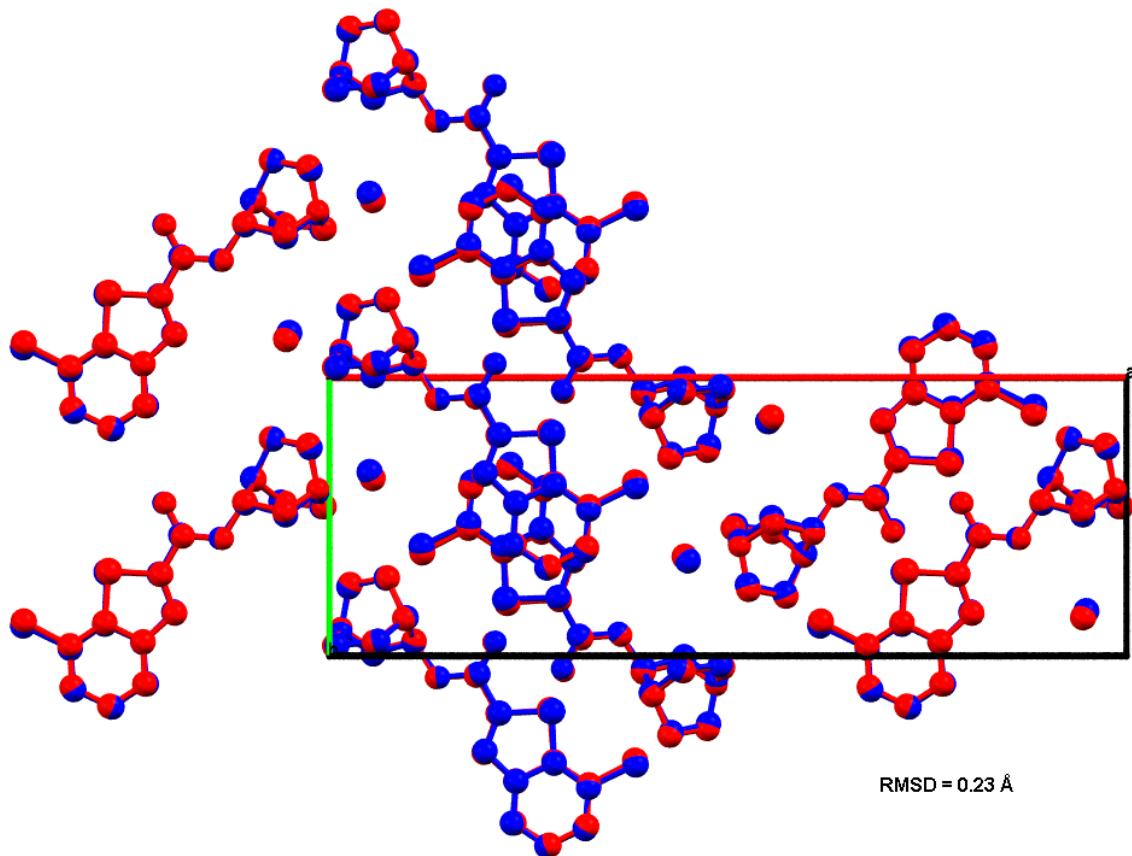


Figure S25. Overlay of form XII_D crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD_{15} value for non-hydrogen atoms.

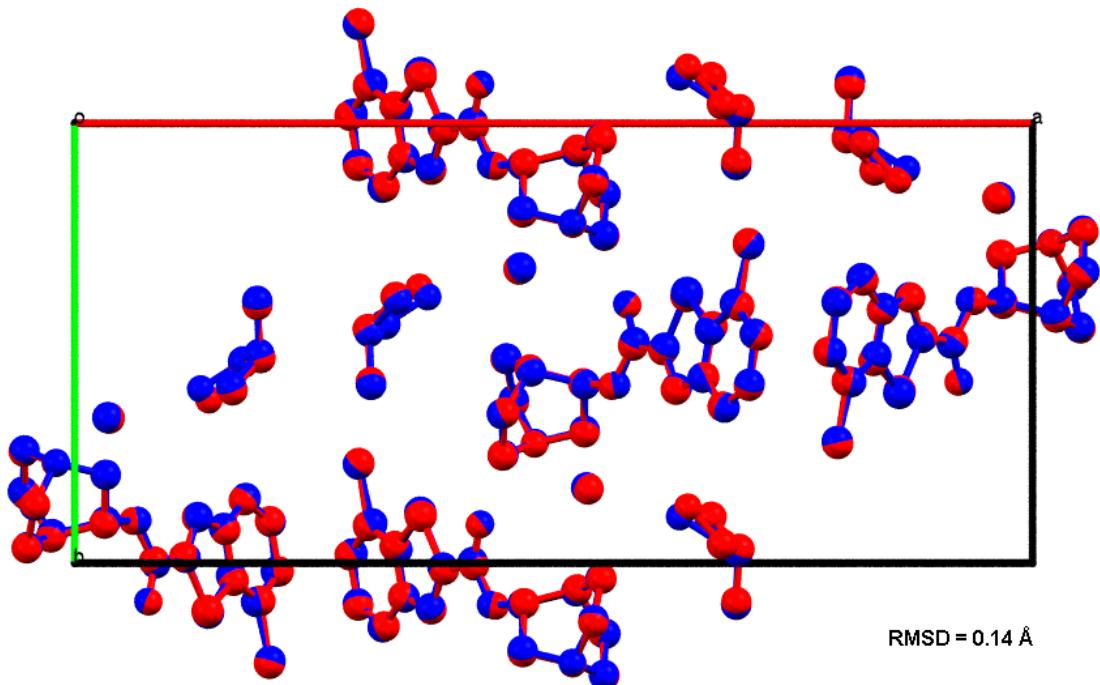


Figure S26. Overlay of form S_IBA crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD_{15} value for non-hydrogen atoms.

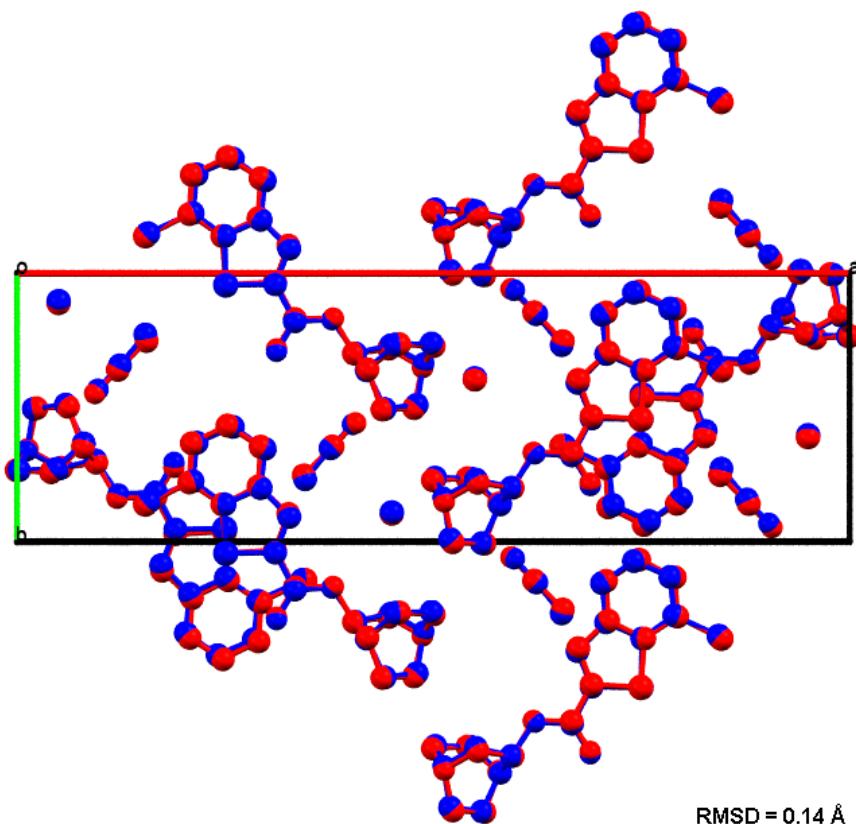


Figure S27. Overlay of form S_{ACN-II} crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD_{15} value for non-hydrogen atoms.

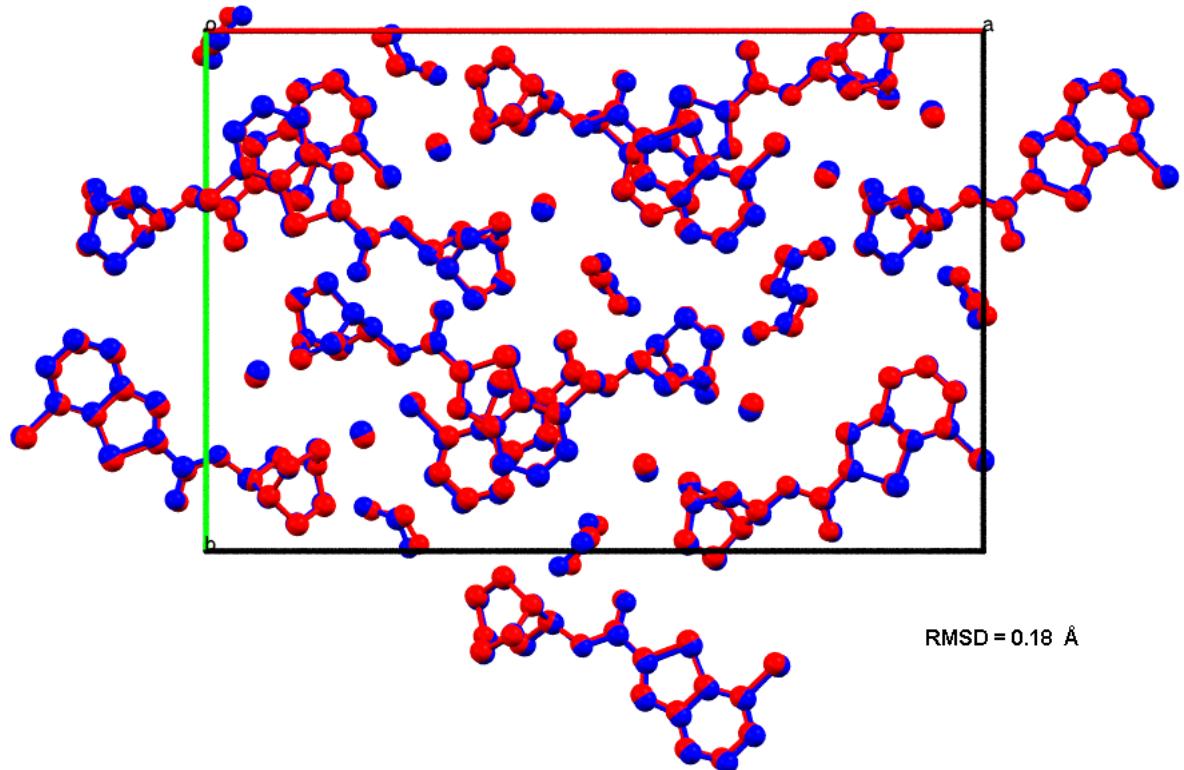


Figure S28. Overlay of form S_{FA} crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD₁₅ value for non-hydrogen atoms.

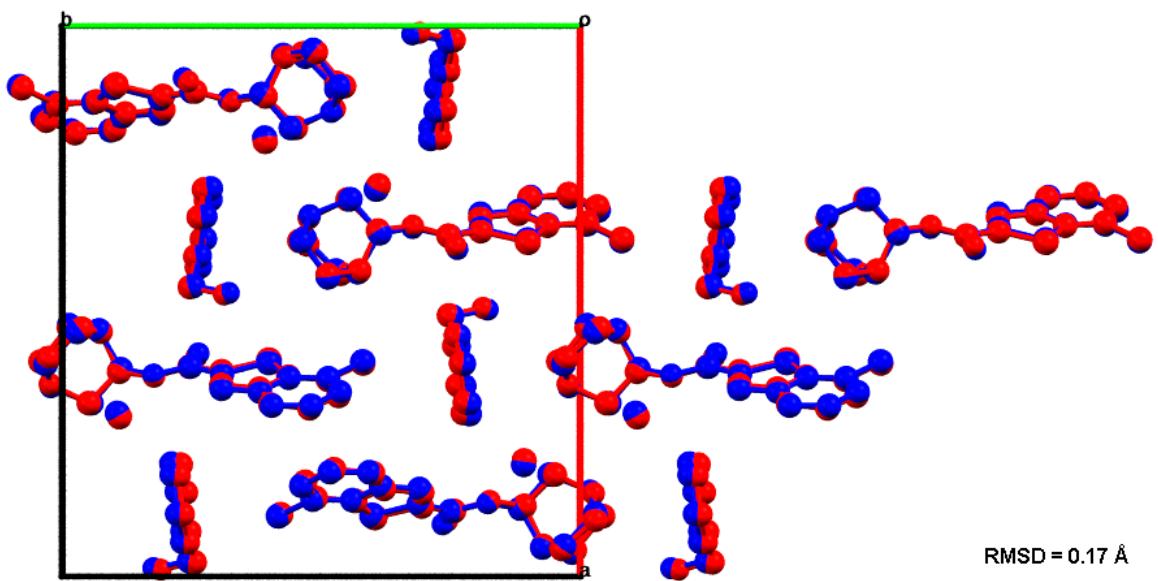


Figure S29. Overlay of form S_{BnOH} crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD₁₅ value for non-hydrogen atoms.

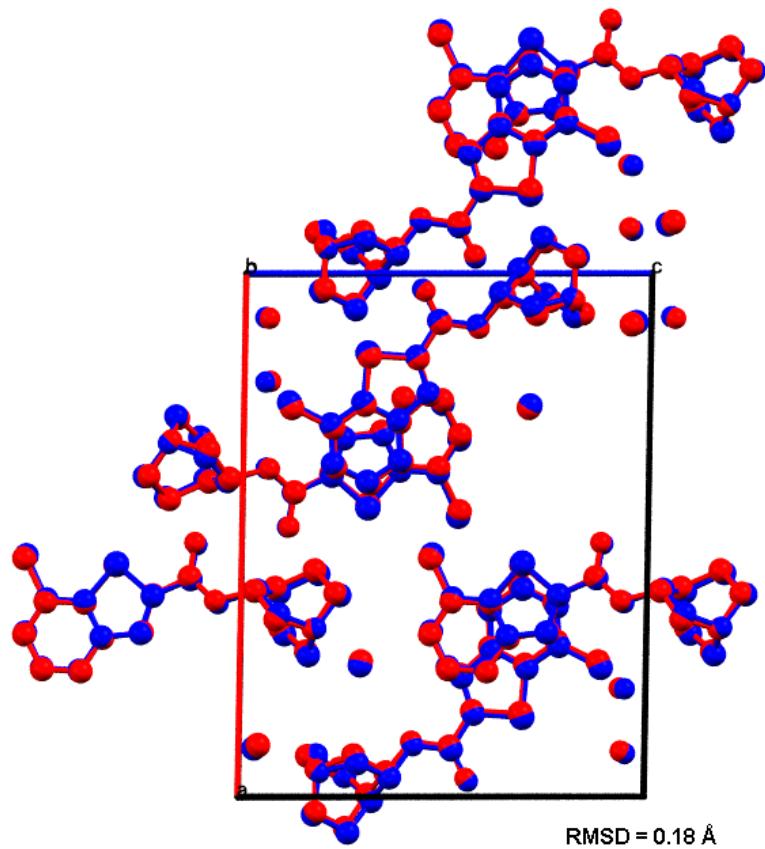


Figure S30. Overlay of form MH-V crystal structure calculated from PXRD data (blue) and crystal structure after DFT optimisation (red) with given RMSD₁₅ value for non-hydrogen atoms.

SI 5. Conformation analysis

Table S4. Enc-HCl cation conformation data.

Phase/ Conformer	τ_1 (C2-C1-C10-O11)/ °	τ_2 (C1-C10-N12-C16)/ °	τ_3 (C10-N12-C16-C15)/ °
IV_D	-160.4	178.7	153.5
V_D	-161.3	173.7	145.3
VI_D	-176.3	179.4	159.9
VII_DA	-173.1	169.3	157.2
VII_DB	-178.6	176.6	164.7
VIII_D	-171.1	167.6	148.2
IX_DA	-178.9	170.7	159.8
IX_DB	-175.7	172.9	148.8
XII_D	-169.1	179.1	166.8
DS_{AA}	-179.2	175.6	143.0
S_{FAA}	-166.9	176.8	142.1
S_{FAB}	-173.1	175.2	155.4
DS_{ACN}	-178.0	175.7	141.8
S_{ACN-I}	-150.8	170.8	144.3
S_{ACN-II}	-178.6	179.2	155.1
S_{IIBA}	-177.8	177.4	153.8
S_{EtoH}	-178.0	176.5	141.7
S_{BnOH}	-173.1	176.8	148.4
Gas phase opt.	-164.9	176.8	160.0

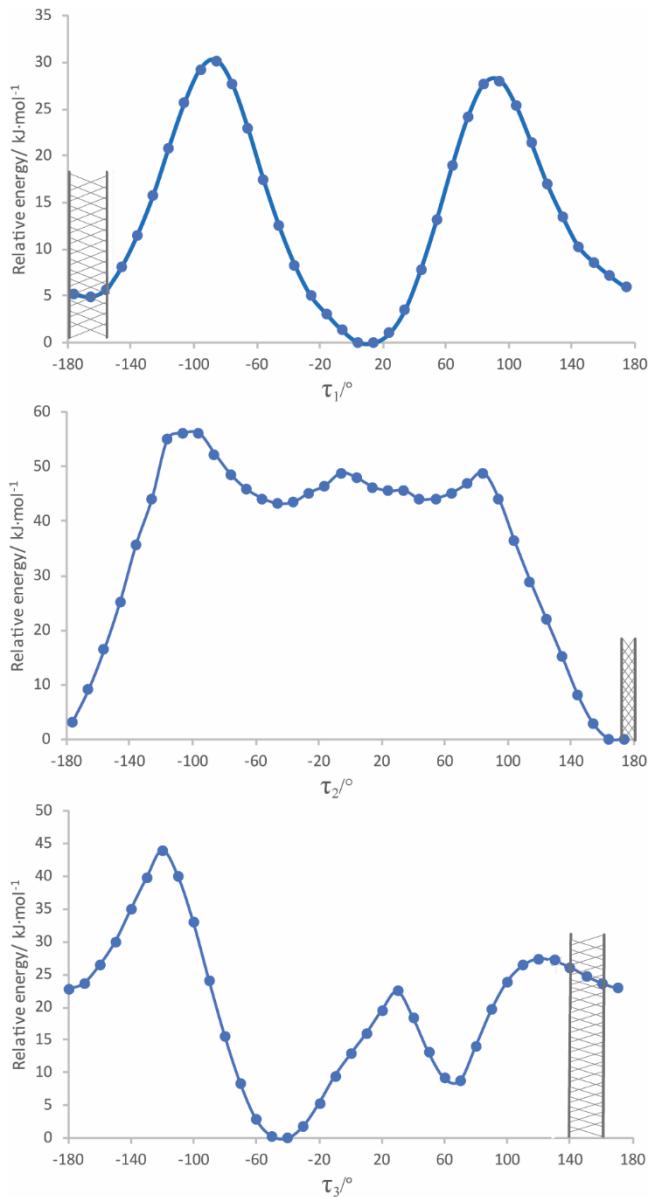
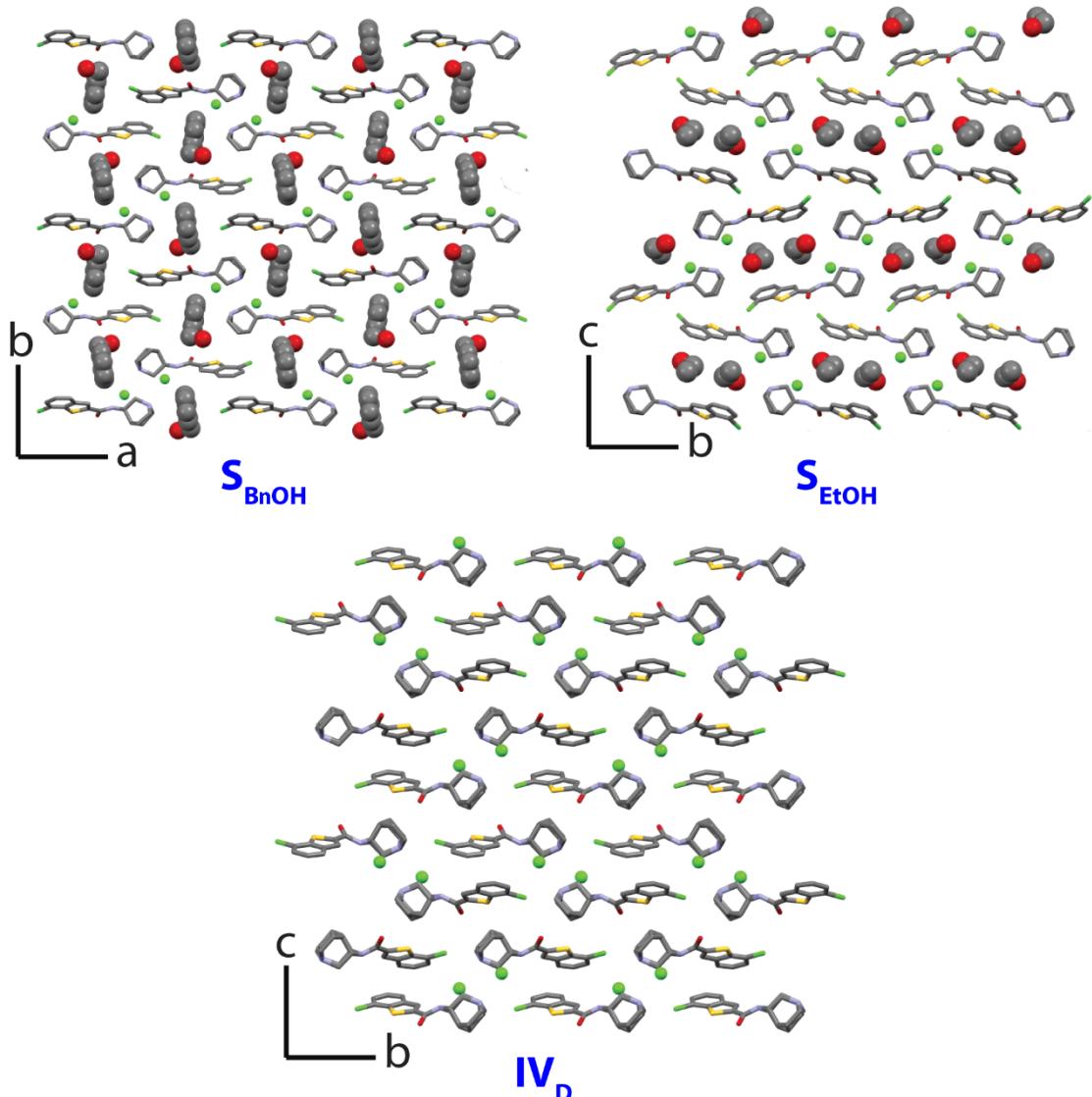
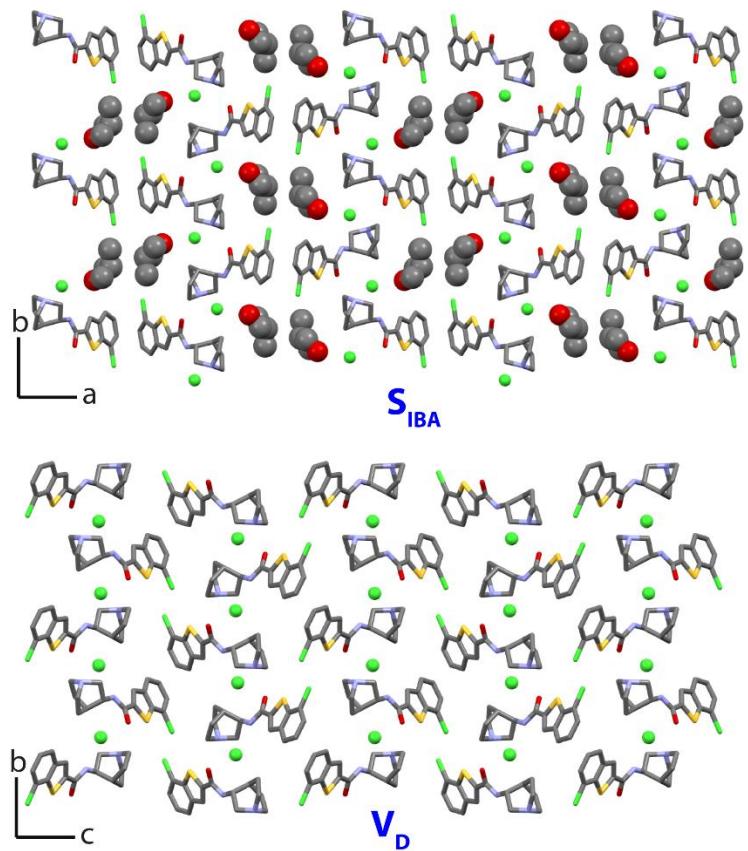


Figure S31. PES scans with respect to torsion angles τ_1 – τ_3 with marked interval of values observed in the experimental crystal structures.

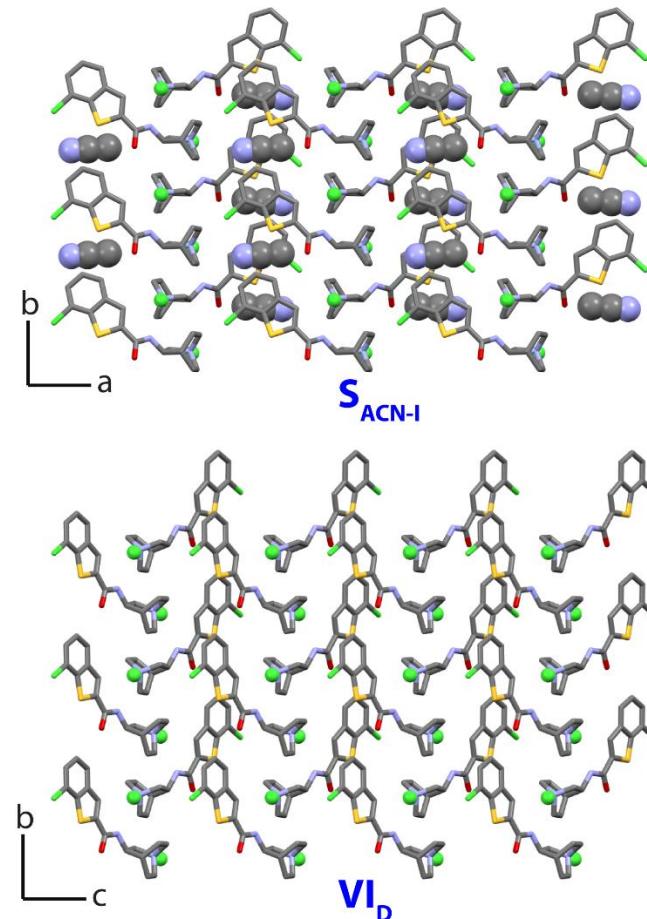
SI 6. Packing analysis



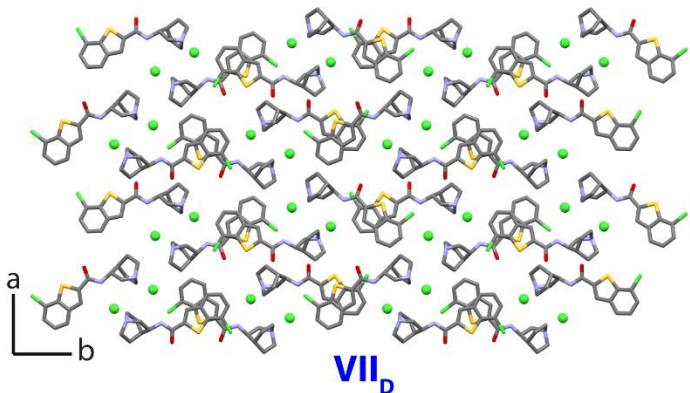
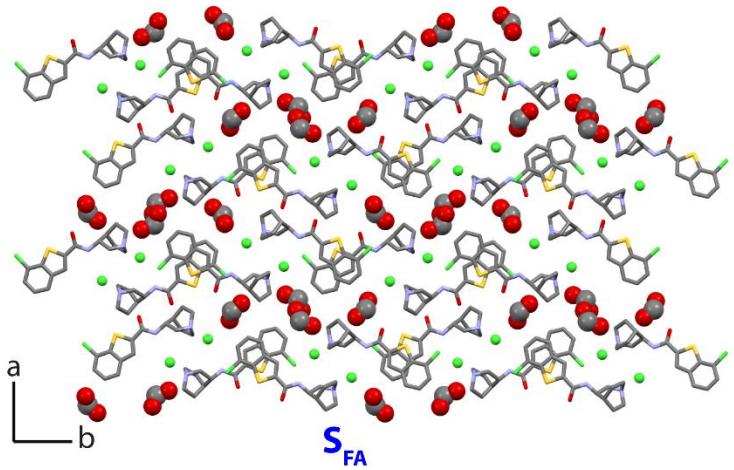
S32. Packing similarity between of Enc-HCl ethanol (S_{EtOH}), benzyl alcohol (S_{BnOH}) monosolvates and polymorph IV_D with omitted hydrogens.



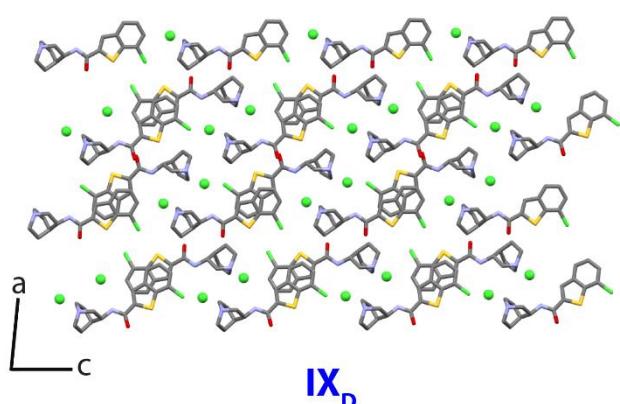
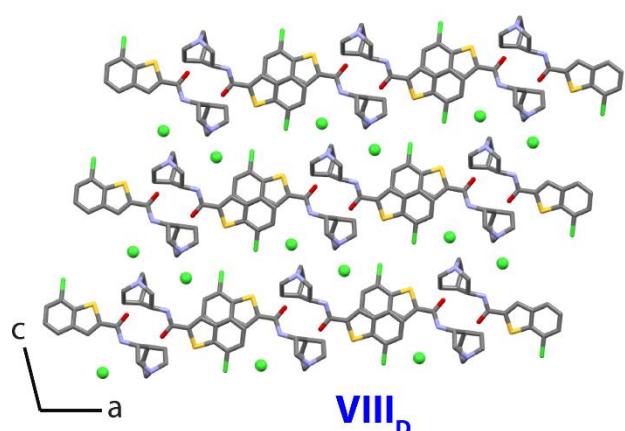
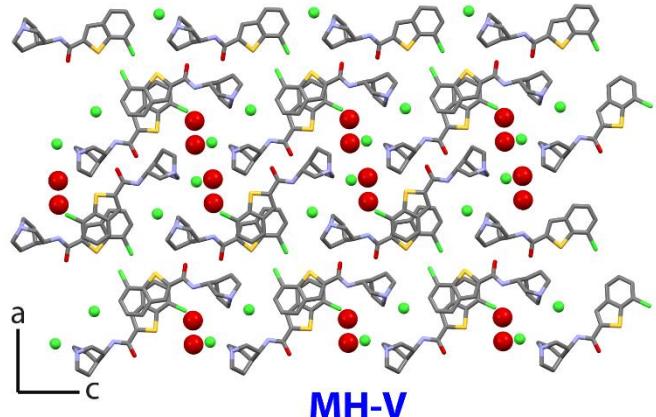
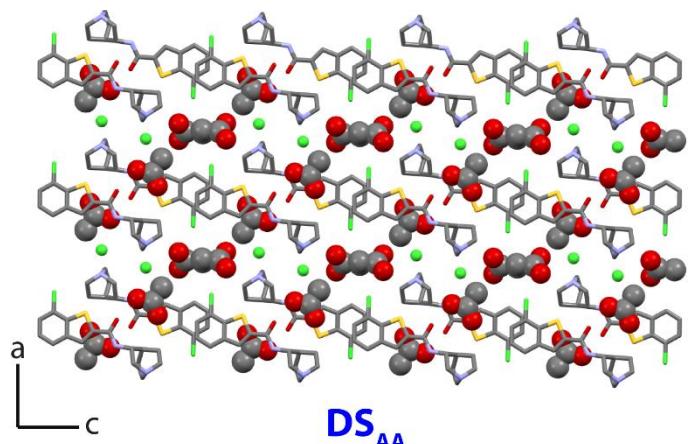
S33. Packing similarity between of Enc-HCl isobutanol monosolvate (S_{IBA}) and polymorph V_D with omitted hydrogens.



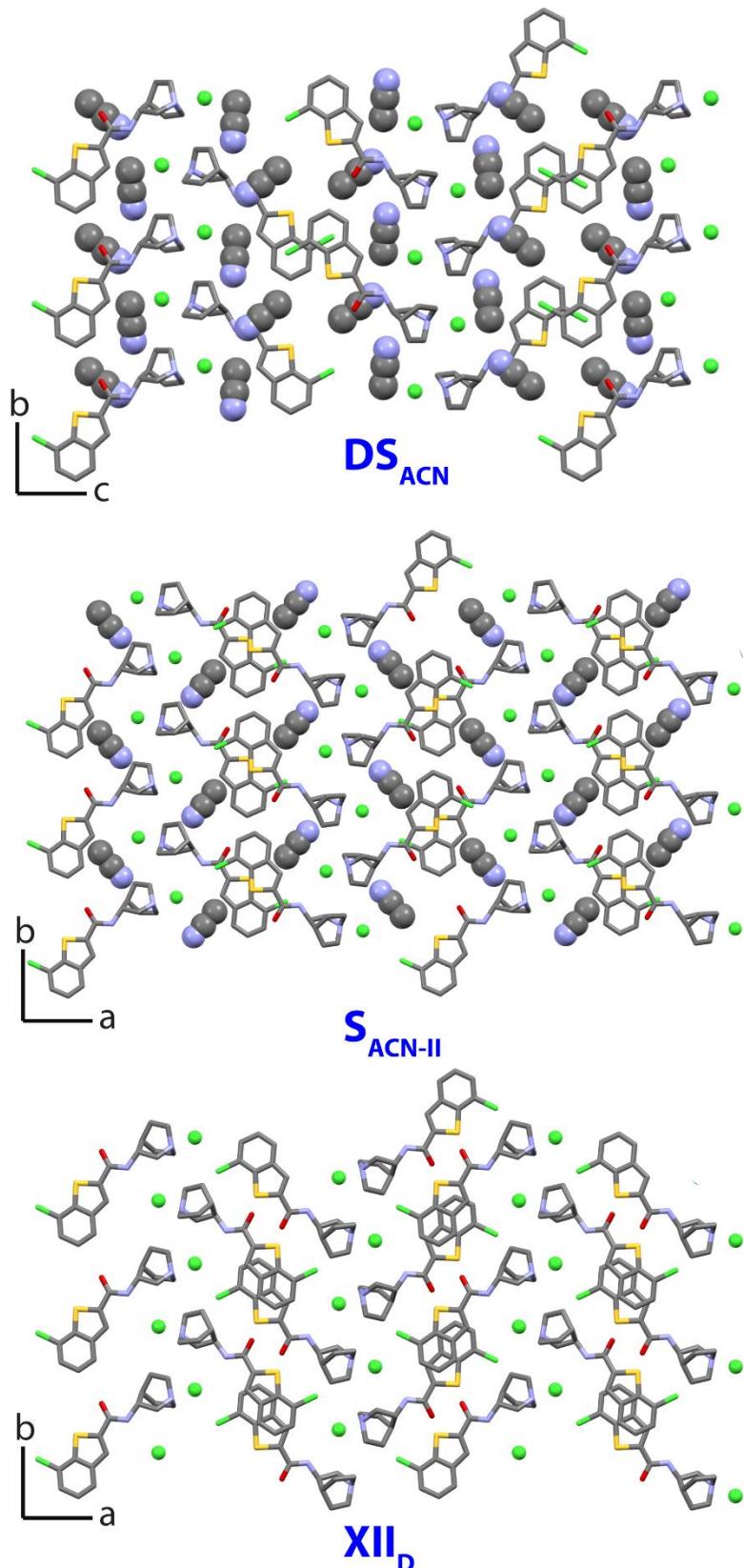
S34. Packing similarity between of Enc-HCl acetonitrile monosolvate 1 ($S_{ACN\text{-}I}$) and polymorph VI_D with omitted hydrogens.



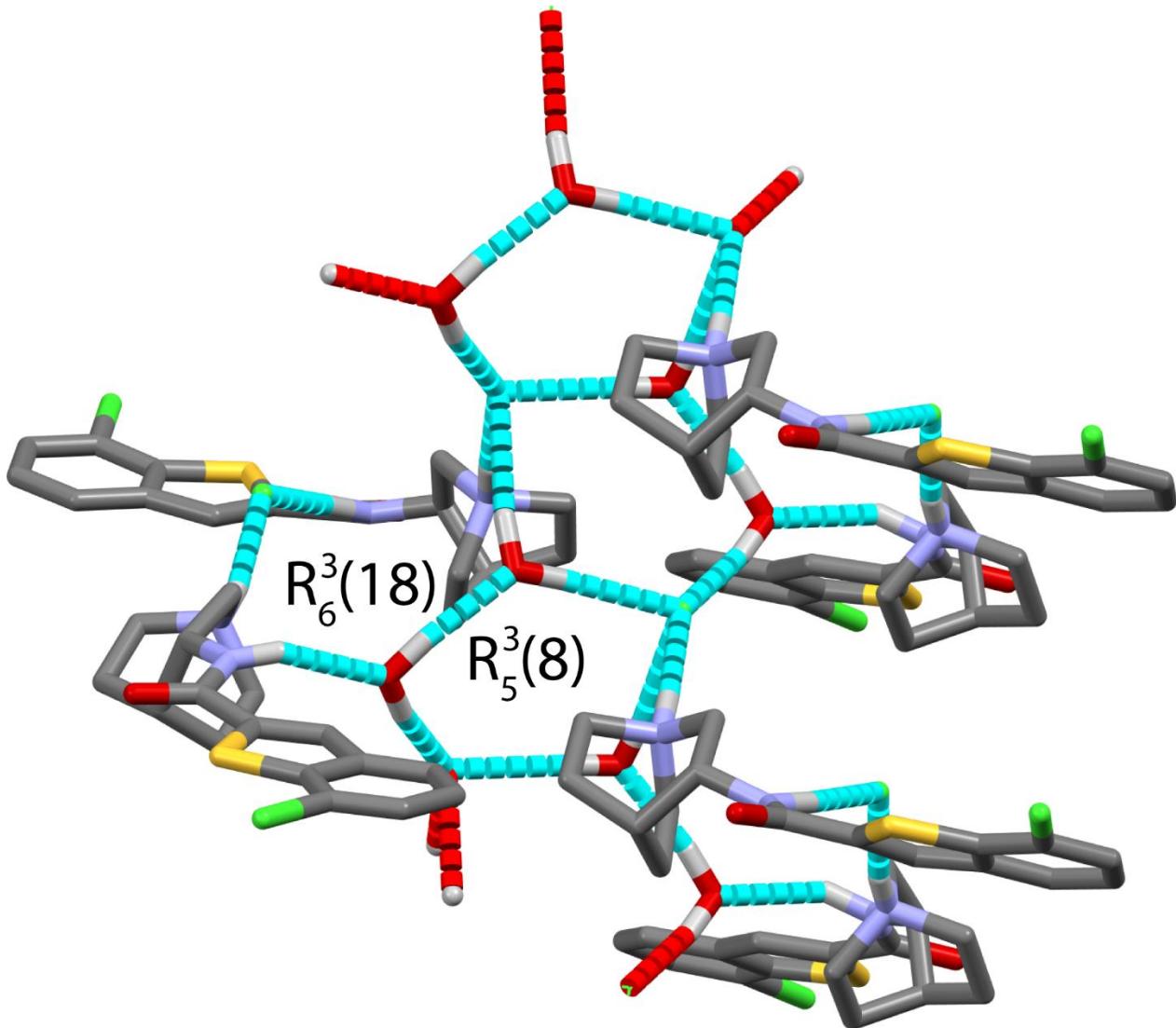
S35. Packing similarity between of Enc-HCl formic acid monosolvate (S_{FA}) and polymorph VII_D with omitted hydrogens.



S36. Packing similarity between of Enc-HCl acetic acid disolvate (DS_{AA}), monohydrate ($MH-V$), polymorphs $VIII_D$ and IX_D with omitted hydrogens.



S37. Packing similarity between of Enc-HCl acetonitrile disolvate (DS_{ACN}), acetonitrile monosolvate 2 (S_{ACN-II}) and polymorph XII_D with omitted hydrogens.



S38. Hydrogen bond network in Enc-HCl MH-V structure.