## Chemical Science

## Electronic supplementary material for:

## Thermal expansion properties of organic crystals: a CSD study

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**Fig. S1**: Fraction of observations that originates from the same study per number of temperatures used to determine the thermal expansion properties of a crystal structure. (lower) Global number of observations per number of temperatures used to determine the thermal expansion properties of a crystal structure **Fig. S2a-b**: Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in in the CSD (version 5.41). Complete dataset

**Fig. S3**: Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in the CSD (version 5.41). Only observations with  $R^2 > 0.75$  were used.

**Fig. S4**: Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in the CSD (version 5.41). Only observations with  $R^2 > 0.80$  were used.

**Fig. S5**: Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in the CSD (version 5.41). Only observations with  $R^2 > 0.85$  were used.

Fig. S6: Best overlay of the structures with *refcodes* VERMAG and VERMAG01

**Table S1**: Mean, median, and standard deviations of distributions of volumetric ( $\alpha_V$ ), uniaxial thermal expansion ( $\alpha_I$ ), and indicatrix anisotropy (IAC) coefficients as a function of  $R^2$ .

**Table S2**: Possible organic compounds with negative volumetric thermal expansion



**Fig. S1**: (upper) Fraction of observations that originates from the same study per number of temperatures used to determine the thermal expansion properties of a crystal structure. (lower) Global number of observations per number of temperatures used to determine the thermal expansion properties of a crystal structure An 'observation' is a set of thermal expansion coefficients for a particular compound.



**Figure S2a** Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in in the CSD (version 5.41). The complete data set was used to calculate these histograms.



**Figure S2b** Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in in the CSD (version 5.41). This figure is the same as Fig S2a but with different (zoomed) x-axes.



**Figure S3** Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in in the CSD (version 5.41). Only observations with  $R^2 > 0.75$  were used and for each data point at least 3 temperature points



**Figure S4** Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in in the CSD (version 5.41). Only observations with  $R^2 > 0.80$  were used and for each data point at least 3 temperature points



**Figure S5** Histograms of the volume and axis expansions and the indicatrix anisotropy coefficient of organic compounds in in the CSD (version 5.41). Only observations with  $R^2 > 0.85$  were used and for each data point at least 3 temperature points



Figure S6 Best overlay of the structures with refcodes VERMAG (red) and VERMAG01 (green). The unit cell of the VERMAG01 structure has been transformed to its conventional setting in  $P2_1/c$ . The rms deviation between the two structures is 0.108 Å. The calculations and visualization have been performed with MERCURY. The two structure packings are strictly the same. The room temperature structure (refcode VERMAG) was determined from Weissenberg camera data<sup>1</sup>, whereas the data for the 235 K structure were collected on a Syntex P2<sub>1</sub> diffractometer<sup>2</sup> (refcode VERMAG01). Interestingly, the densities of both structures were also determined using a standard flotation technique (1.31 and 1.17 gcm<sup>-3</sup> both measured at room temperature for the 235 and 295 K structures, respectively) and found to be in close correspondence with the density calculated from the X-ray crystal structures (1.326 and 1.16 gcm<sup>-3</sup> for the 235 K and 295 K structures, respectively). The difference in measured room temperature densities may seem suspicious or a sign of polymorphism. The latter can be ruled out by superposing the experimental structures, leaving the possibility that the room temperature density in the Selladurai study<sup>1</sup> was incorrectly assessed. Only a new temperature dependent X-ray study on the same crystal using the same diffractometer can validate or invalidate this result.

- 1. S. Selladarai, M. S. Kumar and K. Subramanian, Proc. Indian Acad. Sci. (Chem. Sci.), 1990, **102**, 39-43
- 2. H. W. Thompson, P. A. Vanderhoff and R. A. Lalancette, *Acta Crystallographica Section C Crystal Structure Communications*, 1991, **47**, 1443-1445.

	Global dataset	<i>R</i> <sup>2</sup> > 0.75	<i>R</i> <sup>2</sup> > 0.80	<i>R</i> <sup>2</sup> > 0.85	<i>R</i> <sup>2</sup> > 0.90
$n(\alpha_v)/n(\alpha_1)$	4719/14157	846/1403	828/1329	794/1248	745/1129
$\mu(\alpha_{\rm V})$	155.98	166.62	167.41	167.98	168.60
$\eta(\alpha_{\rm V})$	159.01	166.58	166.93	167.50	168.30
$\sigma(\alpha_{\rm V})$	90.85	73.87	73.05	72.55	72.51
$\mu(\alpha_1)$	51.64	66.20	67.80	69.34	71.43
$\eta(\alpha_1)$	42.57	56.98	58.23	59.74	62.57
$\sigma(\alpha_1)$	64.95	68.64	69.29	69.60	69.90
μ(IAC)	-0.06	-0.05	-0.05	-0.05	-0.05
η( IAC)	-0.07	-0.06	-0.06	-0.05	-0.05
<i>σ</i> ( IAC)	0.40	0.40*	0.40*	0.40*	0.41*

**Table S1** Mean, median, and standard deviations of distributions of volumetric ( $\alpha_V$ ), uniaxial thermal expansion ( $\alpha_l$ ), and indicatrix anisotropy (IAC) coefficients as a function of  $R^2$ .

Notes:  $n(\alpha_V)/n(\alpha)$  are the numbers of observations for the volumetric  $(\alpha_V)$  and axial  $(\alpha)$  thermal expansion coefficients, respectively.  $\mu$  is the mean value of the distribution,  $\eta$  the median value, and  $\sigma$  the standard deviation. The units of  $\alpha_V$  and  $\alpha_I$  are expressed in MK<sup>-1</sup>. The data sets noted by an asterisk are normally distributed. The 'global dataset' includes all temperature data sets, even those measured only at 2 temperatures; the columns  $R^2 > 0.75$ , 0.80, 0.85,0.90 include only data sets measured at at least 3 temperatures, and with goodness of fit values  $R^2$  for a linear dependency of thermal expansion on temperature.

refcode	∝ <sub>V</sub> (MK <sup>-1</sup> )	ΔТ (К)	Nm	χα	Ano	SS	Comments
EGIDUT	-623.12	230-296	2	-4.98			possibly x-ray-induced; polymerization reactions
							taking place; structure has the right metrics for
							photochemical SC-to-SC reactions, according to
							Schmid's rules; no heavy atoms
NEJQEY	-590.25	123-175	3	-3.07			data collection temperature at 175 K is not clear;
							hydrocarbon; $\pi \cdots \pi$ contacts but mostly H-H
							interactions and H-ring interactions
LATDAN	-460.18	120-173	2	-2.44			both are monohydrate, although one is reported as
							dehydrate, can be ok; H-bond backbone, π-π
							contacts but mostly H-H interactions and H-ring
							interactions

**Table S2** Possible organic compounds with negative volumetric thermal expansion.

DOFSUM	-370.75	173-295		-4.52	S		void space present? Can be correct
RALQAW	-318.10	90-173	2	-2.64			can be ok; some H-bonds, $\pi \cdots \pi$ contacts but mostly
							H-H interactions and H-ring interactions
MIPQAG	-306.41	100-273	2	-5.30		0	porous structure, CO <sub>2</sub> loss at RT?; no CO2 in
							structure, no H-bond backbone, H-ring interactions
							(first 3p element)
MEBVOH	-244.94	150-298	2	-3.63		0	can be ok; two 'isoforms' but why was one reported
							at 150 and the other at 298 K, structures seem to be
							the same; weak H-bonds, H-H dihydrogen contacts, 1
MOTOLIO	245.50	460.005	2	2.04			H-π interaction
MOTPUG	-215.59	160-295	2	-2.91			colorless at RT, orange at LT, weak H-bonds, H-H
							dihydrogen contacts, H··· $\pi$ interactions, some $\pi$ ··· $\pi$
41.474.0	211 50	172 201	2	2.50			contacts
ALAZAU	-211.50	1/3-291	Z	-2.50			can be ok; flattest molecule so far, H-bond backbone,
	104.64	160 272	2	2.04	-	-	no dinydrogen contacts, it stacking and H-it contacts
UKUYUU	-194.04	108-273	2	-2.04	5		can be ok; weak H-bonds (N-H, S-H), H-π contacts
	172.06	150 204	r	2.40			and it stacking
WUVIIV	-1/3.06	150-294	Z	-2.49			can be ok; neavy atom (Br), no amide H-bond, but
							in this being the first relatively flat melacule in the
							and this being the first relatively hat molecule in the
	160.07	220.206	r	1 20	6		selles
VEPKEJ	-109.07	220-296	Z	-1.28	5		color of LT crystal is colourless; that of RT crystal dark
							aliphatic chain: no # stacking
VEVOOL	166.25	100 172	2	1 21			can be correct: # stacking Haromatic Haliphatic
AEAQUH	-100.55	100-175	Z	-1.21			interactions weak H-bonds CH
KIZOAO	165 55	172 202	2	1 00			can be correct: Sr calt, if you are really nicky you
KIZQAU	-105.55	1/5-295	Z	-1.99			call be correct; Sr sait, if you are really picky you
							structure with strong H bonds, some H H
							interactions and $H_{-\pi}$ interactions
GEIVIB	-164.03	212-202	2	_1 21			can be correct: IT structure is really near quality: no
GEIVID	-104.05	215-255	2	-1.51			$\pi$ stacking many H-H contacts
SAIFIT	-156.67	215-296	2	-1 27	1	1	RT crystal colourless IT structure bright vellow: H-
575111	150.07	215 250	2	1.27			hand backhone $\pi$ -stacking many H··· $\pi$ and some H-
							H contacts
Ιυμχαο	-144.75	220-295	2	-1.09			can be correct: H-bonds provide the structure, some
	21		-	2.00			$\pi$ -stacking: several CH··· $\pi$ contacts (@ angle with
							respect to c axis)
MOXVIF	-142.37	85-173	2	-1.25		0	can be correct; H-bonds and $\pi$ -stacking form the
							backbone, some CH··· $\pi$ and H···H contacts present
CIWCIV	-135.56	223-296	2	-0.99	s		can be correct: very small difference between the R
					_		of the two structures (6.04% LT vs 6.07%) HT; H-
							bonds and $\pi$ -stacking form the backbone, some
							CH··· $\pi$ and H-H contacts present
KIKJEU	-135.13	125-294	2	-2.28			can be correct; some H-bonds, no $\pi$ -stacking, many
							CH- $\pi$ interactions in the bc plane (might be the
							mechanism)
VUTXUE	-134.68	103-273	2	-2.29			can be correct; however, should be rather classified
							as organometallic: Te covalent bond with C; S-S
							dipolar contacts, I-H H-bonds, some dihydrogen
							contacts
HOTNIO	-127.73	150-296	2	-1.86	s		can be correct; weak H-bonds, π-stacking
ATTDAZ	-121.54	199-295	2	-1.17			can be correct; H-bonds, $\pi$ -stacking, $\pi$ ··· $\pi$
							interactions
LIYLAJ	-120.09	100-173	2	-0.88		0	photo-induced switching, Both temperatures 100,
							173 should be probably 'room temperature'; CH $\cdots\pi$
							backbone in the ab plane (probably the cause of
							NTE), H-bond spiral between the diols along c,
							isolated from the rest of the interactions
NIZXEA	-106.37	150-293	2	-1.52			can be correct; Cl··· $\pi$ interaction, some H-bonds,
							some dihydrogen contacts
MIYBAX	-102.58	123-293	2	-1.74	S		can be correct, H-bonds and dipole stacking in the bc

							plane
QOYJOD	-91.33	223-293	2	-0.64			$\alpha$ polymorph, probably correct, H-bonds and $\pi$ stacking
YEDPAZ	-89.84	100-153	2	-0.48	s		can be correct, H-bond, $\pi$ and dipole stacking, weak
BEWKUM	-89.21	173-290	2	-1.04			can be correct. H-bond. $\pi$ stacking. CH- $\pi$ contact
TIFOOO	-89.21	225-293	2	-0.61			can be correct, $\pi$ -stacking and H-bonds
TUKWOI	-86.87	125-177	2	-0.45			can be correct. H-bonds, dihydrogen contacts
FAGSUB	-86.83	173-293	2	-1.04			can be correct, weak H-bond. CH- $\pi$ contacts
MUBJOJ	-86.72	130-296	3	-1.44		0	isosymmetric phase transition: no agreement
			-			-	between room temperature structure reported in
							Acta Cryst E and that in CrystEngComm, CH- $\pi$
							contacts, weak H-bond,
MATZEO	-86.12	173-293	2	-1.03	s		can be correct, weak CH····O bond, Type II CI-CI balogen bond, weak CH····CI contacts
τγραμη	-85.64	200-295	2	-0.81			can be correct possible regrientation of the H-bond
	03.01	200 255	-	0.01			network
ROHBOF	-82.14	100-173	2	-0.60			can be correct, weak CH···N H-bond, CH- $\pi$ contacts
DADMED	-82.13	150-296	2	-1.20	s		can be correct, some H-bonds, $\pi$ stacking
YUHHUE	-80.13	203-295	2	-0.74		0	can be correct, H-bond (OH···OH), CH···π contacts, $\pi$
							stacking
TIJDAR	-79.47	150-295	2	-1.15			can be correct
CEYYAG	-79.42	173-295	2	-0.97			can be correct, H-bond (OH···OH), CH···π contacts, π
							stacking
LUZXOU	-71.53	173-296	2	-0.88	S		LT crystal colourless, RT crystal yellow, H-bonds, $\pi$
							stacking, π···π contacts
ZUZBUT	-69.38	173-293	2	-0.83	S		can be correct, weak CH…O bonds, dihydrogen
							contacts, CH··· π contacts
CAZKUK	-68.92	150-205	2	-0.38			can be correct, H-bonds, $\pi$ stacking, CHO contacts
ADEREH	-66.98	173-292	2	-0.80			can be correct, $\pi$ stacking, CH…O contacts, CH… $\pi$
							contacts
ELEPOB	-59.65	100-200	3	-0.60			can be correct, H-bonds, $\pi$ stacking, $\pi \cdots \pi$ contacts
NOTION	-54.78	173-296	2	-0.67			can be correct, CH···O contacts, CH··· $\pi$ contacts, $\pi$
							stacking, π···π contacts
FIGKOY	-53.84	155-293	2	-0.74	S	0	can be correct, Cl atom half occupancy
KERNOM	-53.32	200-295	2	-0.51			can be correct, H-bonds, CH…O contacts, $\pi \dots \pi$ contacts
BEWYAE	-50.39	173-293	2	-0.60			can be correct, H-bonds, $\pi \cdots \pi$ contacts (unoccupied
AXUBUR	-49 30	90-293	2	-1 00			can be correct $\pi$ stacking CH···· $\pi$ contacts CH···F
7000000	13.50	50 255	-	1.00			bonds
FELCEH	-48.62	100-293	2	-0.94	1	0	can be correct. CH···· $\pi$ contacts. $\pi$ stacking (double
_						_	bond – ring)
PESKII	-47.25	100-193	2	-0.44			can be correct, π stacking, CH…O contacts,
							dihydrogen contacts
SOVBAG	-46.82	163-273	2	-0.51			can be correct, poorly solved, probably dihydrogen
							contacts
FOMZUD	-43.48	150-293	2	-0.62			can be correct, CH···O contacts, $\pi$ stacking
ASONOM	-43.06	133-296	2	-0.70			can be correct, dihydrogen contacts, CH… $\pi$ contacts
DOCKUE	-42.27	200-293	2	-0.39			can be correct, $\pi$ stacking, dihydrogen contacts,
QEDHUE	-41.96	153-293	2	-0.59			LT crystal yellow, RT colourless, H-bonds, $\pi$ stacking
CAPLAC	-40.34	123-295	2	-0.69			can be correct, H-bonds, dihydrogen contacts (along
							a axis)
TIGYUE	-39.32	150-293	2	-0.56			can be correct, H-bonds, CH $\cdots \pi$ contacts
FOKZUB	-38.98	213-293	2	-0.31	S		can be correct, H-bonds, dihydrogen contacts
WONVEA	-38.47	100-295	2	-0.75			can be correct
KUQSAS	-37.53	173-293	2	-0.45			color RT ligh-green, color LT colorless, CH… $\pi$
		ļ					contacts, $\pi$ stacking, $\pi \cdots \pi$ contacts
DAPSUO	-37.17	295-373	7	-0.29			Isosymmetric phase transition, H-bonds,
					ļ		CH···O/CH···N weak H-bonds, CH··· $\pi$ contacts
WEDMOI	-36.09	20-100	2	-0.29			can be correct, fluorine contacts, odd one out

MAZMAE         -34.18         100-153         2         -0.18         Contacts           NOTLAL         -33.38         113-293         2         -0.60         RT crystal grey, LT crystal gre	π onds
INDITIAL         33.8         103-135         2         0.160         RT crystal grey. IT crystal yellow, m stacking, m-m contacts, CH0 bonds           WIMVIZ         -33.37         173-293         2         -0.40         s         can be correct, T-montacts, CH0 bonds, m stacking, m-m contacts, CH0 bonds, m stacking, m-m contacts, CH0 bonds, m stacking, m-m contacts, CH0 bonds, m stacking, m m contacts, CH0 bonds, m stacking, CHn contacts           ACUFIO         -32.82         150-295         2         -0.48         can be correct, CH0 bonds, m stacking, CHn contacts           KAKCEH         -32.23         150-295         2         -0.47         0         can be correct, H-bonds, CHn m contacts           MEHBOR         -28.17         200-293         2         -0.54         can be correct, H-bonds, CHn m contacts           REXIZ         -28.06         293-546         2         -0.71         s         can be correct, H-bonds, CHn m contacts           MEHBOR         -27.11         173-273         2         -0.23         can be correct, H-bonds, CHn m contacts           NUZQOO         -26.58         173-295         2         -0.46         o         can be correct, H-bonds, m stacking, weak CH0 contacts           NUZQ	π onds
Normalized for the plan plan plan plan plan plan plan plan	onds
WIMVIZ         -33.37         173-293         2         -0.40         s         can be correct, π stacking, π-π contacts, CHI bo GARCAC         -32.95         159-297         2         -0.48         can be correct, H-mods, n-tacking, m-m contacts, CHI bo GARCAC	onds acts
SARCAC         -32.95         129-295         5         -0.55         o         can be correct, CHO contacts, dihydrogen contacts           ACUFIO         -32.82         150-297         2         -0.48         can be correct, CHDods, π stacking,           KAKCEH         -32.23         150-295         2         -0.47         o         can be correct, H-bonds, π stacking, CH-           HELHUD         -28.17         200-293         2         -0.54         can be correct, H-bonds, TH π contacts           REZSIZ         -28.06         293-546         2         -0.71         s         can be correct, H-bonds, CH π contacts           RAKVOR         -28.06         293-546         2         -0.71         s         can be correct, H-bonds, CH π contacts           BATLOZ         -26.58         173-296         2         -0.23         can be correct, H-bonds, CH π contacts, m stacking, dihydrogen contacts, m stacking, possi         pai-stacking, m stacking, possi           VIZGIX         -25.29         100-295         2         -0.20         can be correct, H-bonds, m stacking, weak CHO contacts, m stacking, stacking, meak           NUZQOO         -25.29         100-295 <td>acts</td>	acts
ACUFIO         -32.82         150-297         2         -0.48         can be correct, H-bonds, π stacking,           KAKCEH         -32.23         150-295         2         -0.47         0         can be correct, H-bonds, π stacking, CH-contacts           HELHUD         -28.17         200-293         2         -0.26         can be correct, H-bonds, CH-··π contacts           REZSIZ         -28.06         293-546         2         -0.71         s         can be correct, H-bonds, CH-··π contacts           RAKYOR         -28.06         293-546         2         -0.71         s         can be correct, H-bonds, CH-··π contacts           BATLOZ         -26.58         173-296         2         -0.33         s         can be correct, T, stacking, dihydrogen contacts, CH-··F contacts           NUZQOO         -25.30         130-295         2         -0.46         o         can be correct, T, bonds, CH-··π contacts, m stacking, possil           ANULEN         -25.29         100-295         3         -0.49         can be correct, H-bonds, TH-···C ontacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, H-bonds, fillydrogen contacts           RVFFED         -15.35         123-295         2         -0.20         can be correct, H-bonds, fill	Jucis
KAKCEH         -32.23         150-295         2         -0.47         o         can be correct void space 20-30%, π stacking, CH-contacts           REZSIZ         -28.08         100-293         2         -0.54         can be correct, H-bonds, CH-·· π contacts           REXSIZ         -28.08         100-293         2         -0.54         can be correct, H-bonds, CH-·· π contacts           MEHBOR         -27.11         173-273         2         -0.27         can be correct, H-bonds, CH-·· π contacts           BATLOZ         -26.58         173-296         2         -0.33         s         can be correct, H-bonds, CH-·· π contacts, m stacking, dihydrogen contacts, CH-··F contacts           RETNEM         -26.41         93-180         2         -0.23         can be correct, H-bonds, CH-·· π contacts, m stacking, view contacts, m stacking, contacts, m stacking, contac	
HELHUD         -28.17         200-293         2         -0.26         Can be correct, H-bonds, CH-···π contacts           REZSIZ         -28.08         100-293         2         -0.54         Can be correct, H-bonds, CH-···π contacts           RAKYOR         -28.06         293-546         2         -0.71         s         Can be correct, H-bonds, CH-···π contacts           MEHBOR         -27.11         173-273         2         -0.27         Can be correct, H-bonds, CH-···π contacts           BATLOZ         -26.58         173-296         2         -0.33         s         Can be correct, H-bonds, CH-···π contacts           RETNEM         -26.41         93-180         2         -0.23         Can be correct, π stacking, dihydrogen contacts, π stacking, VIZGIX         -25.46         113-295         2         -0.42         Can be correct, π stacking, dihydrogen contacts, m stacking, veak CH-··O contacts           NUZQOO         -25.30         130-295         2         -0.42         Can be correct, π stacking, weak CH-··O contacts           RURQAX         -16.69         173-295         2         -0.20         Can be correct, π stacking, weak CH-··O contacts           RURQAX         -16.64         100-200         2         -0.17         Can be correct, π stacking, weak CH-···O contacts           VIYFED <td>···π</td>	···π
HELHUD         -28.17         200-293         2         -0.26         can be correct, H-bonds, CH···π contacts           RZSIZ         -28.08         100-293         2         -0.54         can be correct, H-bonds, CH···π contacts           RAKYOR         -28.06         293-546         2         -0.71         s         can be correct, H-bonds, CH···π contacts           MEHBOR         -27.11         173-273         2         -0.27         can be correct, H-bonds, CH···π contacts           BATLOZ         -26.58         173-296         2         -0.33         s         can be correct, H-bonds, CH···π contacts, m contacts, m contacts           VIZGIX         -25.46         113-295         2         -0.42         can be correct, H-bonds           ANULEN         -25.29         100-295         3         -0.49         can be correct, H-bonds, m contacts, m stacking, possil phase transition due to changing occupancies           TEZZUU         -16.69         173-295         2         -0.20         can be correct, H-bonds, dihydrogen contacts, O····B contacts, dihydrogen contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, H-bonds, dihydrogen contacts           QOCXAJ         -14.60         150-295         2         -0.21         can be correct, H-bonds, dihy	
REZSIZ         -28.08         100-293         2         -0.54         can be correct, H-bonds, π stacking, F contacts           RAKYOR         -28.06         293-546         2         -0.71         s         can be correct, H-bonds, CHπ contacts           MEHBOR         -27.11         173-296         2         -0.27         can be correct, H-bonds, CHπ contacts           BATLOZ         -26.58         173-296         2         -0.23         can be correct, T, stacking, dihydrogen contacts, CHF contacts           RETNEM         -26.41         93-180         2         -0.23         can be correct, T-bonds, CH π contacts, π stack           VIZGIX         -25.46         113-295         2         -0.42         can be correct, T-bonds           ANULEN         -25.29         100-295         3         -0.49         can be correct, T-stacking, weak CH0 contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH0 contacts, 0	
RAKYOR         -28.06         293-546         2         -0.71         s         can be correct, H-bonds, CH··· π contacts           MEHBOR         -27.11         173-273         2         -0.27         can be correct, m contacts           BATLOZ         -26.58         173-296         2         -0.33         s         can be correct, m stacking, dihydrogen contacts, m contacts           RETNEM         -26.41         93-180         2         -0.23         can be correct, m stacking, dihydrogen contacts, m stacking, dihydrogen contacts, m contacts           VIZGIX         -25.46         113-295         2         -0.42         can be correct, H-bonds           NUZQOO         -25.30         130-295         2         -0.42         can be correct, H-bonds           ANULEN         -25.29         100-295         3         -0.49         can be correct, H-bonds, macking, weak CH···O contacts           TEZZUU         -16.69         173-295         2         -0.20         can be correct, first ding/orgen contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, first ding/orgen contacts           RURQAX         -16.64         100-200         2         -0.21         can be correct, first ding/orgen contacts           KIYFED	
MEHBOR         -27.11         173-273         2         -0.27         can be correct, H-bonds, CH··· π contacts           BATLOZ         -26.58         173-296         2         -0.33         s         can be correct, π stacking, dihydrogen contacts, CH···F contacts           RETNEM         -26.41         93-180         2         -0.23         can be correct, π stacking, dihydrogen contacts, CH···F contacts           NUZQOO         -25.30         130-295         2         -0.46         o         can be correct, H-bonds, CH··· π contacts, π stacking, dihydrogen contacts, CH···O contacts           NUZQOO         -25.30         130-295         2         -0.42         can be correct, H-bonds           ANULEN         -25.29         100-295         3         -0.49         can be correct, H-bonds, m stacking, weak CH···O contacts           RURQAX         -16.69         173-295         2         -0.20         can be correct, first 3d polymer structure, n stacking, weak CH···O contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, n stacking, weak CH···O contacts           QOXXAJ         -14.60         150-295         2         -0.21         s         can be correct, H-bonds, dihydrogen contacts           BOGCAC         -13.96         90-150         2 <td< td=""><td></td></td<>	
BATLOZ         -26.58         173-296         2         -0.33         s         can be correct, π stacking, dihydrogen contacts, CH···F contacts           RETNEM         -26.41         93-180         2         -0.23         can be correct, π stacking, dihydrogen contacts, π stack           VIZGIX         -25.46         113-295         2         -0.46         o         can be correct, π stacking, dihydrogen contacts, π stack           NUZQOO         -25.30         130-295         2         -0.42         can be correct, π contacts, π stacking, possil           ANULEN         -25.29         100-295         3         -0.49         can be correct, π stacking, weak CH···O contacts           TEZZUU         -16.69         173-295         2         -0.26         can be correct, π stacking, weak CH···O contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH···O contacts           KIYFED         -15.35         123-295         2         -0.21         can be correct, n stacking weak CH···O contacts           QOCXAJ         -14.40         150-296         2         -0.21         s         can be correct, n stacking CH··· π contacts, m stacking           BOGCAC         -13.96         90-150         2         -0.08         can be correct,	
RETNEM         -26.41         93-180         2         -0.23         can be correct, H-bonds, CH-·· π contacts, π stack           VIZGIX         -25.46         113-295         2         -0.46         0         can be correct, π stacking, dihydrogen contacts, CH··· 0 contacts           NUZQOO         -25.30         130-295         2         -0.42         can be correct, H-bonds           ANULEN         -25.29         100-295         3         -0.49         can be correct, CH··· π contacts, π stacking, possil phase transition due to changing occupancies           TEZZUU         -16.69         173-295         2         -0.20         can be correct, H-bonds, π stacking, weak CH···O contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH···O contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, H-bonds, dihydrogen contacts           QOCXAJ         -14.60         150-295         2         -0.21         can be correct, CH··· π contacts, weak CH···O contacts           BOGCAC         -13.96         90-150         2         -0.21         s         can be correct, CH··· π contacts, m stack find ydrogen contacts           HAPLAN         -13.46         173-293         2         -0.16         ca	
VIZGIX         -25.46         113-295         2         -0.46         o         can be correct, π stacking, dihydrogen contacts, CH···O contacts           NUZQOO         -25.30         130-295         2         -0.42         can be correct, π stacking, dihydrogen contacts, π stacking, possil phase transition due to changing occupancies           ANULEN         -25.29         100-295         3         -0.49         can be correct, CH···π contacts, π stacking, weak CH···O contacts           TEZZUU         -16.69         173-295         2         -0.20         can be correct, Thoods, π stacking, weak CH···O contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, Thoods, π stacking, weak CH···O contacts, O····Br contacts, dihydrogen contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, First 3d polymer structure, π stack           QOCXAJ         -14.60         150-295         2         -0.21         s         can be correct, H-bonds, CH··· π contacts, weak CH···O contacts           BOGCAC         -13.96         90-150         2         -0.08         can be correct, H-bonds, CH··· π contacts, m stacking CH··· π contacts, m stacking CH··· π contacts, m stacking CH··· π contacts, m stack           NOBFEP         -13.46         173-293         2         -0.16         can be corr	king
NUZQOO         -25.30         130-295         2         -0.42         can be correct, H-bonds           ANULEN         -25.29         100-295         3         -0.49         can be correct, CH··· π contacts, π stacking, positi phase transition due to changing occupancies           TEZZUU         -16.69         173-295         2         -0.20         can be correct, H-bonds, π stacking, weak CH···O contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH···O contacts, 0···Br contacts, dihydrogen contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, first 3d polymer structure, π stack indy drogen contacts           FAHDOI         -14.60         150-295         2         -0.21         can be correct, first 3d polymer structure, π stack H-bonds, dihydrogen contacts           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, π stacking CH··· π contacts, mastack H-bonds, CH··· π contacts, mastack H-Donds           NOBFEP         -13.46         173-293         2         -0.26         can be correct, CH··· π contacts, mastack H-··O contacts           HAPLAN         -13.44         100-293         2         -0.26         s         can be correct, m stacking CH··· π contacts, mastack H-··O contacts	
NUZQO0         -25.30         130-295         2         -0.42         can be correct, H-bonds           ANULEN         -25.29         100-295         3         -0.49         can be correct, CH··· π contacts, π stacking, posil phase transition due to changing occupancies           TEZZUU         -16.69         173-295         2         -0.20         can be correct, H-bonds, π stacking, weak CH···O contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH···O contacts, O···Br contacts, dihydrogen contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, first 3d polymer structure, π stack H-bonds, dihydrogen contacts           QOCXAJ         -14.60         150-295         2         -0.21         s         can be correct, GH··· π contacts, weak CH···O contacts, dihydrogen contacts           BOGCAC         -13.96         90-150         2         -0.28         can be correct, R+bonds, CH··· π contacts, π stack NOBFEP           -13.46         173-293         2         -0.16         can be correct, CH··· π contacts, weak CH···O contacts, dihydrogen contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, T··· π contacts, weak CH···O contacts, dihydrogen contacts           BOWMIK	
ANULEN         -25.29         100-295         3         -0.49         can be correct, CH··· π contacts, π stacking, possil phase transition due to changing occupancies           TEZZUU         -16.69         173-295         2         -0.20         can be correct, H-bonds, π stacking, weak CH···O contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH···O contacts, O···Br contacts, dihydrogen contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, π stacking, weak CH···O contacts           FAHDOI         -14.60         150-295         2         -0.21         can be correct, first 3d polymer structure, π stack           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, CH··· π contacts, weak CH···O contacts           BOGCAC         -13.96         90-150         2         -0.08         can be correct, R stacking CH··· π contacts, π stack Miydrogen contacts           HAPLAN         -13.44         100-293         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-293         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts, dihydrogen contacts           BOWMI	
TEZZUU         -16.69         173-295         2         -0.20         can be correct, H-bonds, π stacking, weak CH···O contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH···O contacts, 0···Br contacts, dihydrogen contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, first 3d polymer structure, π stacking, weak CH···O contacts, dihydrogen contacts           FAHDOI         -14.60         150-295         2         -0.21         can be correct, first 3d polymer structure, π stack           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, H-bonds, dihydrogen contacts           BOGCAC         -13.96         90-150         2         -0.21         s         can be correct, T-m contacts, m stack           NOBFEP         -13.46         173-293         2         -0.16         can be correct, π stacking CH··· π contacts, dihydrogen contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, R-m contacts, weak CH···O contacts           BOWMIK         -13.44         100-298         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -12.63	ible
TEZZUU         -16.69         173-295         2         -0.20         can be correct, H-bonds, π stacking, weak CH···O contacts           RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH···O contacts, 0···Br contacts, dihydrogen contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, H-bonds, dihydrogen contacts           FAHDOI         -14.60         150-295         2         -0.21         can be correct, first 3d polymer structure, π stack H-bonds, dihydrogen contacts           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, CH··· π contacts, weak CH···O contacts, dihydrogen contacts           BOGCAC         -13.96         90-150         2         -0.08         can be correct, π stacking CH··· π contacts, π stack MOBFEP         173-293         2         -0.16         can be correct, π stacking CH··· π contacts, m stack Mihydrogen contacts           HAPLAN         -13.44         100-293         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-298         2         -0.26         o         False hit, synchrotron powder data, however no carbon present, but only boron; see text           AMASOY         -12.75         90	
RURQAX         -16.64         100-200         2         -0.17         Contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, π stacking, weak CH···O contacts, O···Br contacts, dihydrogen contacts           FAHDOI         -14.60         150-295         2         -0.21         can be correct, H-bonds, dihydrogen contacts           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, CH··· π contacts, weak CH···O contacts           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, H-bonds, CH··· π contacts, weak CH···O contacts           BOGCAC         -13.96         90-150         2         -0.08         can be correct, π stacking CH··· π contacts, m stack           NOBFEP         -13.46         173-293         2         -0.16         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-293         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts, dihydrogen contacts           AMASOY         -12.75         90-296         3 <t< td=""><td>)</td></t<>	)
RURQAX         -16.64         100-200         2         -0.17         can be correct, π stacking, weak CH···O contacts, O···Br contacts, dihydrogen contacts           KIYFED         -15.35         123-295         2         -0.26         can be correct, π stacking, weak CH···O contacts, O···Br contacts, dihydrogen contacts           FAHDOI         -14.60         150-295         2         -0.21         can be correct, first 3d polymer structure, π stack H-bonds, dihydrogen contacts           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, CH··· π contacts, weak CH···O contacts, dihydrogen contacts           BOGCAC         -13.96         90-150         2         -0.08         can be correct, π stacking CH··· π contacts, π stack dihydrogen contacts           HAPLAN         173-293         2         -0.16         can be correct, π stacking CH··· π contacts, weak CH···O contacts           BOWMIK         -13.44         100-293         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts, dihydrogen c	
KIYFED         -15.35         123-295         2         -0.26         can be correct, H-bonds, dihydrogen contacts           FAHDOI         -14.60         150-295         2         -0.21         can be correct, first 3d polymer structure, π stack H-bonds, dihydrogen contacts           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, GH··· π contacts, weak CH···O contacts, dihydrogen contacts           BOGCAC         -13.96         90-150         2         -0.08         can be correct, π stacking CH··· π contacts, π stack dihydrogen contacts           NOBFEP         -13.46         173-293         2         -0.16         can be correct, π stacking CH··· π contacts, dihydrogen contacts           HAPLAN         -13.44         100-293         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -12.75         90-296         3         -0.26         o         False hit, synchrotron powder data, however no carbon present, but only boron; see text           AMASOY         -12.75         90-296         3         -0.26         color red or orange, π stacking CH··· π contacts, dihydrogen contacts, possible phase tr	,
KIYFED         -15.35         123-295         2         -0.26         can be correct, H-bonds, dihydrogen contacts           FAHDOI         -14.60         150-295         2         -0.21         can be correct, first 3d polymer structure, π stack H-bonds, dihydrogen contacts           QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, CH··· π contacts, weak CH···O contacts, dihydrogen contacts           BOGCAC         -13.96         90-150         2         -0.08         can be correct, H-bonds, CH··· π contacts, π stack Gan be correct, H-bonds, CH··· π contacts, π stack           NOBFEP         -13.46         173-293         2         -0.16         can be correct, CH··· π contacts, weak CH···O contacts           HAPLAN         -13.44         100-293         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -12.75         90-296         3         -0.26         o         False hit, synchrotron powder data, however no carbon present, but only boron; see text           AMASOY         -12.75         90-296         3         -0.26         color red or oranage, π stacking CH··· π contacts, dihydrogen contact	
FAHDOI       -14.60       150-295       2       -0.21       Can be correct, first 3d polymer structure, π stack H-bonds, dihydrogen contacts         QOCXAJ       -14.30       150-296       2       -0.21       s       Can be correct, first 3d polymer structure, π stack H-bonds, dihydrogen contacts         BOGCAC       -13.96       90-150       2       -0.08       Can be correct, H-bonds, CH… π contacts, weak CH…O contacts, dihydrogen contacts         NOBFEP       -13.46       173-293       2       -0.16       Can be correct, π stacking CH… π contacts, π stack dihydrogen contacts         HAPLAN       -13.44       100-293       2       -0.26       s       Can be correct, CH… π contacts, weak CH…O contacts         BOWMIK       -13.35       100-298       2       -0.26       s       Can be correct, CH… π contacts, weak CH…O contacts         AMASOY       -12.75       90-296       3       -0.26       O       False hit, synchrotron powder data, however no carbon present, but only boron; see text         AMASOY       -12.63       150-296       2       -0.18       Can be correct, CH… π contacts, weak CH…O contacts         SECHUG       -12.09       150-293       2       -0.17       Can be correct, H-bonds, weak CH…O contacts, π	
QOCXAJ         -14.30         150-296         2         -0.21         s         can be correct, CH… π contacts, weak CH…O contacts, dihydrogen contacts           BOGCAC         -13.96         90-150         2         -0.08         can be correct, CH… π contacts, weak CH…O contacts, dihydrogen contacts           NOBFEP         -13.46         173-293         2         -0.16         can be correct, π stacking CH… π contacts, π stacking CH… π contacts, dihydrogen contacts           HAPLAN         -13.44         100-293         2         -0.26         s         can be correct, CH… π contacts, weak CH…O contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, CH… π contacts, weak CH…O contacts           BOWMIK         -12.75         90-296         3         -0.26         contacts         Color red or orange, π stacking CH… π contacts, dihydrogen contacts, dihydrogen contacts           AMASOY         -12.75         90-296         3         -0.26         O         False hit, synchrotron powder data, however no carbon present, but only boron; see text           AMASOY         -12.63         150-296         2         -0.18         can be correct, CH… π contacts, weak CH…O contacts           SECHUG         -12.09         150-293         2         -0.17         can be correct, H-bonds, weak CH…O c	king,
QOCXAJ       -14.30       150-296       2       -0.21       s       can be correct, CH··· π contacts, weak CH···O contacts, dihydrogen contacts         BOGCAC       -13.96       90-150       2       -0.08       can be correct, H-bonds, CH··· π contacts, π stack         NOBFEP       -13.46       173-293       2       -0.16       can be correct, π stacking CH··· π contacts, dihydrogen contacts         HAPLAN       -13.44       100-293       2       -0.26       s       can be correct, CH··· π contacts, weak CH···O contacts         BOWMIK       -13.35       100-298       2       -0.26       s       can be correct, CH··· π contacts, weak CH···O contacts         BOWMIK       -12.75       90-296       3       -0.26       s       can be correct, CH··· π contacts, possible phase transition (disorder in HT)         ZETFIQ       -12.63       150-296       2       -0.18       can be correct, CH··· π contacts, weak CH···O contacts         SECHUG       -12.09       150-293       2       -0.17       can be correct, H-bonds, weak CH···O contacts, π	
BOGCAC         -13.96         90-150         2         -0.08         can be correct, H-bonds, CH··· π contacts, π stack           NOBFEP         -13.46         173-293         2         -0.16         can be correct, π stacking CH··· π contacts, dihydrogen contacts           HAPLAN         -13.44         100-293         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-298         2         -0.26         s         can be correct, CH··· π contacts, weak CH···O contacts           BOWMIK         -13.35         100-298         2         -0.26         o         False hit, synchrotron powder data, however no carbon present, but only boron; see text           AMASOY         -12.75         90-296         3         -0.26         o         False hit, synchrotron powder data, however no carbon present, but only boron; see text           ZETFIQ         -12.63         150-296         2         -0.18         can be correct, CH··· π contacts, weak CH···O contacts           SECHUG         -12.09         150-293         2         -0.17         can be correct, H-bonds, weak CH···O contacts, π	
BOBCAC-13.3690-1502-0.08Call be correct, H-bolds, CH π contacts, n stackNOBFEP-13.46173-2932-0.16can be correct, π stacking CH π contacts, dihydrogen contactsHAPLAN-13.44100-2932-0.26scan be correct, CH π contacts, weak CHO contactsBOWMIK-13.35100-2982-0.26oFalse hit, synchrotron powder data, however no carbon present, but only boron; see textAMASOY-12.7590-2963-0.26Color red or orange, π stacking CH π contacts, dihydrogen contacts, possible phase transition (disorder in HT)ZETFIQ-12.63150-2962-0.18can be correct, H-bonds, weak CHO contactsSECHUG-12.09150-2932-0.17can be correct, H-bonds, weak CHO contacts, π	lina
NOBPEP-13.40173-2932-0.10Call be correct, it stacking CH··· it contacts, dihydrogen contactsHAPLAN-13.44100-2932-0.26scan be correct, CH··· π contacts, weak CH···O contactsBOWMIK-13.35100-2982-0.26oFalse hit, synchrotron powder data, however no carbon present, but only boron; see textAMASOY-12.7590-2963-0.26Color red or orange, π stacking CH··· π contacts, dihydrogen contacts, possible phase transition (disorder in HT)ZETFIQ-12.63150-2962-0.18can be correct, CH··· π contacts, weak CH···O contactsSECHUG-12.09150-2932-0.17can be correct, H-bonds, weak CH···O contacts, π	ling
HAPLAN-13.44100-2932-0.26scan be correct, CH··· π contacts, weak CH···O contactsBOWMIK-13.35100-2982-0.260False hit, synchrotron powder data, however no carbon present, but only boron; see textAMASOY-12.7590-2963-0.26Color red or orange, π stacking CH··· π contacts, dihydrogen contacts, possible phase transition (disorder in HT)ZETFIQ-12.63150-2962-0.18can be correct, CH··· π contacts, weak CH···O contactsSECHUG-12.09150-2932-0.17can be correct, H-bonds, weak CH···O contacts, π	
BOWMIK       -13.35       100-298       2       -0.26       O       False hit, synchrotron powder data, however no carbon present, but only boron; see text         AMASOY       -12.75       90-296       3       -0.26       Color red or orange, π stacking CH··· π contacts, dihydrogen contacts, possible phase transition (disorder in HT)         ZETFIQ       -12.63       150-296       2       -0.18       can be correct, CH··· π contacts, weak CH···O contacts         SECHUG       -12.09       150-293       2       -0.17       can be correct, H-bonds, weak CH···O contacts, π	
AMASOY     -12.75     90-296     3     -0.26     Color red or orange, π stacking CH··· π contacts, dihydrogen contacts, possible phase transition (disorder in HT)       ZETFIQ     -12.63     150-296     2     -0.18     can be correct, CH··· π contacts, weak CH···O contacts       SECHUG     -12.09     150-293     2     -0.17     can be correct, H-bonds, weak CH···O contacts, π	
AMASOY       -12.75       90-296       3       -0.26       Color red or orange, π stacking CH··· π contacts, dihydrogen contacts, possible phase transition (disorder in HT)         ZETFIQ       -12.63       150-296       2       -0.18       can be correct, CH··· π contacts, weak CH···O contacts         SECHUG       -12.09       150-293       2       -0.17       can be correct, H-bonds, weak CH···O contacts, π	
ZETFIQ     -12.63     150-296     2     -0.18     can be correct, CH··· π contacts, weak CH···O contacts       SECHUG     -12.09     150-293     2     -0.17     can be correct, H-bonds, weak CH···O contacts, π	
ZETFIQ         -12.63         150-296         2         -0.18         Can be correct, CH··· π contacts, weak CH···O contacts           SECHUG         -12.09         150-293         2         -0.17         can be correct, H-bonds, weak CH···O contacts, π	
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SECHUG         -12.09         150-293         2         -0.17         Can be correct, H-bonds, weak CH···O contacts, π	
SECHUG -12.09 150-293 2 -0.17 can be correct, H-bonds, weak CH…O contacts, π	
	τ
stacking	
MORPHM -12.01 25-295 2 -0.32 can be correct, H-bonds, weak CH···O contacts, $\pi$	C
TEINAX -10.71 85-150 2 -0.07 con be correct. Η bonds π stacking (corbox/lates)	c)
TARDOF -10.29 173-295 2 -0.07 Can be correct. H-bonds, it stacking (tabboy) acts	3)
OMERUT -9.58 110-293 2 -0.18 0 CSD notes that there is doubt about the reported	d
temperature of one of the structures, a methylen	ne
bridge between to N changes in a strange manne	er, I
would not include it	
PEVNUC -8.86 280-400 4 -0.11 0 correct, same crystal same machine; longest singl	gle C-
C bond ever reported (makes sense from a chemi	ical
point of view: high steric hinderance), π stacking	i.
CH··· π contacts, dihydrogen contacts	
XOTGIY     -8.59     173-296     2     -0.11     can be correct, π stacking, CH··· π contacts, weak       CH···O contacts     CH···O contacts	<
IFISUP -8.57 110-293 2 -0.16 can be correct. CH····π contacts weak CH····Ο	
contacts	

QAVTUE	-8.39	193-296	2	-0.09			can be correct, H-bonds, $\pi$ stacking, CH··· $\pi$ contacts, weak CH··· $\pi$ contacts
WUXNIM	-7.58	92-180	2	-0.07			can be correct. F: $\pi$ contacts. $\pi$ stacking
QUWYAJ	-6.59	123-293	2	-0.11			can be correct, H-bonds, π stacking
ETDAMS	-6.54	100-295	2	-0.13			can be correct, H-bonds, dihydrogen contacts
NEZGAD	-6.04	105-295	2	-0.11		0	can be correct, H-bonds, $\pi$ stacking, weak CH···O/
							CH…F contacts
SILTEP	-5.94	160-273	2	-0.07			can be correct, CH··· $\pi$ contacts, dihydrogen contacts
HOQMIL	-5.76	173-293	3	-0.07			can be correct, porous structure with solvent inside
							(DMF)
JAGRIV	-5.75	150-298	2	-0.09	s		can be correct, H-bonds, $\pi$ stacking, CH… $\pi$ contacts ,
							weak CH…O contacts
ROYHAQ	-5.65	100-294	2	-0.11			can be correct, H-bonds, $\pi$ stacking, CH… $\pi$ contacts ,
							weak CH…O contacts, dihydrogen contacts
XUKKET	-3.83	170-593	3	-0.16		0	can be correct, second 3D polymeric structure, some
							π stacking (acetylenes)
KOZTOK	-3.35	90-293	2	-0.07			LT color yellow; RT light colorless, $\pi$ stacking, CH… $\pi$
							contacts , weak CH…O contacts
BAXSIC	-3.19	80-295	2	-0.07			can be correct, H-bonds, weak CH…O contacts
ITIRAE	-2.71	140-296	2	-0.04	S		can be correct H-bonds, weak CH···O contacts, $\pi$ ··· $\pi$
	0.00		-				contacts (B to Ph)
DOPLOL	-2.69	120-451	2	-0.09		0	can be correct, H-bonds, $\pi$ stacking
CELLOB	-2.27	120-1/3	3	-0.01			can be correct, H-bonds, weak CH···O contacts
JOYQUJ	-2.06	1/3-295	2	-0.03			can be correct, H-bonds, $\pi$ stacking, CHBr contacts
ODAKOV	-2.04	173-293	2	-0.02	S		can be correct, H-bonds, dihydrogen contacts,
							possible rearrangement of the of the H-bond
	1 9 2	200 202	r	0.02			network inside the channel
	-1.82	152 202	2	-0.02			can be correct. H bonds # stacking weak CHO
101101	-1.04	133-295	5	-0.02			contacts
	-1 5/	153-295	2	-0.02			contacts
CEIVICAL	1.54	155 255	2	0.02			dihydrogen contacts
GLUTAS	-1.29	100-295	3	-0.03			can be correct. H-bonds, weak CH···O contacts
KIMSUU	-1.17	122-295	2	-0.02			can be correct. CH··· $\pi$ contacts, dihydrogen contacts
DAYGUH	-1.04	100-296	2	-0.02	s		can be correct, et a contacts, university to be contacts and the contacts $\pi$
27110011	2.0	100 100	-	0.02	Ũ		stacking
НЕКМЕР	-0.59	121-295	2	-0.01			can be correct. H-bonds, weak CH…O contacts.
							dihydrogen contacts
WADQEY	-0.36	220-295	2	-0.00			can be correct, CH··· $\pi$ contacts, weak CH···O contacts
KOYRUN	-0.26	10-100	2	-0.00		0	can be correct, paper also gives powder data at 245 K
							which proves uniaxial thermal expansion along c, H-
							bonds
EZUWAX	-0.21	120-293	2	-0.00			can be correct, H-bonds, dihydrogen contacts
WEJZUH	-0.21	175-293	2	-0.00			can be correct, H-bonds, weak CH…O contacts, CH…
							π contacts
CLAHMB	-0.00	123-423	2	-0.00			probably not correct – identical cell parameters at
							123 and 423 K

Notes: The thermal expansion coefficient, temperature range, Nm and the NTE capacity parameter  $\chi\alpha$  are provided for each individual CSD entry. The column Ano indicates whether there is a <Ueq>s anomaly, where higher temperatures correspond to lower values, and such entries are denoted with an 's'. Datasets collected in the same study are denoted with an 'o' in the SS column and other general remarks on the structure are included as well. The comments column describes some structural features and the possibility whether the negative volumetric expansion is correct or not.