

Tetragonal polymorph of 5,5-dichlorobarbituric acid

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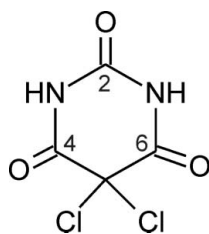
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Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.020; wR factor = 0.050; data-to-parameter ratio = 12.2.

The tetragonal polymorph of 5,5-dichlorobarbituric acid (m.p. 478 K), $\text{C}_4\text{H}_2\text{Cl}_2\text{N}_2\text{O}_3$, forms an $\text{N}-\text{H}\cdots\text{O}$ hydrogen-bonded tape structure along [001]. Two tapes related by a twofold rotation axis are associated *via* $\text{Cl}\cdots\text{O}$ contacts [3.201 (1) Å], and four such chain pairs are arranged around a fourfold roto-inversion axis. The crystal structures of the monoclinic and orthorhombic polymorphs have been reported previously [Gelbrich *et al.* (2011). *CrystEngComm*, **13**, 5502–5509].

Related literature

The polymorphic nature of 5,5-dichlorobarbituric acid was mentioned in Groth's compendium on the chemical crystallography of organic compounds, published more than a hundred years ago (Groth, 1910). For the monoclinic and orthorhombic polymorphs, see: Gelbrich *et al.* (2011). For related structures, see: Gartland & Craven (1971); Gelbrich *et al.* (2007, 2010, 2010a,b); Nichol & Clegg (2007); Zencirci *et al.* (2009, 2010); DesMarteau *et al.* (1994). For a description of the synthesis, see: Ziegler *et al.* (1962). For hydrogen-bond motifs, see: Bernstein *et al.* (1995); Etter *et al.* (1990).



Experimental

Crystal data

$\text{C}_4\text{H}_2\text{Cl}_2\text{N}_2\text{O}_3$
 $M_r = 196.98$
Tetragonal, $P4_21c$
 $a = 13.8883$ (3) Å
 $c = 6.9126$ (2) Å
 $V = 1333.34$ (6) Å³

$Z = 8$
Mo $K\alpha$ radiation
 $\mu = 0.92$ mm⁻¹
 $T = 173$ K
 $0.20 \times 0.05 \times 0.05$ mm

Data collection

Oxford Diffraction Xcalibur Ruby Gemini ultra diffractometer
Absorption correction: multi-scan (*CrysAlis RED*; Oxford Diffraction, 2003)
 $T_{\min} = 0.837$, $T_{\max} = 0.955$
11025 measured reflections
1310 independent reflections
1242 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.020$
 $wR(F^2) = 0.050$
 $S = 1.07$
1310 reflections
107 parameters
2 restraints
All H-atom parameters refined
 $\Delta\rho_{\max} = 0.20$ e Å⁻³
 $\Delta\rho_{\min} = -0.16$ e Å⁻³
Absolute structure: Flack (1983), 541 Friedel pairs
Flack parameter: -0.08 (7)

Table 1
Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N3}-\text{H3}\cdots\text{O6}^i$	0.87 (1)	2.07 (1)	2.923 (2)	167 (2)
$\text{N1}-\text{H1}\cdots\text{O2}^{ii}$	0.86 (1)	2.05 (1)	2.881 (2)	165 (2)

Symmetry codes: (i) $x, y, z - 1$; (ii) $y + \frac{1}{2}, x - \frac{1}{2}, z + \frac{1}{2}$

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2003); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis RED* (Oxford Diffraction, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2351).

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supplementary materials

Acta Cryst. (2012). E68, o235-o236 [doi:10.1107/S1600536811054626]

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Comment

The polymorphic nature of 5,5-dichlorobarbituric acid (I) is already mentioned in Groth's compendium on the chemical crystallography of organic compounds, published more than a hundred years ago (Groth, 1910). As part of our wider investigation of solid state forms of barbiturates, we have recently determined the crystal structures of a monoclinic (Ia) and an orthorhombic (Ib) form (Gelbrich *et al.*, 2011), and herein we report on the tetragonal polymorph (Ic) of the title compound. The equilibrium melting points of (Ia), (Ib) and (Ic), determined by hot-stage microscopy, are 477, 490 and 478 K, respectively. All three modifications were obtained in sublimation experiments; (Ia) as plates, (Ib) as prisms and (Ic) as long needles.

The molecular structure of (Ic) is illustrated in Fig. 1. The crystal structure of (Ic) consists of N—H \cdots O=C bonded tapes (see Fig. 2) that belong to the C-3 type in the classification scheme proposed (Gelbrich *et al.*, 2011) for the H-bonded structures of 5,5-substituted derivatives of barbituric acid. By contrast, all of the other five known crystal structures of 5,5-dihalogen analogues form either an N—H \cdots O=C bonded layer (L) or a framework (F) structure (DesMarteau *et al.*, 1994; Gelbrich *et al.*, 2011). In particular, the monoclinic polymorph (Ia) and the orthorhombic form (Ib) display the layer types L-6 and L-5, respectively (Gelbrich *et al.*, 2011). C-3 tapes have been reported previously for solid forms of γ -methylamobarbital (Gartland & Craven, 1971), butobarbital (Gelbrich *et al.*, 2007), quinal barbitone (Nichol & Clegg, 2007) and alphenal (Zencirci *et al.*, 2009).

In the crystal structure of (Ic), a single tape consists of two parallel strands. Neighbouring molecules forming a single strand are N—H \cdots O=C bonded to one another *via* their C6 carbonyl groups. Two strands of a tape are linked together by a second set of N—H \cdots O=C interactions in which the C2 carbonyl group is involved. These interactions result in two independent $R^3_3(12)$ rings (Etter *et al.*, 1990; Bernstein *et al.*, 1995). The molecules of a single strand are related to one another by a translation along [001]. Additionally, the tape possesses a glide mirror plane that is oriented perpendicular to its mean plane. The C4 carbonyl group is not involved in hydrogen bonding.

The cross section of the H-bonded tape structure is somewhat bent, so that the mean planes of the two strands form an angle of 23.7 (2)°. Two neighbouring H-bonded tapes, which are related to one another by a twofold rotation, form an assembly exhibiting short intermolecular C12 \cdots O4(-y+1/2, -x+1/2, z+1/2) distances of 3.201 (1) Å. As illustrated in Fig. 3, four such two-tape assemblies are situated around the fourfold roto-inversion axis in such a way that the C11 sites of four neighbouring molecules are the vertices of an almost ideal tetrahedron, whose edges are the intermolecular contacts C11 \cdots C11(-y+1, x, -z+1) and C11 \cdots C11(-x+1, -y+1, z) of 3.3873 (8) Å and 3.4462 (9) Å, respectively.

The three polymorphs of (I) can be readily distinguished from each other by their FT—IR spectra, which are depicted in Fig. 4. The calculated densities (Mg m⁻³) at -100 K for the three polymorphs (Ia), (Ib) and (Ic) are 1.984, 1.842 and 1.963, respectively. Therefore, the order of decreasing densities is (Ia) > (Ic) >> (Ib). The density of the tetragonal form (Ic) is 1% lower than that of the monoclinic form (Ia) and 6% higher than that of the orthorhombic polymorph (Ib), which is also the form of (I) with the most complex H-bonded structure.

Experimental

Needle-shaped crystals of (Ic) were obtained in a sublimation experiment carried out at 473 K. On heating, (Ic) undergoes a transformation into (Ib). However, the melting of (Ic) can be observed in a thermomicroscopic experiment if the crystals are placed on a hot stage that is preheated to just below the melting temperature of (Ic).

The FT—IR spectrum of (Ic) (see Fig. 4) shows a strong and sharp N—H vibration at 3258 cm^{-1} and a weak one at 3152 cm^{-1} . In the C=O region the spectrum exhibits a weaker band at 1756 cm^{-1} with a shoulder at about 1769 cm^{-1} and a stronger band at 1729 cm^{-1} . These characteristics are consistent with the G5b-type spectrum in the IR classification scheme for barabitarates (Zencirci *et al.*, 2009). This type indicates the presence of the H-bonded tape connectivity C-3. Previous G5b examples include form I of alphenal and the metastable polymorph VIII of phenobarbital (Zencirci *et al.*, 2009).

Refinement

The NH H-atoms were located in a difference Fourier map. They were refined with a distance restraint: N—H = $0.88(1)\text{ \AA}$, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$.

Figures

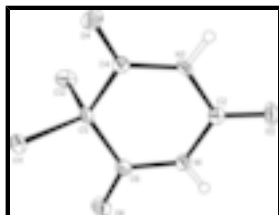


Fig. 1. The molecular structure of the title compound, (Ic), with atom numbering. Displacement ellipsoids are drawn at the 50% probability level, with hydrogen atoms shown as spheres of arbitrary size.

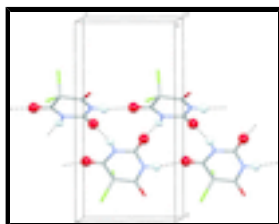


Fig. 2. A view along the *b* axis of the H-bonded C-3 tape structure of (Ic). H, O and N atoms directly involved in N—H...O interactions (dashed lines) are drawn as balls.

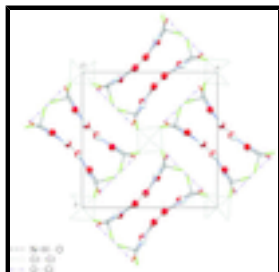


Fig. 3. A portion of the crystal structure of (Ic), view along [001] (translation direction of the C-3 tapes) with four pairs of tapes. The N—H...O bonds and short intermolecular Cl...Cl and Cl...O contacts are indicated.

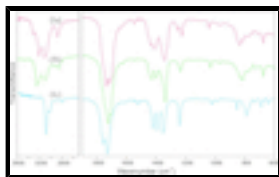


Fig. 4. FT—IR spectra of the polymorphs (Ia), (Ib) and (Ic) of 5,5-dichlorobarbituric acid.

5,5-dichlorobarbituric acid

Crystal data

$C_4H_2Cl_2N_2O_3$

$M_r = 196.98$

Tetragonal, $P4_21c$

Hall symbol: P -4 2n

$a = 13.8883 (3) \text{ \AA}$

$c = 6.9126 (2) \text{ \AA}$

$V = 1333.34 (6) \text{ \AA}^3$

$Z = 8$

$F(000) = 784$

$D_x = 1.963 \text{ Mg m}^{-3}$

Melting point: 478 K

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 5173 reflections

$\theta = 2.9\text{--}29.3^\circ$

$\mu = 0.92 \text{ mm}^{-1}$

$T = 173 \text{ K}$

Needle, colourless

$0.20 \times 0.05 \times 0.05 \text{ mm}$

Data collection

Oxford Diffraction Xcalibur Ruby Gemini ultra diffractometer

Radiation source: Enhance Ultra (Mo) X-ray Source mirror

ω scans

Absorption correction: multi-scan (*CrysAlis RED*; Oxford Diffraction, 2003)

$T_{\min} = 0.837$, $T_{\max} = 0.955$

11025 measured reflections

1310 independent reflections

1242 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.041$

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 2.9^\circ$

$h = -17 \rightarrow 17$

$k = -16 \rightarrow 17$

$l = -7 \rightarrow 8$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.020$

$wR(F^2) = 0.050$

$S = 1.07$

1310 reflections

107 parameters

2 restraints

Primary atom site location: structure-invariant direct methods

Hydrogen site location: inferred from neighbouring sites

All H-atom parameters refined

$w = 1/[\sigma^2(F_o^2) + (0.0266P)^2 + 0.2183P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.20 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.16 \text{ e \AA}^{-3}$

Extinction correction: *SHELXL97* (Sheldrick, 2008),

$F_c^* = kF_c[1 + 0.001 \times F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0046 (8)

Absolute structure: Flack (1983), **541 Friedel pairs**

supplementary materials

Secondary atom site location: difference Fourier map Flack parameter: -0.08 (7)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.40679 (3)	0.41812 (3)	0.32982 (7)	0.01953 (13)
C12	0.26242 (3)	0.27671 (3)	0.22840 (7)	0.01938 (13)
O2	0.53042 (10)	0.10721 (9)	-0.1061 (2)	0.0225 (3)
O4	0.37178 (9)	0.39086 (9)	-0.0966 (2)	0.0178 (3)
O6	0.44620 (11)	0.22096 (11)	0.4808 (2)	0.0260 (4)
N1	0.48923 (11)	0.16639 (11)	0.1869 (2)	0.0158 (3)
N3	0.44480 (11)	0.24555 (11)	-0.0993 (2)	0.0127 (3)
C2	0.49068 (12)	0.16895 (13)	-0.0125 (3)	0.0134 (4)
C4	0.40209 (13)	0.32207 (12)	-0.0102 (3)	0.0120 (4)
C5	0.38637 (12)	0.30971 (12)	0.2083 (3)	0.0131 (4)
C6	0.44390 (12)	0.22935 (13)	0.3078 (3)	0.0148 (4)
H1	0.5202 (13)	0.1184 (10)	0.233 (3)	0.018*
H3	0.4508 (14)	0.2465 (15)	-0.2243 (14)	0.018*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0247 (2)	0.0140 (2)	0.0199 (2)	0.0036 (2)	-0.0065 (2)	-0.00722 (19)
C12	0.0156 (2)	0.0263 (2)	0.0162 (2)	-0.00433 (19)	0.00163 (18)	0.00387 (19)
O2	0.0286 (8)	0.0197 (7)	0.0193 (7)	0.0098 (6)	0.0067 (6)	-0.0037 (6)
O4	0.0212 (7)	0.0149 (7)	0.0172 (7)	0.0031 (5)	0.0002 (6)	0.0056 (6)
O6	0.0414 (9)	0.0266 (8)	0.0100 (7)	0.0112 (7)	-0.0030 (6)	0.0013 (6)
N1	0.0221 (8)	0.0124 (7)	0.0128 (8)	0.0072 (6)	-0.0019 (7)	0.0002 (7)
N3	0.0165 (8)	0.0151 (8)	0.0065 (7)	0.0004 (6)	0.0010 (6)	-0.0011 (7)
C2	0.0122 (9)	0.0134 (9)	0.0148 (10)	-0.0017 (7)	0.0013 (8)	0.0001 (8)
C4	0.0091 (9)	0.0132 (9)	0.0136 (9)	-0.0027 (7)	-0.0006 (8)	-0.0006 (7)
C5	0.0133 (8)	0.0130 (8)	0.0129 (9)	0.0006 (7)	0.0007 (7)	-0.0046 (7)
C6	0.0173 (9)	0.0140 (9)	0.0130 (10)	-0.0002 (7)	-0.0021 (7)	-0.0006 (8)

Geometric parameters (\AA , $^\circ$)

C11—C5	1.7471 (18)	N1—H1	0.856 (9)
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Cl2—C5	1.7868 (18)	N3—C4	1.364 (2)
O2—C2	1.208 (2)	N3—C2	1.378 (2)
O4—C4	1.203 (2)	N3—H3	0.868 (9)
O6—C6	1.202 (2)	C4—C5	1.536 (3)
N1—C6	1.363 (2)	C5—C6	1.535 (2)
N1—C2	1.379 (2)		
C6—N1—C2	127.07 (17)	N3—C4—C5	114.75 (16)
C6—N1—H1	120.1 (14)	C6—C5—C4	116.60 (15)
C2—N1—H1	112.9 (14)	C6—C5—C11	109.07 (12)
C4—N3—C2	127.31 (17)	C4—C5—C11	110.70 (13)
C4—N3—H3	118.6 (14)	C6—C5—C12	106.27 (12)
C2—N3—H3	113.6 (14)	C4—C5—C12	104.00 (12)
O2—C2—N3	121.74 (17)	C11—C5—C12	109.88 (10)
O2—C2—N1	121.60 (18)	O6—C6—N1	122.37 (18)
N3—C2—N1	116.65 (16)	O6—C6—C5	122.00 (17)
O4—C4—N3	123.15 (18)	N1—C6—C5	115.60 (16)
O4—C4—C5	121.84 (17)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N3—H3 \cdots O6 ⁱ	0.87 (1)	2.07 (1)	2.923 (2)	167.(2)
N1—H1 \cdots O2 ⁱⁱ	0.86 (1)	2.05 (1)	2.881 (2)	165 (2)

Symmetry codes: (i) *x*, *y*, *z*-1; (ii) *y*+1/2, *x*-1/2, *z*+1/2.

Fig. 1

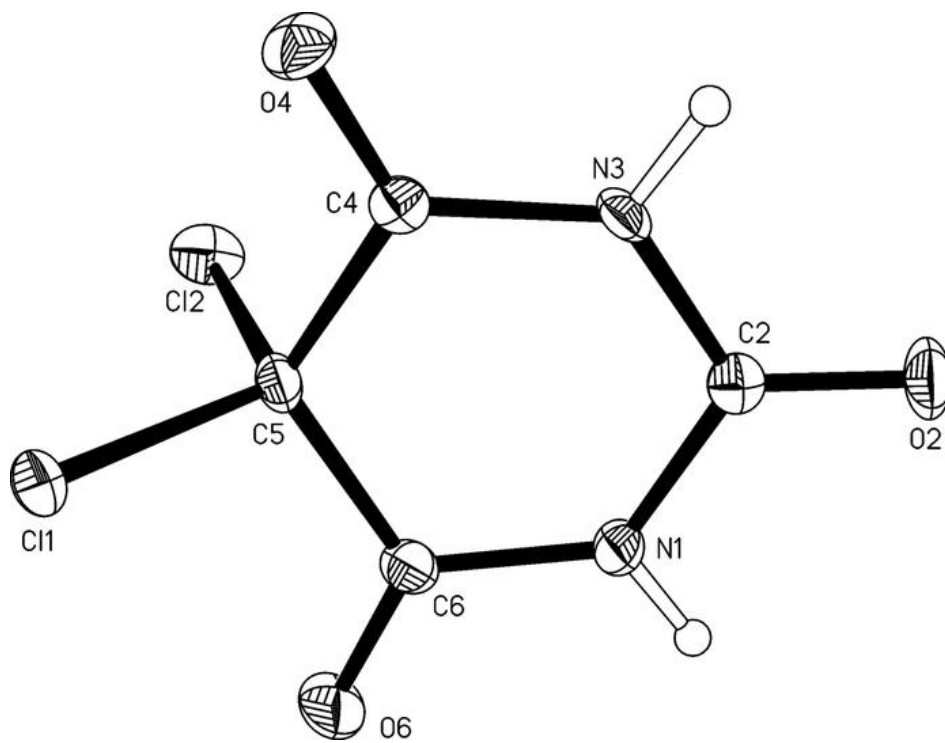


Fig. 2

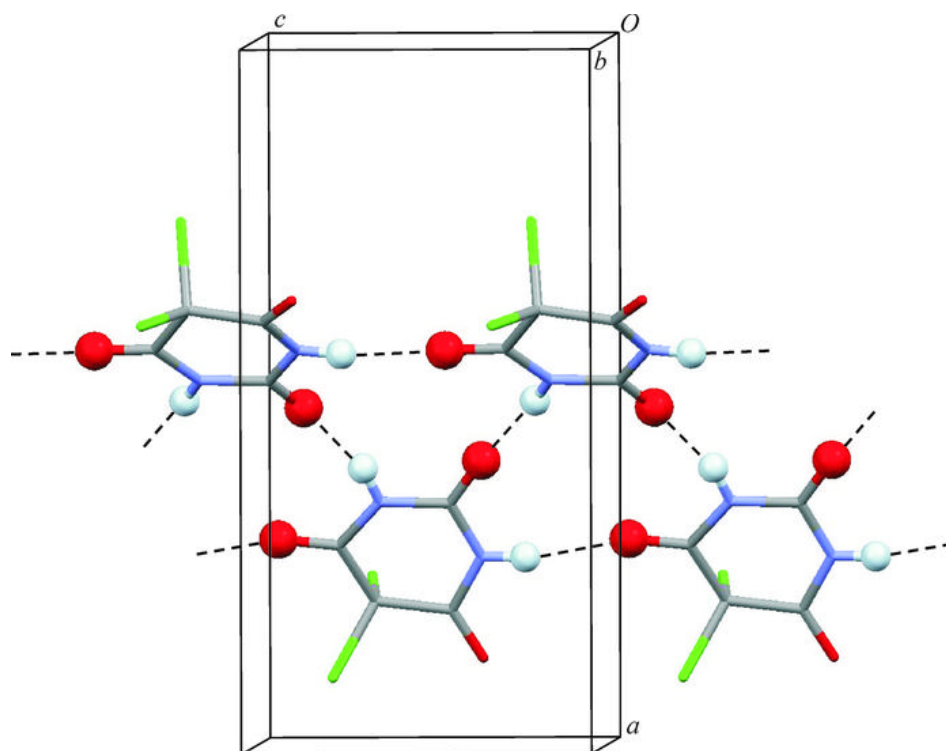


Fig. 3

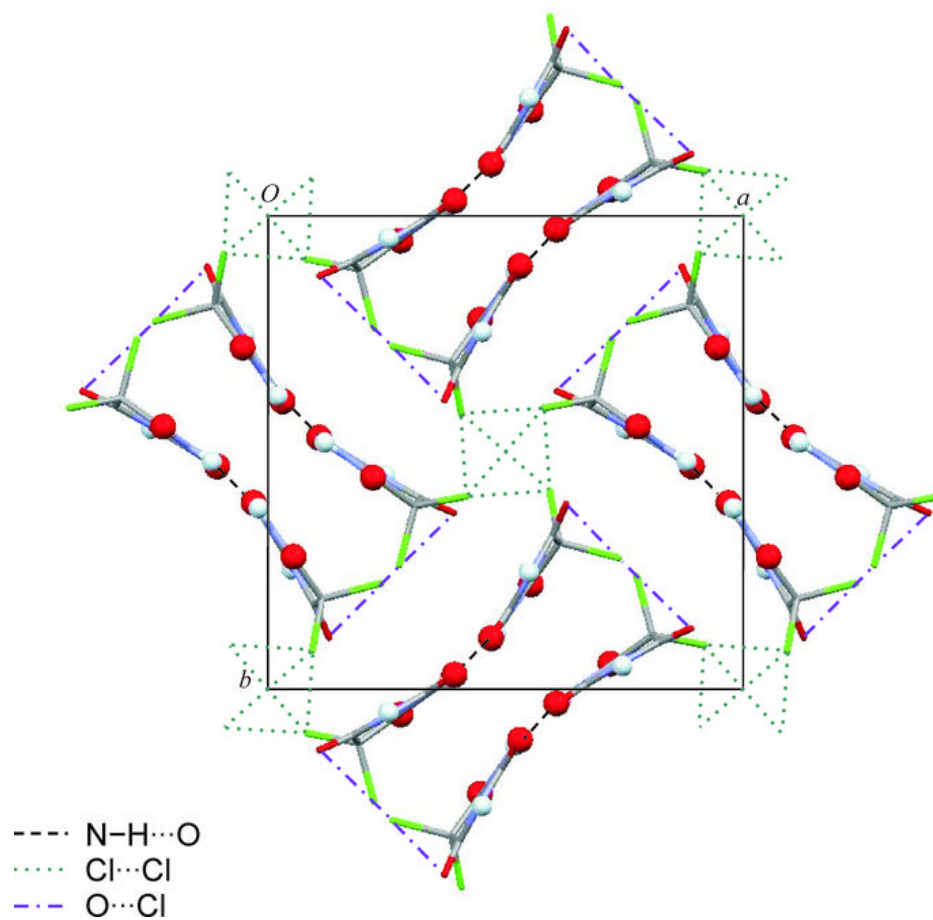


Fig. 4

