organic compounds

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4-Carbamoylpyridin-1-ium 2,2,2-trichloroacetate—isonicotinamide (1/1)

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Key indicators: single-crystal X-ray study; T = 293 K; mean σ (C–C) = 0.004 Å; R factor = 0.040; wR factor = 0.091; data-to-parameter ratio = 16.7.

In the crystal structure of the title 1:1 co-crystal, $C_6H_7N_2O^+ \cdot C_2Cl_3O_2^- \cdot C_6H_6N_2O$, the amide groups of the 4carbamoylpyridin-1-ium ion and the isonicotinamide molecule are twisted out of the plane of the aromatic ring with C-C-C-N torsion angles of 21.5 (4) and -33.5 (4)°, respectively. The 4-carbamoylpyridin-1-ium and isonicotinamide amide groups form $R_2^2(8)$ hydrogen-bonded dimers *via* N- $H \cdot \cdot \cdot O$ =C interactions. The two remaining amide H atoms (i) link dimers *via* the cation to an isonicotinamide and (ii) from the isonicotinamide to a trichloroacetate anion. The pyridinium H atom also forms an N-H···O hydrogen bond with the trichloroacetate anion. Due to the extended hydrogen bonding, including C-H···O and C-H···Cl interactions, all components in the structure aggregate into a three-dimensional supramolecular framework.

Related literature

For applications of co-crystals, see: Karki *et al.* (2009); Friščić & Jones (2010). For related structures, see: Madeley *et al.* (2011).



Experimental

Crystal data $C_6H_7N_2O^+ \cdot C_2CI_3O_2^- \cdot C_6H_6N_2O$ $M_r = 407.63$ Orthorhombic, *Pna2*₁ a = 13.7910 (3) Å b = 22.6680 (5) Å c = 5.6340 (1) Å

 $V = 1761.27 \text{ (6) } \text{Å}^{3}$ Z = 4Mo K\alpha radiation $\mu = 0.55 \text{ mm}^{-1}$ T = 293 K $0.4 \times 0.1 \times 0.1 \text{ mm}$



Data collection

Agilent SuperNova, Dual, Cu at zero, Atlas diffractometer Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2011) $T_{min} = 0.811, T_{max} = 0.947$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.04$	
$wR(F^2) = 0.091$	
S = 1.04	
4017 reflections	
241 parameters	
1 restraint	

16297 measured reflections 4017 independent reflections 3575 reflections with $I > 2\sigma(I)$ $R_{int} = 0.031$

H atoms treated by a mixture of independent and constrained refinement
$$\begin{split} &\Delta\rho_{max}=0.41\ e\ \mathring{A}^{-3}\\ &\Delta\rho_{min}=-0.57\ e\ \mathring{A}^{-3}\\ &Absolute\ structure:\ Flack\ (1983),\\ &1791\ Friedel\ pairs\\ &Flack\ parameter:\ 0.01\ (6) \end{split}$$

Table 1 Hydrogen-bond geometry (Å, $^\circ).$

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1 - H15 \cdots O2^{i}$	0.90 (3)	1.78 (3)	2.679 (3)	175 (3)
$N2-H16A\cdots N3^{ii}$	0.87 (3)	2.11 (3)	2.958 (3)	164 (3)
$N2-H16B\cdots O3$	0.90 (4)	1.99 (4)	2.887 (3)	178 (3)
$N4-H17A\cdots O4$	0.91 (4)	2.08 (4)	2.972 (3)	167 (4)
$N4-H17B\cdots O1$	0.89 (4)	1.98 (4)	2.839 (3)	160 (3)
$C1 - H1 \cdot \cdot \cdot O1^{i}$	0.93	2.58	3.211 (3)	126
$C2-H2 \cdot \cdot \cdot O4^{iii}$	0.93	2.55	3.358 (3)	146
$C7 - H7 \cdot \cdot \cdot O3^{iv}$	0.93	2.58	3.489 (3)	166
$C11 - H11 \cdots Cl2^{v}$	0.93	2.82	3.711 (3)	162

Symmetry codes: (i) $-x, -y + 1, z + \frac{3}{2}$, (ii) $x - \frac{1}{2}, -y + \frac{3}{2}, z + 1$; (iii) $-x, -y + 1, z + \frac{1}{2}$, (iv) $x + \frac{1}{2}, -y + \frac{3}{2}, z$; (v) $x - \frac{1}{2}, -y + \frac{3}{2}, z - 1$.

Data collection: CrysAlis PRO (Agilent, 2011); cell refinement: CrysAlis PRO; data reduction: CrysAlis PRO; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997) and DIAMOND (Brandenburg, 1999); software used to prepare material for publication: WinGX (Farrugia, 1999) and publCIF (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GG2095).

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4-Carbamoylpyridin-1-ium 2,2,2-trichloroacetate-isonicotinamide (1/1)

Franc Perdih

S1. Comment

Co-crystals have attracted much attention in recent years owing to their contributions to crystal engineering and pharmaceutical chemistry. They were found to be useful in improving the stability, solubility, dissolution rate and mechanical properties (Karki *et al.*, 2009; Friščić & Jones, 2010). Here we present the structure obtained by reacting isonicotinamide and trichloroacetic acid in 2:1 molar ratio.

The asymmetric unit of (I) consists of one 4-carbamoylpyridin-1-ium cation, one trichloroacetate anion and one isonicotinamide molecule (Fig. 1). The amide groups of 4-carbamoylpyridin-1-ium ion and isonicotinamide molecule are twisted out of the plane of the aromatic ring with a C—C—C—N torsion angle of 21.5 (4)° and -33.5 (4)°, respectively. Similar twisting was observed for example in isonicotinamide–2-naphthoic acid (1/1) (Madeley *et al.*, 2011). Aromatic rings of 4-carbamoylpyridin-1-ium ion and isonicotinamide molecule are not coplanar, but are inclined by 35.05 (12)°. In the crystal, all the components of the structure are associated *via* the extended system of hydrogen bonds (N—H…O and N—H…N) and weak C—H…O and C—H…Cl interactions into extended three-dimensional supramolecular framework (Figs. 2, 3). The 4-carbamoylpyridin-1-ium ion is hydrogen bonded *via* N—H…O hydrogen bonding of the pyridinium unit to the trichloroacetate ion. The amide groups from 4-carbamoylpyridin-1-ium and isonicotinamide form a dimer *via* N—H…O hydrogen bonding, that is a typical supramolecular hydrogen-bonded synthon observed for amide-amide homodimers. Furthermore, the amide group of the cation is hydrogen bonded to the pyridine unit of isonicotinamide and the amide group of the isonicotinamide is hydrogen bonded to the trichloroacetate ion.

S2. Experimental

Crystals of the title compound were obtained by slow evaporation of a 2:1 mol. mixture of isonicotinamide and trichloroacetic acid in methanol at room temperature.

S3. Refinement

All H atoms were initially located in a difference Fourier maps. H atoms attached to N atoms were refined isotropically with $U_{iso}(H) = 1.5U_{eq}(N)$. Other H atoms were treated as riding atoms in geometrically idealized positions, with C—H = 0.93 Å, and with $U_{iso}(H) = 1.2U_{eq}(C)$.





The asymmetric unit of the title compound with displacement ellipsoids drawn at the 50% probability level.



Figure 2

Hydrogen bonding diagram. Dashed lines indicate intermolecular N—H···O, N—H···N, C—H···O and C—H···Cl hydrogen bonding. For the sake of clarity, H atoms not involved in the motif shown have been omitted. Symmetry codes: ⁱ -x, -y + 1, z + 3/2; ⁱⁱ x - 1/2, -y + 3/2, z + 1; ^{iv} x + 1/2, -y + 3/2, z; ^v x - 1/2, -y + 3/2, z - 1.



Figure 3

Crystal packing of the title compound. For the sake of clarity, hydrogen bonding is not presented.

4-Carbamoylpyridin-1-ium 2,2,2-trichloroacetate-isonicotinamide (1/1)

Crystal data	
$C_6H_7N_2O^+ \cdot C_2Cl_3O_2^- \cdot C_6H_6N_2O$	F(000) = 832
$M_r = 407.63$	$D_{\rm x} = 1.537 { m Mg} { m m}^{-3}$
Orthorhombic, <i>Pna</i> 2 ₁	Mo <i>K</i> α radiation, $\lambda = 0.71073$ Å
Hall symbol: P 2c -2n	Cell parameters from 7898 reflections
a = 13.7910(3) Å	$\theta = 3.1 - 30.4^{\circ}$
b = 22.6680 (5) Å	$\mu=0.55~\mathrm{mm^{-1}}$
c = 5.6340(1) Å	T = 293 K
V = 1761.27 (6) Å ³	Prism, colourless
Z = 4	$0.4 \times 0.1 \times 0.1 \text{ mm}$

Data collection

Agilent SuperNova, Dual, Cu at zero, Atlas	$T_{\min} = 0.811, T_{\max} = 0.947$
diffractometer	16297 measured reflections
Radiation source: SuperNova (Mo) X-ray	4017 independent reflections
Source	3575 reflections with $I > 2\sigma(I)$
Mirror monochromator	$R_{\text{int}} = 0.031$
Detector resolution: 10.4933 pixels mm ⁻¹	$\theta_{\text{max}} = 27.5^{\circ}, \theta_{\text{min}} = 3.1^{\circ}$
ω scans	$h = -17 \rightarrow 17$
Absorption correction: multi-scan	$k = -29 \rightarrow 29$
(<i>CrysAlis PRO</i> ; Agilent, 2011)	$l = -7 \rightarrow 7$
Refinement	
Refinement on F^2	Hydrogen site location: inferred from
Least-squares matrix: full	neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.04$	H atoms treated by a mixture of independent
$wR(F^2) = 0.091$	and constrained refinement
S = 1.03	$w = 1/[\sigma^2(F_o^2) + (0.0359P)^2 + 0.8943P]$
4017 reflections	where $P = (F_o^2 + 2F_c^2)/3$
241 parameters	$(\Delta/\sigma)_{max} < 0.001$
1 restraint	$\Delta\rho_{max} = 0.41$ e Å ⁻³
Primary atom site location: structure-invariant	$\Delta\rho_{min} = -0.57$ e Å ⁻³
direct methods	Absolute structure: Flack (1983), 1791 Friedel
Secondary atom site location: difference Fourier	pairs
map	Absolute structure parameter: 0.01 (6)

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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 $F^2 > 2\sigma(F^2)$ 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 $(Å^2)$

	r	12	7	11*/11	
	<i>A</i>	<i>y</i>	2		
CII	0.49348 (5)	0.58322 (3)	0.45563 (14)	0.04547 (17)	
C12	0.44990 (9)	0.67585 (4)	0.11741 (18)	0.0850 (4)	
C13	0.32105 (7)	0.65069 (5)	0.50880 (15)	0.0770 (3)	
N1	-0.29717 (16)	0.50785 (9)	1.1681 (4)	0.0377 (5)	
H15	-0.330 (2)	0.4892 (15)	1.285 (6)	0.057*	
N2	-0.16743 (16)	0.62843 (10)	0.4929 (5)	0.0396 (5)	
H16A	-0.217 (2)	0.6478 (14)	0.551 (7)	0.059*	
H16B	-0.135 (3)	0.6394 (15)	0.362 (6)	0.059*	
N3	0.19060 (17)	0.78523 (10)	-0.3220 (5)	0.0425 (5)	
N4	0.06680 (19)	0.61873 (11)	0.1998 (5)	0.0495 (7)	
H17A	0.030 (3)	0.5952 (17)	0.294 (8)	0.074*	
H17B	0.131 (3)	0.6133 (15)	0.197 (7)	0.074*	
01	0.25905 (13)	0.57929 (9)	0.1146 (4)	0.0524 (5)	
O2	0.40234 (13)	0.54301 (8)	0.0115 (4)	0.0426 (4)	

03	-0.06660 (12)	0.66646 (9)	0.0707 (4)	0.0473 (5)	
O4	-0.05512 (13)	0.55797 (8)	0.5595 (4)	0.0488 (5)	
C1	-0.20142 (19)	0.49801 (11)	1.1584 (5)	0.0384 (6)	
H1	-0.1721	0.4742	1.2722	0.046*	
C2	-0.14636 (17)	0.52287 (10)	0.9812 (5)	0.0356 (5)	
H2	-0.0801	0.5155	0.9729	0.043*	
C3	-0.19079 (17)	0.55906 (10)	0.8150 (4)	0.0293 (5)	
C4	-0.29018 (18)	0.56864 (11)	0.8322 (5)	0.0335 (5)	
H4	-0.3214	0.5929	0.7233	0.04*	
C5	-0.34193 (18)	0.54211 (11)	1.0107 (5)	0.0387 (6)	
Н5	-0.4085	0.5481	1.0217	0.046*	
C6	-0.13130 (17)	0.58311 (10)	0.6111 (5)	0.0320 (5)	
C7	0.2246 (2)	0.76108 (12)	-0.1226 (6)	0.0454 (7)	
H7	0.2848	0.7735	-0.0675	0.055*	
C8	0.17586 (17)	0.71890 (11)	0.0064 (5)	0.0396 (6)	
H8	0.2034	0.7027	0.1424	0.048*	
C9	0.08482 (16)	0.70091 (10)	-0.0701 (5)	0.0305 (5)	
C10	0.0493 (2)	0.72548 (12)	-0.2753 (5)	0.0384 (6)	
H10	-0.0114	0.7145	-0.3326	0.046*	
C11	0.1043 (2)	0.76648 (11)	-0.3955 (5)	0.0433 (6)	
H11	0.0797	0.7819	-0.5361	0.052*	
C12	0.02224 (18)	0.65981 (11)	0.0724 (5)	0.0358 (6)	
C13	0.34726 (17)	0.57605 (10)	0.1255 (4)	0.0306 (5)	
C14	0.40010 (19)	0.61971 (11)	0.2972 (5)	0.0368 (6)	

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0384 (3)	0.0586 (4)	0.0395 (3)	0.0069 (3)	-0.0118 (3)	0.0020 (3)
Cl2	0.1224 (9)	0.0644 (5)	0.0681 (6)	-0.0526 (6)	-0.0447 (6)	0.0302 (5)
Cl3	0.0848 (6)	0.0950 (6)	0.0510 (5)	0.0508 (5)	-0.0198 (4)	-0.0362 (5)
N1	0.0414 (12)	0.0353 (11)	0.0365 (13)	-0.0078 (9)	0.0084 (10)	0.0028 (9)
N2	0.0375 (12)	0.0379 (11)	0.0435 (13)	0.0082 (9)	0.0147 (11)	0.0077 (11)
N3	0.0423 (13)	0.0351 (11)	0.0503 (14)	-0.0051 (9)	0.0080 (11)	0.0057 (10)
N4	0.0317 (12)	0.0490 (14)	0.0678 (17)	0.0069 (10)	0.0141 (12)	0.0256 (13)
O1	0.0277 (9)	0.0589 (12)	0.0706 (14)	0.0023 (8)	-0.0027 (10)	-0.0192 (12)
O2	0.0346 (9)	0.0506 (10)	0.0426 (10)	0.0042 (8)	-0.0017 (8)	-0.0179 (9)
O3	0.0278 (9)	0.0557 (11)	0.0585 (13)	0.0039 (8)	0.0068 (9)	0.0201 (10)
O4	0.0367 (10)	0.0514 (11)	0.0585 (14)	0.0153 (8)	0.0200 (9)	0.0148 (10)
C1	0.0444 (15)	0.0377 (13)	0.0332 (15)	0.0002 (11)	-0.0051 (11)	0.0055 (11)
C2	0.0307 (11)	0.0348 (12)	0.0413 (14)	0.0014 (9)	0.0001 (12)	0.0002 (12)
C3	0.0297 (12)	0.0279 (11)	0.0304 (12)	-0.0023 (9)	0.0021 (10)	-0.0034 (9)
C4	0.0300 (13)	0.0340 (13)	0.0364 (13)	0.0012 (10)	0.0031 (11)	0.0044 (10)
C5	0.0347 (13)	0.0381 (13)	0.0435 (14)	-0.0010 (10)	0.0099 (12)	0.0013 (12)
C6	0.0295 (12)	0.0343 (12)	0.0321 (12)	-0.0013 (9)	0.0071 (10)	-0.0003 (11)
C7	0.0327 (14)	0.0460 (16)	0.0577 (18)	-0.0081 (12)	-0.0007 (12)	0.0027 (14)
C8	0.0317 (12)	0.0450 (14)	0.0422 (15)	0.0006 (10)	-0.0029 (11)	0.0061 (13)
C9	0.0288 (11)	0.0289 (11)	0.0337 (12)	0.0018 (9)	0.0050 (10)	0.0020 (10)

supporting information

C10	0.0337 (13)	0.0412 (14)	0.0404 (14)	-0.0027 (11)	-0.0053 (11)	0.0025 (12)	
C11	0.0490 (15)	0.0441 (14)	0.0369 (14)	0.0005 (12)	-0.0016 (13)	0.0113 (13)	
C12	0.0313 (13)	0.0343 (12)	0.0418 (15)	0.0018 (10)	0.0064 (11)	0.0060 (11)	
C13	0.0323 (12)	0.0325 (11)	0.0269 (11)	-0.0016 (9)	-0.0010 (10)	-0.0008 (10)	
C14	0.0435 (15)	0.0342 (14)	0.0327 (12)	0.0039 (11)	-0.0083 (11)	-0.0023 (11)	

Geometric parameters (Å, °)

Cl1—C14	1.772 (3)	C1—H1	0.93	
Cl2—C14	1.765 (3)	C2—C3	1.387 (4)	
Cl3—C14	1.762 (3)	C2—H2	0.93	
N1—C5	1.331 (4)	C3—C4	1.391 (3)	
N1—C1	1.340 (3)	C3—C6	1.513 (3)	
N1—H15	0.90 (3)	C4—C5	1.372 (4)	
N2—C6	1.322 (3)	C4—H4	0.93	
N2—H16A	0.87 (3)	С5—Н5	0.93	
N2—H16B	0.90 (4)	C7—C8	1.376 (4)	
N3—C11	1.330 (4)	C7—H7	0.93	
N3—C7	1.335 (4)	C8—C9	1.389 (3)	
N4—C12	1.326 (3)	C8—H8	0.93	
N4—H17A	0.91 (4)	C9—C10	1.374 (4)	
N4—H17B	0.89 (4)	C9—C12	1.503 (3)	
O1—C13	1.220 (3)	C10—C11	1.378 (4)	
O2—C13	1.245 (3)	C10—H10	0.93	
O3—C12	1.234 (3)	C11—H11	0.93	
O4—C6	1.230 (3)	C13—C14	1.564 (3)	
C1—C2	1.375 (4)			
C5—N1—C1	121.8 (2)	N3—C7—C8	123.9 (3)	
C5—N1—H15	122 (2)	N3—C7—H7	118	
C1—N1—H15	116 (2)	C8—C7—H7	118	
C6—N2—H16A	120 (2)	C7—C8—C9	118.8 (3)	
C6—N2—H16B	116 (2)	C7—C8—H8	120.6	
H16A—N2—H16B	124 (3)	C9—C8—H8	120.6	
C11—N3—C7	116.4 (2)	C10—C9—C8	117.7 (2)	
C12—N4—H17A	118 (2)	C10—C9—C12	119.7 (2)	
C12—N4—H17B	123 (2)	C8—C9—C12	122.4 (2)	
H17A—N4—H17B	119 (3)	C9—C10—C11	119.4 (2)	
N1-C1-C2	120.3 (2)	C9—C10—H10	120.3	
N1-C1-H1	119.8	C11—C10—H10	120.3	
C2—C1—H1	119.8	N3—C11—C10	123.7 (3)	
C1—C2—C3	119.2 (2)	N3—C11—H11	118.1	
C1—C2—H2	120.4	C10—C11—H11	118.1	
C3—C2—H2	120.4	O3—C12—N4	123.4 (2)	
C2—C3—C4	118.7 (2)	O3—C12—C9	119.3 (2)	
C2—C3—C6	119.1 (2)	N4—C12—C9	117.3 (2)	
C4—C3—C6	122.1 (2)	O1—C13—O2	128.2 (2)	
C5—C4—C3	119.7 (2)	O1—C13—C14	117.2 (2)	

С5—С4—Н4	120.2	O2—C13—C14	114.6 (2)
C3—C4—H4	120.2	C13—C14—Cl3	112.49 (18)
N1—C5—C4	120.2 (2)	C13—C14—Cl2	106.43 (18)
N1—C5—H5	119.9	Cl3—C14—Cl2	109.97 (15)
С4—С5—Н5	119.9	C13—C14—Cl1	110.78 (17)
O4—C6—N2	124.3 (2)	Cl3—C14—Cl1	107.15 (15)
O4—C6—C3	118.4 (2)	Cl2—C14—Cl1	110.04 (15)
N2—C6—C3	117.3 (2)		
C5—N1—C1—C2	-0.8 (4)	C7—C8—C9—C12	-173.6 (2)
N1—C1—C2—C3	1.1 (4)	C8—C9—C10—C11	-0.1 (4)
C1—C2—C3—C4	-0.5 (4)	C12—C9—C10—C11	175.2 (2)
C1—C2—C3—C6	-176.0 (2)	C7—N3—C11—C10	1.5 (4)
C2—C3—C4—C5	-0.4 (4)	C9—C10—C11—N3	-1.5 (4)
C6—C3—C4—C5	175.0 (2)	C10—C9—C12—O3	-30.3 (4)
C1—N1—C5—C4	-0.1 (4)	C8—C9—C12—O3	144.8 (3)
C3—C4—C5—N1	0.7 (4)	C10—C9—C12—N4	151.4 (3)
C2—C3—C6—O4	20.0 (4)	C8—C9—C12—N4	-33.5 (4)
C4—C3—C6—O4	-155.4 (3)	O1—C13—C14—Cl3	17.9 (3)
C2—C3—C6—N2	-163.1 (2)	O2—C13—C14—Cl3	-163.93 (19)
C4—C3—C6—N2	21.5 (4)	O1—C13—C14—Cl2	-102.6 (3)
C11—N3—C7—C8	0.1 (4)	O2-C13-C14-Cl2	75.6 (2)
N3—C7—C8—C9	-1.6 (4)	O1-C13-C14-Cl1	137.8 (2)
C7—C8—C9—C10	1.5 (4)	O2-C13-C14-Cl1	-44.0 (3)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H··· <i>A</i>	D—H	H···A	D····A	D—H··· A
N1—H15…O2 ⁱ	0.90 (3)	1.78 (3)	2.679 (3)	175 (3)
N2—H16A…N3 ⁱⁱ	0.87 (3)	2.11 (3)	2.958 (3)	164 (3)
N2—H16 <i>B</i> ···O3	0.90 (4)	1.99 (4)	2.887 (3)	178 (3)
N4—H17 <i>A</i> …O4	0.91 (4)	2.08 (4)	2.972 (3)	167 (4)
N4—H17 <i>B</i> …O1	0.89 (4)	1.98 (4)	2.839 (3)	160 (3)
C1—H1···O1 ⁱ	0.93	2.58	3.211 (3)	126
C2—H2···O4 ⁱⁱⁱ	0.93	2.55	3.358 (3)	146
C7—H7···O3 ^{iv}	0.93	2.58	3.489 (3)	166
C11—H11····Cl2 ^v	0.93	2.82	3.711 (3)	162

Symmetry codes: (i) -*x*, -*y*+1, *z*+3/2; (ii) *x*-1/2, -*y*+3/2, *z*+1; (iii) -*x*, -*y*+1, *z*+1/2; (iv) *x*+1/2, -*y*+3/2, *z*; (v) *x*-1/2, -*y*+3/2, *z*-1.