

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

4-Carbamoylpyridin-1-ium 2,2,2-trichloroacetate

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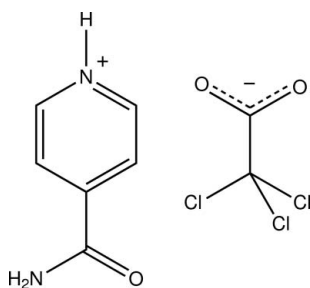
Received 9 August 2012; accepted 11 August 2012

 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.007$ Å; R factor = 0.069; wR factor = 0.199; data-to-parameter ratio = 18.0.

In the asymmetric unit of the title salt, $\text{C}_6\text{H}_7\text{N}_2\text{O}^+ \cdot \text{C}_2\text{Cl}_3\text{O}_2^-$, there are two crystallographic independent ion pairs. The amide groups of the 4-carbamoylpyridin-1-ium ions are slightly twisted out of the plane of the aromatic ring with $\text{C}-\text{C}-\text{C}-\text{N}$ torsion angles of 8.8 (9°) and 4.6 (8°). In the crystal, the 4-carbamoylpyridin-1-ium ion is $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonded to the trichloroacetate ion *via* the pyridinium unit and amide group. Layers parallel to the *ac* plane are formed due to the $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonding of the adjacent amide groups of 4-carbamoylpyridin-1-ium ions. Weak $\text{C}-\text{H} \cdots \text{O}$ interactions also occur.

Related literature

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



Experimental

Crystal data

 $\text{C}_6\text{H}_7\text{N}_2\text{O}^+ \cdot \text{C}_2\text{Cl}_3\text{O}_2^-$
 $M_r = 285.51$

 Monoclinic, *Pc*
 $a = 9.8768$ (3) Å
 $b = 9.4403$ (3) Å
 $c = 12.5157$ (3) Å
 $\beta = 90.240$ (2°)

 $V = 1166.95$ (6) Å³
 $Z = 4$

 Mo $K\alpha$ radiation
 $\mu = 0.78$ mm⁻¹
 $T = 293$ K
 $0.2 \times 0.2 \times 0.2$ mm

Data collection

 Agilent SuperNova, Dual, Cu at zero, Atlas diffractometer
 Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2011)
 $T_{\min} = 0.860$, $T_{\max} = 0.860$

 11114 measured reflections
 5195 independent reflections
 4452 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.026$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.069$
 $wR(F^2) = 0.199$
 $S = 1.04$
 5195 reflections
 289 parameters
 2 restraints

 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.53$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.47$ e Å⁻³
 Absolute structure: Flack (1983), 2512 Friedel pairs
 Flack parameter: 0.08 (12)

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{N1}-\text{H1A} \cdots \text{O2}^{\text{i}}$	0.86	1.78	2.643 (5)	176
$\text{N2}-\text{H2A} \cdots \text{O1}^{\text{ii}}$	0.86	2.03	2.883 (6)	170
$\text{N2}-\text{H2B} \cdots \text{O6}^{\text{iii}}$	0.86	2.06	2.854 (7)	152
$\text{N3}-\text{H3A} \cdots \text{O3}^{\text{iv}}$	0.86	1.8	2.661 (5)	178
$\text{N4}-\text{H4A} \cdots \text{O4}^{\text{v}}$	0.86	2.01	2.858 (6)	169
$\text{N4}-\text{H4B} \cdots \text{O5}^{\text{v}}$	0.86	2.03	2.858 (7)	160
$\text{C2}-\text{H2} \cdots \text{O6}^{\text{iii}}$	0.93	2.34	3.251 (7)	166
$\text{C5}-\text{H5} \cdots \text{O3}$	0.93	2.6	3.422 (8)	148
$\text{C8}-\text{H8} \cdots \text{O5}^{\text{v}}$	0.93	2.36	3.270 (6)	166

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

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* publication routines (Farrugia, 1999) and *pubCIF* (Westrip, 2010).

The author thanks the Ministry of Education, Science, Culture and Sport of the Republic of Slovenia and the Slovenian Research Agency for financial support through grants P1-0230-0175 as well as the EN-FIST Centre of Excellence, Dunajska 156, 1000 Ljubljana, Slovenia for use of the Supernova diffractometer.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BQ2374).

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supporting information

Acta Cryst. (2012). E68, o2733 [doi:10.1107/S1600536812035507]

4-Carbamoylpyridin-1-ium 2,2,2-trichloroacetate

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S1. Comment

Salts or co-crystals represent two possible ways to produce functional pharmaceutical materials. The salt or co-crystal formation is dependent on a pK_a difference between the acid and the base (Karki *et al.*, 2009; Friščić & Jones, 2010). Here we present the structure obtained by contacting isonicotinamide and trichloroacetic acid in 1:1 molar ratio.

The asymmetric unit of (I) consists of two crystallographic independent 4-carbamoylpyridin-1-ium cations and two trichloroacetate anions (Fig. 1). The amide groups of 4-carbamoylpyridin-1-ium ions are only slightly twisted out of the plane of the aromatic ring with a C—C—C—N torsion angle of 8.8 (9)° and 4.6 (8)°, respectively. The 4-carbamoylpyridin-1-ium ion is N—H···O hydrogen bonded to the trichloroacetate ion *via* the pyridinium unit and amide group (Fig. 2). Two-dimensional framework is formed due to the N—H···O hydrogen bonding of the adjacent amide groups of 4-carbamoylpyridin-1-ium ions. This layer formation is further stabilized by weak C—H···O interactions. In the structure of (I) typical amide-amide hydrogen-bonded homodimer is not present as is for example in the 4-carbamoylpyridin-1-ium 3-carboxypicolinate (Das & Baruah, 2011).

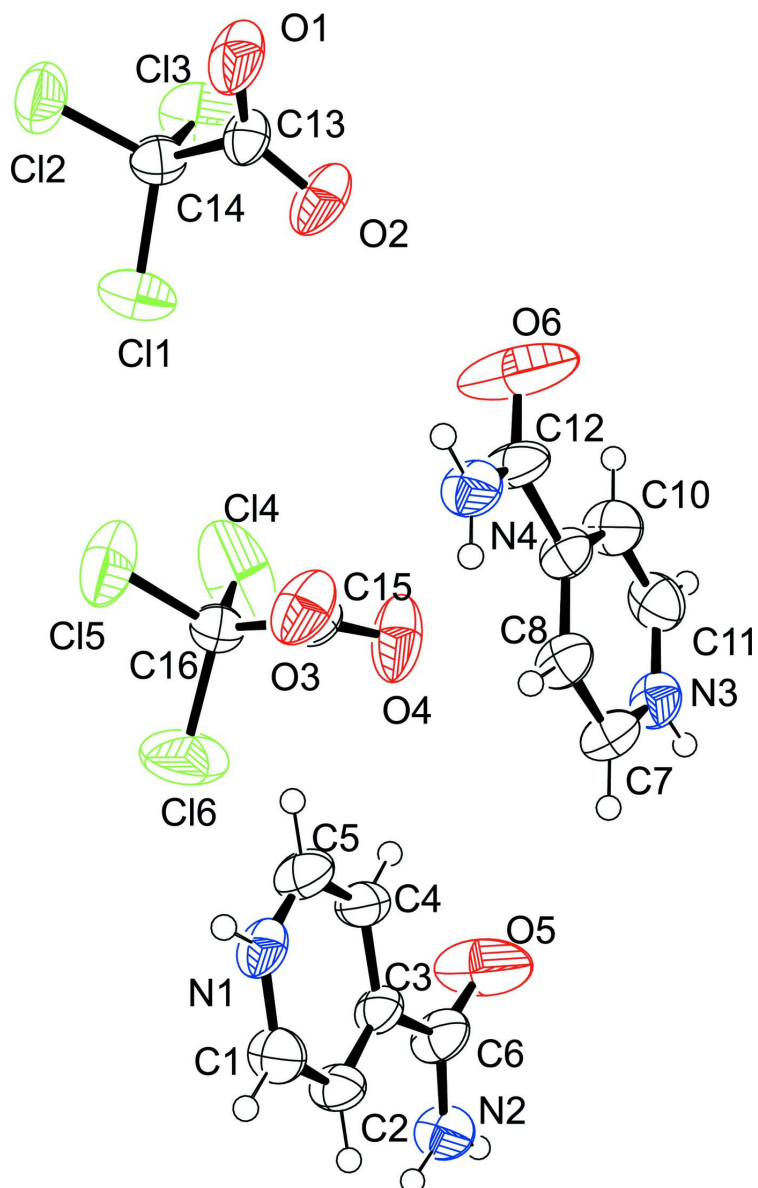
S2. Experimental

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

S3. Refinement

The presence of atoms H1A and H3B bonded to N1 and N3, respectively was confirmed by the observation of peaks in those locations in an electron-density map. All H atoms were then added at calculated positions and refined using a riding model, with C—H = 0.93 Å and N—H = 0.86 Å, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{N})$. To improve the refinement results, two reflection with too high value of $\delta(F^2)/\text{e.s.d.}$ and with $F_o^2 < F_c^2$ were deleted from the refinement.

Displacement ellipsoid of O5 and O6 are large compared to the other atoms, however the treatment of O5 and O6 as disordered over two positions did not improve the model.

**Figure 1**

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

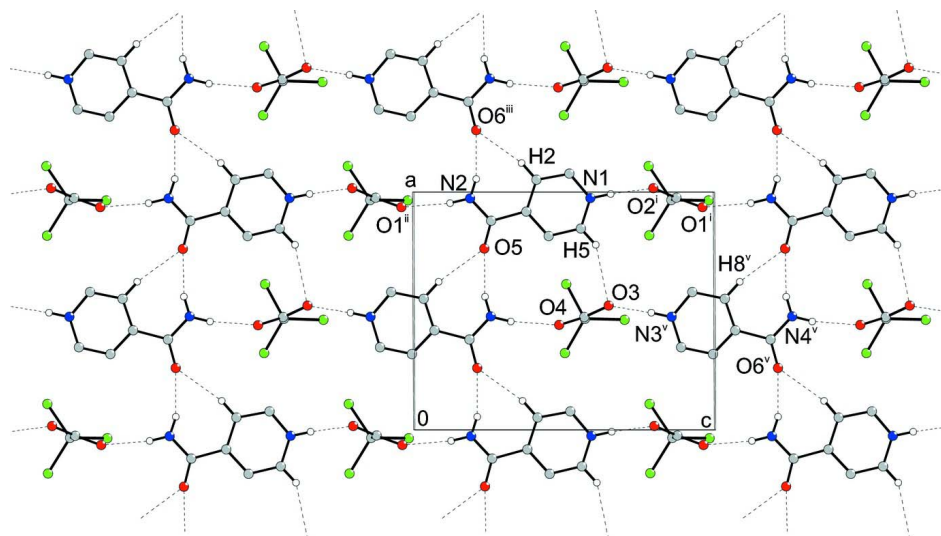


Figure 2

Layer formation. Dashed lines indicate intermolecular N—H \cdots O and C—H \cdots O hydrogen bonding. For the sake of clarity, H atoms not involved in the motif shown have been omitted. Symmetry codes: ⁱ $x + 1, y, z$; ⁱⁱ $x + 1, y, z - 1$; ⁱⁱⁱ $x + 1, -y + 1, z - 1/2$; ^v $x, -y + 1, z + 1/2$.

4-Carbamoylpyridin-1-ium 2,2,2-trichloroacetate

Crystal data

$C_6H_7N_2O^+ \cdot C_2Cl_3O_2^-$

$M_r = 285.51$

Monoclinic, Pc

Hall symbol: $P -2yc$

$a = 9.8768$ (3) Å

$b = 9.4403$ (3) Å

$c = 12.5157$ (3) Å

$\beta = 90.240$ (2)°

$V = 1166.95$ (6) Å³

$Z = 4$

$F(000) = 576$

$D_x = 1.625$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 4989 reflections

$\theta = 3.0\text{--}30.3^\circ$

$\mu = 0.78$ mm⁻¹

$T = 293$ K

Cube, colourless

$0.2 \times 0.2 \times 0.2$ mm

Data collection

Agilent SuperNova, Dual, Cu at zero, Atlas diffractometer

Radiation source: SuperNova (Mo) X-ray Source

Mirror monochromator

Detector resolution: 10.4933 pixels mm⁻¹

ω scans

Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2011)

$T_{\min} = 0.860, T_{\max} = 0.860$

11114 measured reflections

5195 independent reflections

4452 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.026$

$\theta_{\max} = 27.5^\circ, \theta_{\min} = 3.0^\circ$

$h = -12 \rightarrow 12$

$k = -12 \rightarrow 12$

$l = -16 \rightarrow 16$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.069$

$wR(F^2) = 0.199$

$S = 1.04$

5195 reflections

289 parameters

2 restraints

Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.104P)^2 + 1.1455P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.53 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.47 \text{ e } \text{\AA}^{-3}$$

Absolute structure: Flack (1983), 2512 Friedel pairs

Absolute structure parameter: 0.08 (12)

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.10761 (17)	0.0588 (2)	0.78518 (16)	0.0826 (5)
C12	-0.0364 (2)	0.03005 (15)	0.98391 (14)	0.0768 (5)
C13	-0.18351 (17)	0.0792 (2)	0.78632 (16)	0.0819 (5)
C14	0.3217 (3)	0.0487 (2)	0.5048 (3)	0.1287 (12)
C15	0.4613 (3)	0.04358 (18)	0.70263 (17)	0.0950 (7)
C16	0.6112 (3)	0.0464 (2)	0.5097 (3)	0.1206 (10)
N1	0.9738 (5)	0.3469 (4)	0.5899 (3)	0.0465 (9)
H1A	0.9911	0.3487	0.6573	0.07*
N2	0.9725 (5)	0.3126 (5)	0.1897 (3)	0.0545 (11)
H2A	0.9523	0.3112	0.1228	0.082*
H2B	1.0545	0.2975	0.21	0.082*
N3	0.4737 (5)	0.6594 (5)	0.3579 (3)	0.0489 (10)
H3A	0.4908	0.6608	0.2906	0.073*
N4	0.4753 (5)	0.6746 (5)	0.7577 (3)	0.0519 (10)
H4A	0.4549	0.6785	0.8243	0.078*
H4B	0.5587	0.6786	0.7384	0.078*
O1	-0.0643 (6)	0.3280 (4)	0.9614 (3)	0.0789 (15)
O2	0.0158 (5)	0.3511 (5)	0.7986 (3)	0.0713 (13)
O3	0.5208 (6)	0.3362 (5)	0.6487 (3)	0.0713 (12)
O4	0.4394 (7)	0.3324 (4)	0.4841 (3)	0.0834 (16)
O5	0.7614 (5)	0.3595 (9)	0.2357 (4)	0.116 (3)
O6	0.2588 (5)	0.6567 (10)	0.7106 (4)	0.138 (3)
C1	1.0769 (5)	0.3404 (6)	0.5206 (4)	0.0512 (12)
H1	1.1657	0.3372	0.5456	0.061*
C2	1.0507 (5)	0.3384 (6)	0.4112 (4)	0.0451 (10)
H2	1.1213	0.3352	0.3623	0.054*
C3	0.9178 (5)	0.3415 (5)	0.3768 (4)	0.0410 (9)
C4	0.8152 (5)	0.3479 (5)	0.4506 (4)	0.0471 (10)
H4	0.7253	0.3503	0.4283	0.057*

C5	0.8467 (6)	0.3506 (6)	0.5575 (4)	0.0555 (13)
H5	0.7775	0.3551	0.6076	0.067*
C6	0.8784 (6)	0.3376 (6)	0.2605 (4)	0.0538 (13)
C7	0.5779 (6)	0.6614 (6)	0.4269 (4)	0.0519 (12)
H7	0.6667	0.6613	0.4026	0.062*
C8	0.5502 (5)	0.6637 (5)	0.5346 (4)	0.0470 (11)
H8	0.6208	0.6664	0.5839	0.056*
C9	0.4176 (5)	0.6621 (5)	0.5696 (4)	0.0418 (10)
C10	0.3170 (6)	0.6561 (6)	0.4940 (4)	0.0543 (12)
H10	0.2271	0.6525	0.5155	0.065*
C11	0.3464 (6)	0.6552 (6)	0.3887 (4)	0.0556 (13)
H11	0.2771	0.6518	0.3382	0.067*
C12	0.3792 (5)	0.6626 (7)	0.6854 (4)	0.0558 (13)
C13	-0.0250 (5)	0.2831 (5)	0.8761 (3)	0.0436 (10)
C14	-0.0344 (5)	0.1177 (5)	0.8593 (4)	0.0481 (10)
C15	0.4722 (5)	0.2781 (5)	0.5682 (3)	0.0419 (9)
C16	0.4658 (5)	0.1124 (5)	0.5706 (4)	0.0463 (10)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0647 (9)	0.0950 (12)	0.0882 (12)	0.0167 (8)	0.0213 (8)	-0.0217 (10)
C12	0.1036 (12)	0.0537 (7)	0.0730 (10)	0.0026 (8)	0.0071 (9)	0.0188 (7)
C13	0.0599 (8)	0.1001 (12)	0.0856 (11)	-0.0164 (8)	-0.0161 (7)	-0.0240 (10)
C14	0.1135 (16)	0.0708 (11)	0.201 (3)	-0.0082 (11)	-0.0941 (19)	-0.0250 (14)
C15	0.1471 (19)	0.0620 (9)	0.0760 (11)	-0.0091 (10)	0.0003 (12)	0.0210 (8)
C16	0.1032 (15)	0.0709 (12)	0.188 (3)	0.0098 (11)	0.0773 (17)	-0.0256 (14)
N1	0.066 (3)	0.049 (2)	0.0250 (16)	-0.0007 (19)	0.0018 (17)	0.0029 (14)
N2	0.059 (2)	0.083 (3)	0.0216 (17)	0.008 (2)	-0.0014 (16)	-0.0018 (19)
N3	0.065 (3)	0.053 (2)	0.0280 (18)	0.004 (2)	-0.0005 (17)	0.0013 (16)
N4	0.057 (2)	0.069 (3)	0.0296 (19)	0.003 (2)	-0.0002 (17)	-0.0002 (18)
O1	0.146 (5)	0.059 (2)	0.0319 (18)	0.012 (3)	0.008 (2)	0.0011 (16)
O2	0.116 (4)	0.071 (3)	0.0270 (16)	-0.017 (2)	0.002 (2)	0.0035 (16)
O3	0.117 (4)	0.065 (2)	0.0324 (18)	-0.017 (2)	0.004 (2)	-0.0052 (16)
O4	0.158 (5)	0.052 (2)	0.040 (2)	0.007 (3)	-0.014 (3)	0.0037 (17)
O5	0.049 (2)	0.251 (8)	0.047 (2)	0.012 (4)	-0.0102 (19)	0.015 (4)
O6	0.051 (3)	0.312 (11)	0.051 (3)	-0.018 (4)	0.011 (2)	0.008 (4)
C1	0.047 (2)	0.067 (3)	0.039 (2)	0.005 (2)	-0.005 (2)	0.001 (2)
C2	0.040 (2)	0.065 (3)	0.030 (2)	0.003 (2)	0.0028 (17)	0.0013 (19)
C3	0.043 (2)	0.052 (3)	0.0288 (18)	-0.0013 (19)	0.0020 (16)	0.0073 (16)
C4	0.040 (2)	0.057 (3)	0.044 (2)	0.001 (2)	0.0088 (19)	0.003 (2)
C5	0.061 (3)	0.065 (3)	0.041 (3)	0.000 (2)	0.018 (2)	0.004 (2)
C6	0.051 (3)	0.082 (4)	0.029 (2)	-0.007 (3)	-0.0046 (19)	0.010 (2)
C7	0.053 (3)	0.063 (3)	0.040 (2)	-0.008 (2)	0.010 (2)	0.004 (2)
C8	0.049 (3)	0.063 (3)	0.029 (2)	-0.008 (2)	-0.0054 (19)	0.0021 (19)
C9	0.047 (2)	0.051 (3)	0.0277 (19)	-0.0028 (19)	-0.0021 (17)	0.0005 (17)
C10	0.049 (3)	0.071 (3)	0.043 (2)	0.002 (2)	-0.005 (2)	-0.001 (2)
C11	0.056 (3)	0.065 (3)	0.046 (3)	0.012 (3)	-0.010 (2)	-0.001 (2)

C12	0.044 (2)	0.093 (4)	0.030 (2)	-0.004 (3)	0.0067 (19)	0.003 (2)
C13	0.056 (2)	0.046 (2)	0.028 (2)	-0.001 (2)	-0.0072 (18)	0.0027 (17)
C14	0.047 (2)	0.051 (3)	0.046 (2)	-0.002 (2)	0.0039 (19)	-0.006 (2)
C15	0.058 (2)	0.044 (2)	0.0239 (18)	-0.004 (2)	0.0062 (17)	-0.0036 (17)
C16	0.044 (2)	0.048 (2)	0.047 (2)	0.001 (2)	0.0028 (19)	-0.005 (2)

Geometric parameters (Å, °)

C11—C14	1.774 (5)	O5—C6	1.213 (7)
C12—C14	1.766 (5)	O6—C12	1.233 (7)
C13—C14	1.768 (5)	C1—C2	1.393 (6)
C14—C16	1.748 (5)	C1—H1	0.93
C15—C16	1.777 (6)	C2—C3	1.380 (7)
C16—C16	1.743 (5)	C2—H2	0.93
N1—C5	1.318 (7)	C3—C4	1.375 (6)
N1—C1	1.342 (7)	C3—C6	1.505 (6)
N1—H1A	0.86	C4—C5	1.372 (8)
N2—C6	1.308 (7)	C4—H4	0.93
N2—H2A	0.86	C5—H5	0.93
N2—H2B	0.86	C7—C8	1.377 (7)
N3—C11	1.317 (7)	C7—H7	0.93
N3—C7	1.340 (7)	C8—C9	1.382 (7)
N3—H3A	0.86	C8—H8	0.93
N4—C12	1.314 (7)	C9—C10	1.371 (7)
N4—H4A	0.86	C9—C12	1.499 (6)
N4—H4B	0.86	C10—C11	1.350 (8)
O1—C13	1.213 (6)	C10—H10	0.93
O2—C13	1.233 (6)	C11—H11	0.93
O3—C15	1.242 (6)	C13—C14	1.578 (7)
O4—C15	1.215 (6)	C15—C16	1.566 (7)
C5—N1—C1	121.8 (4)	C7—C8—C9	120.2 (4)
C5—N1—H1A	119.1	C7—C8—H8	119.9
C1—N1—H1A	119.1	C9—C8—H8	119.9
C6—N2—H2A	120	C10—C9—C8	117.8 (4)
C6—N2—H2B	120	C10—C9—C12	118.8 (4)
H2A—N2—H2B	120	C8—C9—C12	123.4 (4)
C11—N3—C7	122.9 (4)	C11—C10—C9	121.1 (5)
C11—N3—H3A	118.6	C11—C10—H10	119.5
C7—N3—H3A	118.6	C9—C10—H10	119.5
C12—N4—H4A	120	N3—C11—C10	119.6 (5)
C12—N4—H4B	120	N3—C11—H11	120.2
H4A—N4—H4B	120	C10—C11—H11	120.2
N1—C1—C2	119.8 (5)	O6—C12—N4	121.5 (5)
N1—C1—H1	120.1	O6—C12—C9	119.7 (5)
C2—C1—H1	120.1	N4—C12—C9	118.8 (4)
C3—C2—C1	118.6 (4)	O1—C13—O2	128.1 (5)
C3—C2—H2	120.7	O1—C13—C14	116.4 (4)

C1—C2—H2	120.7	O2—C13—C14	115.4 (4)
C4—C3—C2	119.6 (4)	C13—C14—C12	110.3 (3)
C4—C3—C6	117.6 (4)	C13—C14—C13	108.7 (3)
C2—C3—C6	122.8 (4)	C12—C14—C13	110.3 (3)
C5—C4—C3	119.5 (5)	C13—C14—C11	109.5 (3)
C5—C4—H4	120.3	C12—C14—C11	109.1 (3)
C3—C4—H4	120.3	C13—C14—C11	108.9 (3)
N1—C5—C4	120.7 (5)	O4—C15—O3	128.1 (5)
N1—C5—H5	119.6	O4—C15—C16	115.3 (4)
C4—C5—H5	119.6	O3—C15—C16	116.2 (4)
O5—C6—N2	122.4 (5)	C15—C16—C16	108.4 (3)
O5—C6—C3	119.0 (5)	C15—C16—C14	111.6 (4)
N2—C6—C3	118.6 (5)	C16—C16—C14	110.0 (3)
N3—C7—C8	118.4 (5)	C15—C16—C15	112.6 (3)
N3—C7—H7	120.8	C16—C16—C15	107.4 (3)
C8—C7—H7	120.8	C14—C16—C15	106.8 (3)
C5—N1—C1—C2	0.5 (8)	C7—N3—C11—C10	1.5 (8)
N1—C1—C2—C3	-0.8 (8)	C9—C10—C11—N3	0.4 (9)
C1—C2—C3—C4	0.7 (8)	C10—C9—C12—O6	0.5 (10)
C1—C2—C3—C6	-179.1 (5)	C8—C9—C12—O6	-177.8 (7)
C2—C3—C4—C5	-0.3 (8)	C10—C9—C12—N4	-177.1 (6)
C6—C3—C4—C5	179.6 (5)	C8—C9—C12—N4	4.6 (8)
C1—N1—C5—C4	0.0 (8)	O1—C13—C14—C12	-23.3 (6)
C3—C4—C5—N1	-0.1 (8)	O2—C13—C14—C12	159.9 (4)
C4—C3—C6—O5	9.9 (9)	O1—C13—C14—C13	97.8 (5)
C2—C3—C6—O5	-170.3 (7)	O2—C13—C14—C13	-79.1 (5)
C4—C3—C6—N2	-171.0 (5)	O1—C13—C14—C11	-143.3 (5)
C2—C3—C6—N2	8.8 (9)	O2—C13—C14—C11	39.8 (6)
C11—N3—C7—C8	-2.1 (8)	O4—C15—C16—C16	80.2 (6)
N3—C7—C8—C9	0.8 (8)	O3—C15—C16—C16	-93.8 (5)
C7—C8—C9—C10	0.9 (8)	O4—C15—C16—C14	-41.1 (6)
C7—C8—C9—C12	179.2 (5)	O3—C15—C16—C14	145.0 (5)
C8—C9—C10—C11	-1.6 (9)	O4—C15—C16—C15	-161.2 (5)
C12—C9—C10—C11	-180.0 (5)	O3—C15—C16—C15	24.9 (6)

Hydrogen-bond geometry (\AA , $^\circ$)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots O2 ⁱ	0.86	1.78	2.643 (5)	176
N2—H2A \cdots O1 ⁱⁱ	0.86	2.03	2.883 (6)	170
N2—H2B \cdots O6 ⁱⁱⁱ	0.86	2.06	2.854 (7)	152
N3—H3A \cdots O3 ^{iv}	0.86	1.8	2.661 (5)	178
N4—H4A \cdots O4 ^v	0.86	2.01	2.858 (6)	169
N4—H4B \cdots O5 ^v	0.86	2.03	2.858 (7)	160
C2—H2 \cdots O6 ⁱⁱⁱ	0.93	2.34	3.251 (7)	166

C5—H5···O3	0.93	2.6	3.422 (8)	148
C8—H8···O5 ^v	0.93	2.36	3.270 (6)	166

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