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4-[(Adamantan-1-yl)carbamoyl]-pyridinium chloride

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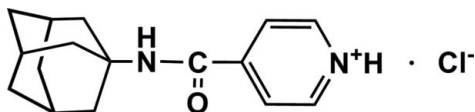
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 Key indicators: single-crystal X-ray study; $T = 93$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.044; wR factor = 0.157; data-to-parameter ratio = 18.1.

In the title salt, $\text{C}_{16}\text{H}_{21}\text{N}_2\text{O}^+\cdot\text{Cl}^-$, the amide group makes a dihedral angle of $24.98(2)^\circ$ with respect to the pyridinium ring. In the crystal, both the amide and pyridinium N atoms are involved in $\text{N}-\text{H}\cdots\text{Cl}$ hydrogen bonding. Weak intermolecular $\text{C}-\text{H}\cdots\text{Cl}$ and $\text{C}-\text{H}\cdots\text{O}$ interactions also occur.

Related literature

For the structures and properties of related compounds, see: Fu *et al.* (2011a,b,c); Dai & Chen (2011); Xu *et al.* (2011); Zheng (2011).



Experimental

Crystal data

 $\text{C}_{16}\text{H}_{21}\text{N}_2\text{O}^+\cdot\text{Cl}^-$
 $M_r = 292.80$

 Monoclinic, $P2_1/n$
 $a = 7.072(4)$ Å

 $b = 22.691(13)$ Å

 $c = 8.905(6)$ Å

 $\beta = 91.703(7)^\circ$
 $V = 1428.3(15)$ Å³
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 0.27$ mm⁻¹
 $T = 93$ K

 $0.10 \times 0.03 \times 0.03$ mm

Data collection

 Rigaku Mercury2 diffractometer
 Absorption correction: multi-scan
 (*CrystalClear*; Rigaku, 2005)
 $T_{\min} = 0.910$, $T_{\max} = 1.000$

 13912 measured reflections
 3278 independent reflections
 2863 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.157$
 $S = 1.19$

3278 reflections

181 parameters

H-atom parameters constrained

 $\Delta\rho_{\max} = 0.54$ e Å⁻³
 $\Delta\rho_{\min} = -0.43$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1A}\cdots\text{Cl1}^{\text{i}}$	0.86	2.19	3.020 (2)	161
$\text{N2}-\text{H2A}\cdots\text{Cl1}$	0.86	2.51	3.272 (2)	148
$\text{C1}-\text{H1B}\cdots\text{Cl1}^{\text{ii}}$	0.95	2.77	3.520 (3)	136
$\text{C2}-\text{H2B}\cdots\text{Cl1}^{\text{iii}}$	0.95	2.76	3.494 (3)	134
$\text{C5}-\text{H5A}\cdots\text{O1}^{\text{iv}}$	0.95	2.31	3.149 (3)	147

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

Data collection: *CrystalClear* (Rigaku, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5543).

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supplementary materials

Acta Cryst. (2012). E68, o1895 [doi:10.1107/S1600536812022866]

4-[(Adamantan-1-yl)carbamoyl]pyridinium chloride**Ying-Chun Wang****Comment**

Organic amino compounds attracted more attention as phase transition dielectric materials for its application in memory storage (Fu *et al.*, 2011*a, b* and *c*). With the purpose of obtaining phase transition crystals of amino compounds, various amines have been studied and we have elaborated a series of new materials with this organic molecules (Dai & Chen 2011; Xu, *et al.* 2011; Zheng 2011). In this study, we describe the crystal structure of the title compound, 4-[(adamantyl)carbamoyl]pyridinium chloride.

The asymmetric unit is composed of one 4-[(adamantyl)carbamoyl]pyridinium cation and one chloride anion. The pyridine N atom was protonated, and the amide group makes a dihedral angle of 24.98 (2)° with respect to the pyridinium ring (Fig.1). The torsion angles of C2—C3—C6—O1 and C2—C3—C6—N2 are 23.6 (3)° and -158.1 (4)°, C4—C3—C6—O1 and C4—C3—C6—N2 are -152.3 (2)° and 26.0 (3)°. Intermolecular N—H···Cl bonds and weak C—H···Cl and C—H···O contacts link ions into a network parallel to (0 1 0) plane (Table 1 and Fig 2).

Experimental

Isonicotinic acid 5 g was added in thionyl chloride (50 ml), and the mixture reacted at 353 K for 5 h. Then the solvate was removed under reduced pressure, the isonicotinoyl chloride was obtained. The 1-aminodiamantane hydrochloride (10 mmol) and triethylamine 2.02 g (20 mmol) dissolved in chloroform (40 ml) at 273 K, then the isonicotinoyl chloride 1.51 g (10 mmol) was added. Then the reactant mixture was stirred for 7 h at room temperature and some flaxen solid appeared. After filtering the mixture, the solid was dissolved in water and was neutralized with sodium carbonate, The mixed solution was extracted by dichloromethane. The N-(1-adamantyl)isonicotinamide was obtained when the dichloromethane was evaporated under reduced pressure. A mixture of N(1-adamantyl)isonicotinamide 2.56 g (10 mmol), HCl (2.0 mL, 6 mol/L), and 20 mL ethanol were added into a 50ml flask and refluxed for 5 hours, then cooled and filtrated. The solution was evaporated slowly in the air. Colorless block crystals suitable for X-ray analysis were obtained after one week.

Refinement

All H atoms attached to C atoms were fixed geometrically and treated as riding with C-H = 0.95 Å (aromatic), C-H = 0.99 Å (methylene) and C-H = 1.00 Å (methine) with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. H atoms bonded to N atoms were located in difference Fourier maps and restrained with the H—N = 0.86 (2)Å. In the last stage of refinement they were treated as riding on the N atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$.

Computing details

Data collection: *CrystalClear* (Rigaku, 2005); cell refinement: *CrystalClear* (Rigaku, 2005); data reduction: *CrystalClear* (Rigaku, 2005); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for

publication: *SHELXTL* (Sheldrick, 2008).

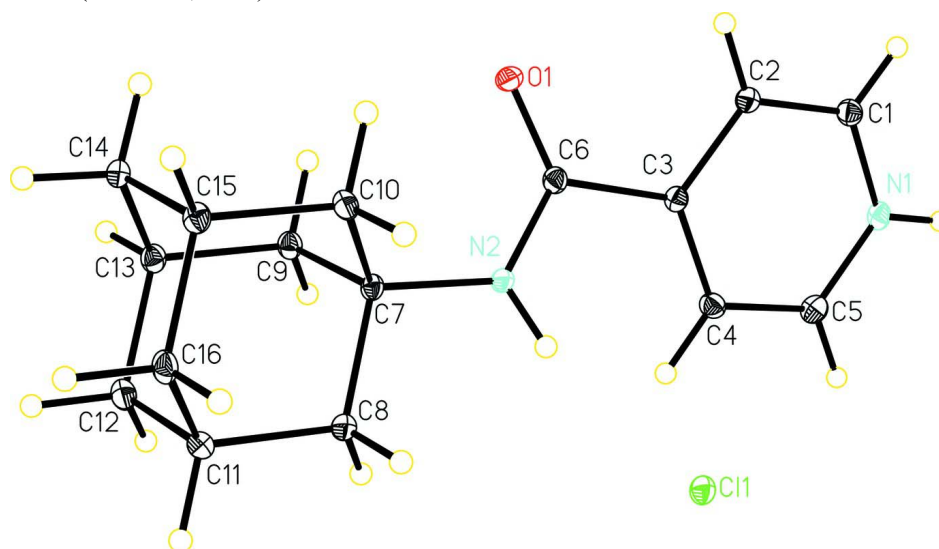


Figure 1

Molecular view of the title compound with the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

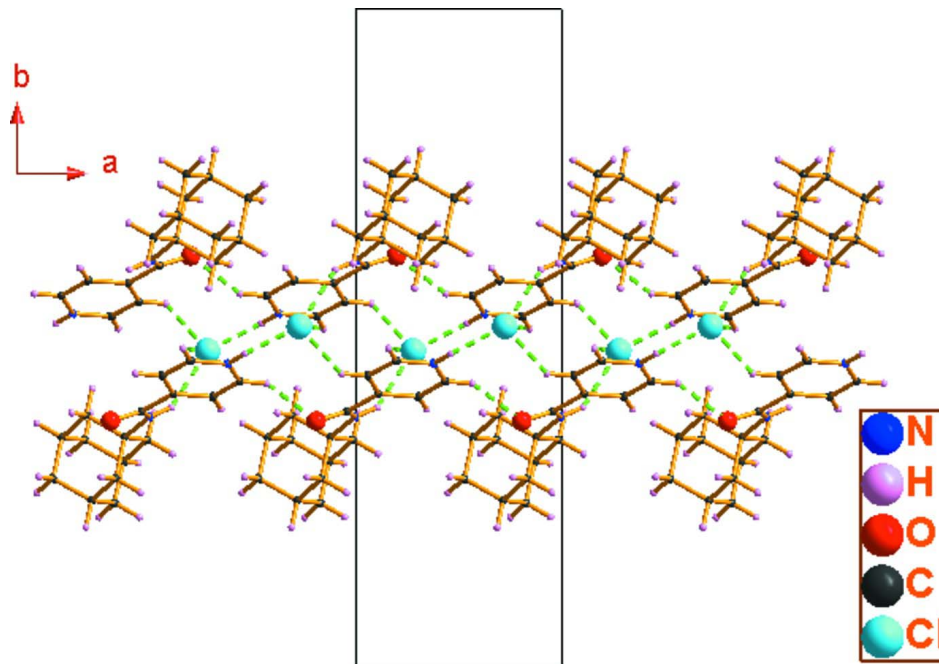


Figure 2

The crystal packing of the title compound viewed along the *c* axis showing the hydrogen bondings unit (dashed line).

4-[(Adamantan-1-yl)carbamoyl]pyridinium chloride

Crystal data

$C_{16}H_{21}N_2O^+ \cdot Cl^-$
 $M_r = 292.80$

Monoclinic, $P2_1/n$
 Hall symbol: -P 2yn

$a = 7.072$ (4) Å
 $b = 22.691$ (13) Å
 $c = 8.905$ (6) Å
 $\beta = 91.703$ (7)°
 $V = 1428.3$ (15) Å³
 $Z = 4$
 $F(000) = 624$
 $D_x = 1.362$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 3278 reflections
 $\theta = 1.8$ – 27.5 °
 $\mu = 0.27$ mm⁻¹
 $T = 93$ K
 Block, colourless
 $0.10 \times 0.03 \times 0.03$ mm

Data collection

Rigaku Mercury2
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 Detector resolution: 13.6612 pixels mm⁻¹
 CCD profile fitting scans
 Absorption correction: multi-scan
 (*CrystalClear*; Rigaku, 2005)
 $T_{\min} = 0.910$, $T_{\max} = 1.000$

13912 measured reflections
 3278 independent reflections
 2863 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$
 $\theta_{\max} = 27.5$ °, $\theta_{\min} = 1.8$ °
 $h = -9 \rightarrow 9$
 $k = -29 \rightarrow 29$
 $l = -11 \rightarrow 11$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.157$
 $S = 1.19$
 3278 reflections
 181 parameters
 0 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
 $w = 1/[\sigma^2(F_o^2) + (0.0868P)^2 + 0.3419P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.54$ e Å⁻³
 $\Delta\rho_{\min} = -0.43$ e Å⁻³

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N2	0.4961 (2)	0.11853 (7)	0.77561 (19)	0.0136 (4)
H2A	0.6014	0.1044	0.7457	0.016*
O1	0.3053 (2)	0.12500 (7)	0.97887 (17)	0.0195 (4)
N1	0.8767 (2)	0.03797 (7)	1.21603 (19)	0.0150 (4)
H1A	0.9604	0.0230	1.2768	0.018*
C4	0.8026 (3)	0.08813 (9)	0.9896 (2)	0.0157 (4)
H4A	0.8425	0.1069	0.9006	0.019*
C8	0.4853 (3)	0.15358 (9)	0.5188 (2)	0.0154 (4)
H8A	0.5154	0.1135	0.4830	0.018*

H8B	0.6058	0.1748	0.5389	0.018*
C12	0.3246 (3)	0.24905 (9)	0.4539 (2)	0.0169 (4)
H12A	0.4446	0.2706	0.4735	0.020*
H12B	0.2500	0.2709	0.3762	0.020*
C6	0.4548 (3)	0.11078 (9)	0.9207 (2)	0.0143 (4)
C7	0.3740 (3)	0.14958 (8)	0.6638 (2)	0.0123 (4)
C3	0.6104 (3)	0.08371 (8)	1.0195 (2)	0.0127 (4)
C2	0.5567 (3)	0.05623 (9)	1.1516 (2)	0.0153 (4)
H2B	0.4267	0.0531	1.1743	0.018*
C9	0.3315 (3)	0.21225 (8)	0.7186 (2)	0.0146 (4)
H9A	0.2611	0.2103	0.8129	0.018*
H9B	0.4515	0.2336	0.7395	0.018*
C13	0.2134 (3)	0.24535 (8)	0.5982 (2)	0.0151 (4)
H13A	0.1859	0.2861	0.6346	0.018*
C5	0.9336 (3)	0.06493 (9)	1.0909 (2)	0.0169 (4)
H5A	1.0648	0.0680	1.0722	0.020*
C10	0.1881 (3)	0.11626 (8)	0.6305 (2)	0.0143 (4)
H10A	0.1160	0.1126	0.7237	0.017*
H10B	0.2160	0.0761	0.5938	0.017*
C16	0.1811 (3)	0.15352 (9)	0.3664 (2)	0.0161 (4)
H16A	0.2081	0.1133	0.3296	0.019*
H16B	0.1052	0.1744	0.2877	0.019*
C15	0.0694 (3)	0.14989 (9)	0.5109 (2)	0.0146 (4)
H15A	-0.0520	0.1284	0.4902	0.017*
C1	0.6932 (3)	0.03367 (9)	1.2493 (2)	0.0165 (4)
H1B	0.6576	0.0151	1.3398	0.020*
C14	0.0263 (3)	0.21217 (9)	0.5695 (2)	0.0157 (4)
H14A	-0.0534	0.2337	0.4946	0.019*
H14B	-0.0438	0.2095	0.6639	0.019*
C11	0.3670 (3)	0.18655 (9)	0.3972 (2)	0.0163 (4)
H11A	0.4396	0.1890	0.3028	0.020*
Cl1	0.77720 (7)	0.01868 (2)	0.63897 (5)	0.01459 (17)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N2	0.0114 (8)	0.0184 (8)	0.0110 (8)	0.0036 (6)	0.0002 (7)	0.0022 (6)
O1	0.0137 (8)	0.0295 (8)	0.0154 (8)	0.0054 (6)	0.0025 (6)	0.0056 (6)
N1	0.0154 (9)	0.0175 (8)	0.0120 (8)	0.0031 (6)	-0.0045 (7)	0.0002 (6)
C4	0.0164 (11)	0.0172 (10)	0.0135 (10)	-0.0002 (7)	0.0005 (8)	0.0011 (7)
C8	0.0145 (10)	0.0183 (10)	0.0134 (10)	0.0036 (7)	0.0008 (8)	0.0020 (7)
C12	0.0178 (11)	0.0159 (10)	0.0167 (10)	0.0001 (7)	-0.0017 (8)	0.0032 (7)
C6	0.0126 (10)	0.0156 (10)	0.0146 (10)	0.0001 (7)	-0.0017 (8)	0.0015 (7)
C7	0.0109 (9)	0.0141 (9)	0.0117 (9)	0.0010 (7)	-0.0020 (7)	0.0007 (7)
C3	0.0134 (10)	0.0133 (9)	0.0114 (9)	0.0009 (7)	-0.0013 (8)	-0.0009 (7)
C2	0.0137 (10)	0.0173 (9)	0.0149 (10)	0.0018 (7)	0.0004 (8)	0.0019 (8)
C9	0.0161 (10)	0.0145 (10)	0.0130 (10)	0.0011 (7)	-0.0015 (8)	-0.0002 (7)
C13	0.0175 (10)	0.0119 (9)	0.0158 (10)	0.0018 (7)	0.0000 (8)	0.0001 (7)
C5	0.0136 (10)	0.0183 (10)	0.0186 (11)	-0.0007 (7)	-0.0024 (8)	-0.0018 (7)
C10	0.0150 (10)	0.0138 (9)	0.0141 (10)	-0.0011 (7)	-0.0015 (8)	0.0030 (7)

C16	0.0179 (10)	0.0177 (10)	0.0124 (10)	0.0029 (8)	-0.0036 (8)	0.0008 (7)
C15	0.0129 (10)	0.0163 (10)	0.0142 (10)	-0.0014 (7)	-0.0034 (8)	0.0000 (7)
C1	0.0196 (11)	0.0163 (10)	0.0138 (10)	0.0020 (8)	0.0014 (8)	0.0003 (8)
C14	0.0140 (10)	0.0183 (10)	0.0148 (10)	0.0040 (7)	0.0004 (8)	0.0038 (7)
C11	0.0163 (10)	0.0193 (10)	0.0133 (10)	0.0023 (8)	0.0014 (8)	0.0027 (7)
Cl1	0.0136 (3)	0.0161 (3)	0.0139 (3)	0.00244 (16)	-0.00125 (19)	0.00123 (16)

Geometric parameters (Å, °)

N2—C6	1.345 (3)	C2—C1	1.378 (3)
N2—C7	1.478 (2)	C2—H2B	0.9500
N2—H2A	0.8600	C9—C13	1.536 (3)
O1—C6	1.234 (3)	C9—H9A	0.9900
N1—C1	1.343 (3)	C9—H9B	0.9900
N1—C5	1.343 (3)	C13—C14	1.537 (3)
N1—H1A	0.8600	C13—H13A	1.0000
C4—C5	1.378 (3)	C5—H5A	0.9500
C4—C3	1.396 (3)	C10—C15	1.539 (3)
C4—H4A	0.9500	C10—H10A	0.9900
C8—C7	1.534 (3)	C10—H10B	0.9900
C8—C11	1.542 (3)	C16—C11	1.531 (3)
C8—H8A	0.9900	C16—C15	1.532 (3)
C8—H8B	0.9900	C16—H16A	0.9900
C12—C13	1.529 (3)	C16—H16B	0.9900
C12—C11	1.538 (3)	C15—C14	1.540 (3)
C12—H12A	0.9900	C15—H15A	1.0000
C12—H12B	0.9900	C1—H1B	0.9500
C6—C3	1.518 (3)	C14—H14A	0.9900
C7—C9	1.536 (3)	C14—H14B	0.9900
C7—C10	1.538 (3)	C11—H11A	1.0000
C3—C2	1.394 (3)		
C6—N2—C7	124.78 (17)	C12—C13—C9	109.36 (17)
C6—N2—H2A	117.6	C12—C13—C14	110.43 (17)
C7—N2—H2A	117.6	C9—C13—C14	108.90 (16)
C1—N1—C5	122.08 (18)	C12—C13—H13A	109.4
C1—N1—H1A	118.9	C9—C13—H13A	109.4
C5—N1—H1A	119.0	C14—C13—H13A	109.4
C5—C4—C3	119.15 (19)	N1—C5—C4	120.3 (2)
C5—C4—H4A	120.4	N1—C5—H5A	119.8
C3—C4—H4A	120.4	C4—C5—H5A	119.8
C7—C8—C11	109.77 (17)	C7—C10—C15	109.57 (16)
C7—C8—H8A	109.7	C7—C10—H10A	109.8
C11—C8—H8A	109.7	C15—C10—H10A	109.8
C7—C8—H8B	109.7	C7—C10—H10B	109.8
C11—C8—H8B	109.7	C15—C10—H10B	109.8
H8A—C8—H8B	108.2	H10A—C10—H10B	108.2
C13—C12—C11	109.58 (16)	C11—C16—C15	109.68 (16)
C13—C12—H12A	109.8	C11—C16—H16A	109.7
C11—C12—H12A	109.8	C15—C16—H16A	109.7

C13—C12—H12B	109.8	C11—C16—H16B	109.7
C11—C12—H12B	109.8	C15—C16—H16B	109.7
H12A—C12—H12B	108.2	H16A—C16—H16B	108.2
O1—C6—N2	125.65 (19)	C16—C15—C10	108.92 (17)
O1—C6—C3	118.52 (18)	C16—C15—C14	110.33 (16)
N2—C6—C3	115.80 (18)	C10—C15—C14	109.25 (16)
N2—C7—C8	106.92 (16)	C16—C15—H15A	109.4
N2—C7—C9	110.06 (16)	C10—C15—H15A	109.4
C8—C7—C9	108.83 (16)	C14—C15—H15A	109.4
N2—C7—C10	112.01 (16)	N1—C1—C2	119.81 (19)
C8—C7—C10	108.93 (17)	N1—C1—H1B	120.1
C9—C7—C10	110.00 (16)	C2—C1—H1B	120.1
C2—C3—C4	118.94 (18)	C13—C14—C15	109.11 (16)
C2—C3—C6	117.35 (18)	C13—C14—H14A	109.9
C4—C3—C6	123.59 (18)	C15—C14—H14A	109.9
C1—C2—C3	119.7 (2)	C13—C14—H14B	109.9
C1—C2—H2B	120.2	C15—C14—H14B	109.9
C3—C2—H2B	120.2	H14A—C14—H14B	108.3
C13—C9—C7	109.76 (16)	C16—C11—C12	109.60 (17)
C13—C9—H9A	109.7	C16—C11—C8	109.48 (17)
C7—C9—H9A	109.7	C12—C11—C8	108.91 (17)
C13—C9—H9B	109.7	C16—C11—H11A	109.6
C7—C9—H9B	109.7	C12—C11—H11A	109.6
H9A—C9—H9B	108.2	C8—C11—H11A	109.6
C7—N2—C6—O1	4.8 (3)	C7—C9—C13—C14	60.4 (2)
C7—N2—C6—C3	-173.33 (16)	C1—N1—C5—C4	-1.6 (3)
C6—N2—C7—C8	174.15 (18)	C3—C4—C5—N1	0.6 (3)
C6—N2—C7—C9	56.1 (2)	N2—C7—C10—C15	-178.66 (16)
C6—N2—C7—C10	-66.6 (2)	C8—C7—C10—C15	-60.6 (2)
C11—C8—C7—N2	-179.08 (15)	C9—C7—C10—C15	58.6 (2)
C11—C8—C7—C9	-60.2 (2)	C11—C16—C15—C10	-60.5 (2)
C11—C8—C7—C10	59.7 (2)	C11—C16—C15—C14	59.4 (2)
C5—C4—C3—C2	0.4 (3)	C7—C10—C15—C16	61.0 (2)
C5—C4—C3—C6	176.27 (18)	C7—C10—C15—C14	-59.6 (2)
O1—C6—C3—C2	23.6 (3)	C5—N1—C1—C2	1.5 (3)
N2—C6—C3—C2	-158.06 (18)	C3—C2—C1—N1	-0.4 (3)
O1—C6—C3—C4	-152.3 (2)	C12—C13—C14—C15	58.6 (2)
N2—C6—C3—C4	26.0 (3)	C9—C13—C14—C15	-61.5 (2)
C4—C3—C2—C1	-0.5 (3)	C16—C15—C14—C13	-58.5 (2)
C6—C3—C2—C1	-176.61 (18)	C10—C15—C14—C13	61.2 (2)
N2—C7—C9—C13	176.98 (16)	C15—C16—C11—C12	-59.6 (2)
C8—C7—C9—C13	60.1 (2)	C15—C16—C11—C8	59.8 (2)
C10—C7—C9—C13	-59.1 (2)	C13—C12—C11—C16	59.6 (2)
C11—C12—C13—C9	60.3 (2)	C13—C12—C11—C8	-60.1 (2)
C11—C12—C13—C14	-59.5 (2)	C7—C8—C11—C16	-59.5 (2)
C7—C9—C13—C12	-60.4 (2)	C7—C8—C11—C12	60.3 (2)

Hydrogen-bond geometry (Å, °)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots C11 ⁱ	0.86	2.19	3.020 (2)	161
N2—H2A \cdots C11	0.86	2.51	3.272 (2)	148
C1—H1B \cdots C11 ⁱⁱ	0.95	2.77	3.520 (3)	136
C2—H2B \cdots C11 ⁱⁱⁱ	0.95	2.76	3.494 (3)	134
C5—H5A \cdots O1 ^{iv}	0.95	2.31	3.149 (3)	147

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