



Crystal structure of 1-(1-chloroethyl)-6,7-dimethoxy-1,2,3,4-tetrahydroisoquinolinium chloride

Ziroat E. Urunbaeva,^a Kambarali K. Turgunov,^{b,c,*} Abdusalom Sh. Saidov^a and Valentina I. Vinogradova^b

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^aSamarkand State University named after Sh. Rashidov, University blv. 15, Samarkand 140104, Uzbekistan, ^bS. Yunusov Institute of the Chemistry of Plant Substances Academy of Sciences of Uzbekistan, Mirzo Ulugbek Str., 77, Tashkent 100170, Uzbekistan, and ^cTurin Polytechnic University in Tashkent 100095, 17 Little Ring Road, Tashkent, Uzbekistan.

*Correspondence e-mail: kk_turgunov@rambler.ru

1-(1-Chloroethyl)-6,7-dimethoxy-1,2,3,4-tetrahydroisoquinoline was synthesized through the reaction of homoveratrylamine with racemic lactic acid. The formation of two enantiomers, *RR* and *SS*, was detected by performing X-ray diffraction analysis on their chloride salts. The asymmetric unit of the crystal consists of a $C_{13}H_{19}ClNO_2^+$ molecular cation and a Cl^- anion. Two protonated enantiomers of the title compound, with *RR* and *SS* configurations of the chiral atoms, are connected into hydrogen-bonded dimers bridged by Cl^- anions. Weak $C-H \cdots Cl$ interactions lead to the formation of a chain running along the *a*-axis direction of the unit cell, which corresponds to the longest dimension (the preferential growth direction) of the needle-shaped monocystal. The crystal studied was refined as a two-component twin.

1. Chemical context

Isoquinoline alkaloids, widely distributed in the plant and animal kingdoms, have received much attention because of their important biological activities (Lundstrom, 1983). For example, 1,2,3,4-tetrahydroisoquinoline and 2-methyl-1,2,3,4-tetrahydroisoquinoline, present in mammalian brains, are known to induce Parkinson's disease (Ohta *et al.*, 1987; Niwa *et al.*, 1987). Effective synthetic methods for preparing of 1,2,3,4-tetrahydroisoquinoline derivatives have been found (Shinohara *et al.*, 1997). 1-Substituted-1,2,3,4-tetrahydroisoquinolines are especially intriguing among the synthetic derivatives of the isoquinoline alkaloid. They feature biologically active compounds, for example, an antiepileptic agent (Gitto *et al.*, 2003) and a derivative with inhibitory activity against bladder contraction (Naito *et al.*, 2005). A lot of work has been done on the synthesis and structural studies of 1-substituted-1,2,3,4-tetrahydroisoquinolines in search of active compounds (Olszak *et al.*, 1996; Pashev *et al.*, 2020; Turgunov *et al.*, 2016).

In this context, we treated homoveratrylamine with lactic acid and obtained the corresponding amide intermediate. Cyclization of the amide with $POCl_3$ and $NaBH_4$ afforded the

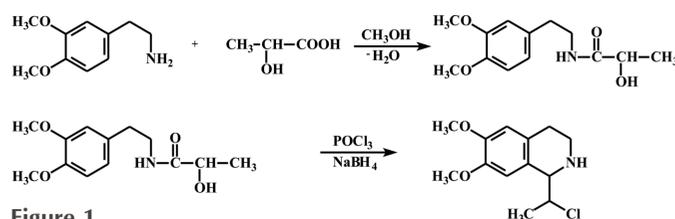
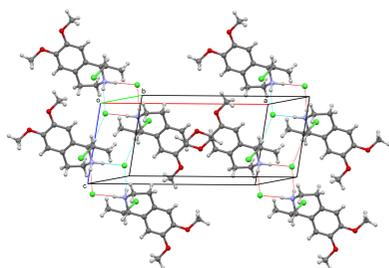
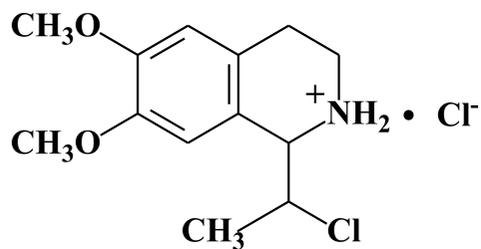


Figure 1
Synthesis scheme of the title compound.



title compound (Fig. 1). Racemic lactic acid was used in the synthesis, so four stereoisomers (*RR*; *RS*; *SS*; *SR*) of 1-(1-chloroethyl)-6,7-dimethoxy-1,2,3,4-tetrahydroisoquinoline were expected. Currently, we have detected the formation of two enantiomers, *RR* and *SS*, packed in a single crystal by X-ray diffraction (XRD) analysis. A detailed analysis of the reaction products is ongoing and will be published in our future work. To obtain good crystals suitable for XRD analysis, hydrochlorides of the isoquinolines were used.



2. Structural commentary

The title compound crystallizes in the monoclinic $P2_1/c$ (No. 14) space group. The asymmetric unit of the crystal contains one independent molecule with an *IS*, *IIS* configuration of chiral carbon atoms, so the crystal consists of *RR* and *SS* enantiomers. The C4A/C4–C8/C8A aromatic ring is twisted slightly with a slightly high value for the r.m.s. deviation (0.0245 Å) of the fitted atoms from the mean plane of the ring. The methoxy groups at C6 and C7 atoms are slightly rotated around the C6–O1 and C7–O2 bonds (Fig. 2), the dihedral angles between the plane of the aromatic ring and the planes defined by atoms C6/O1/C9 and C7/O2/C10 being 13.0 (3) and 6.5 (3)°, respectively. The C4A–C4 and C8A–C1 bonds are

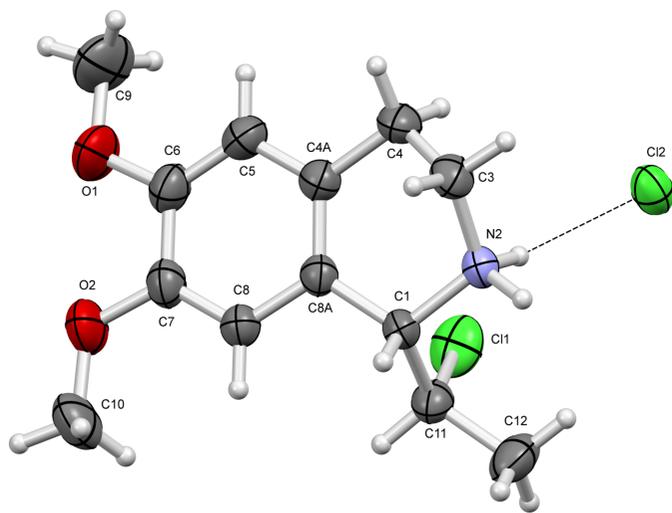


Figure 2
Displacement ellipsoid plot of the title compound with atom labels. Ellipsoids are drawn at the 50% probability level. The hydrogen bond formed between the molecular cation and the chloride anion is shown as a dashed line.

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C1–H1A \cdots Cl2 ⁱ	0.98	2.62	3.450 (2)	143
N2–H2A \cdots Cl2 ⁱⁱ	0.95 (2)	2.14 (2)	3.0895 (19)	177 (2)
N2–H2B \cdots Cl1	0.92 (2)	2.75 (3)	3.1828 (19)	110 (2)
N2–H2B \cdots Cl2	0.92 (2)	2.27 (2)	3.0751 (19)	146 (2)
C11–H11A \cdots Cl1 ⁱⁱⁱ	0.98	2.93	3.767 (2)	144
C12–H12A \cdots Cl2 ⁱⁱⁱ	0.96	2.75	3.710 (3)	174

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

slightly out of the plane, the deviations of C1 and C4 from the mean plane of aromatic ring being 0.206 (2) and -0.147 (2) Å, respectively. The heterocyclic ring of tetrahydroisoquinoline adopts a half chair conformation.

3. Supramolecular features

The presence of both enantiomers of the title compound in the crystal allows the molecules to link into inversion dimers through N2–H2A \cdots Cl2 and N2–H2B \cdots Cl2ⁱ [symmetry code: (i) $2 - x, 1 - y, 2 - z$] intermolecular interactions, forming rings with the graph-set motif $R_2^2(8)$ (Fig. 3, Table 1) where the Cl2 anions act as double hydrogen-bond acceptors. In addition, pairs of C1–H1A \cdots Cl2 weak interactions lead to chain formation along the *a*-axis direction, which is the longest cell dimension (preferential growth direction) of the monocystal. A C12–H12A \cdots Cl2 weak interaction leads to the formation of hydrogen-bonded layers parallel to the *bc* plane.

4. Database survey

A search in the Cambridge Structural Database (CSD, version 5.43, update of November 2022; Groom *et al.*, 2016) revealed 123 structures of 1-substituted and 2-substituted 6,7-dimethoxy-1,2,3,4-tetrahydroisoquinolines. Among these, 15 structures correspond to 1-substituted 6,7-dimethoxy-1,2,3,4-tetrahydroisoquinolines. Enantiopure crystal structures were determined for (*R*)-1-hydroxymethyl-6,7-dimethoxy-1,2,3,4-tetrahydroisoquinoline (refcode: BIMCEG) and (*S*)-1-hy-

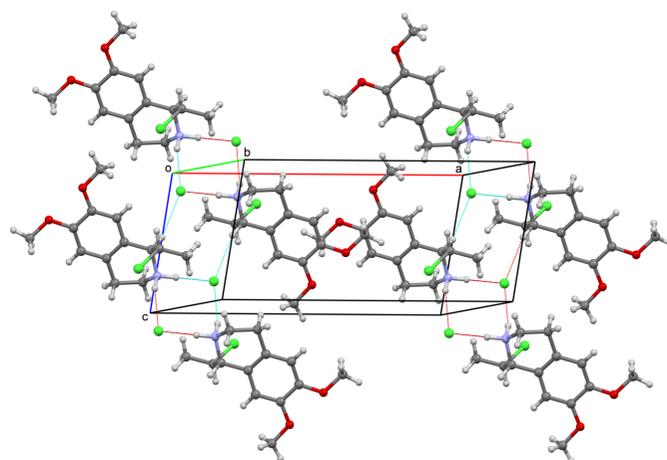


Figure 3
Hydrogen bonding in the crystal of the title compound.

droxymethyl-6,7-dimethoxy-1,2,3,4-tetrahydroisoquinoline chloride (refcode: BIMCIK), alkaloids isolated from seeds of *Calycotome Villosa* (Antri *et al.*, 2004). A search in the Cambridge Structural Database for the cationic form of 6,7-dimethoxy-1,2,3,4-tetrahydroisoquinoline resulted in 13 hits. Ten of these, where the molecule contains a chiral atom, are enantiopure crystals containing only proper symmetry elements. Therefore, in these crystal structures, the interlinking of molecules by hydrogen bonds differs from our case.

5. Synthesis and crystallization

N-(3,4-Dimethoxyphenylethyl)-2-hydroxypropanamide. A mixture of 1.81 g (0.01 mol) of homoveratrilamine and 0.9 g (0.01 mol) of lactic acid was dissolved in 5 ml of methanol. Self-heating occurred. Then the mixture was heated in an oil bath for 2 h at a temperature of 451–453 K. The progress of the reaction was monitored by TLC. The reaction mixture was dissolved in 100 mL of chloroform. The chloroform layer was first washed three times with 3% hydrochloric acid. The chloroform solution was then washed with water until neutral, followed by washing with 2% sodium hydroxide solution and water until neutral. The resulting chloroform solution was dried over Na₂SO₄ and then evaporated. The residue was crystallized from a mixture (acetone-hexane). White crystals with m.p. 343–344 K. Yield 70% (1.77 g). $R_f = 0.40$ chloroform-methanol (8:2).

¹H NMR: (400 MHz, CDCl₃, δ , ppm., J/Hz): 1.34 (3H, *d*, $J = 6.7$, H-3'), 2.73 (2H, *t*, $J = 7.1$, H- α), 3.46 (2H, *q*, $J = 6.7$, H- β), 3.81 (3H, *s*, OCH₃), 3.82 (3H, *s*, OCH₃), 4.10 (1H, *wide s*, OH), 4.17 (1H, *q*, $J = 6.8$ H -2'), 6.69 (2H, *top-top*, H-2,6), 6.77 (1H, *d*, $J = 8.6$, H-5), 6.90 (1H, *wide s*, NH).

¹³C NMR: 21.26 C-3, 35.19 C- α ; 40.60 C- β , 55.96 C-OCH₃, 55.96 C-OCH₃, 68.10 C-2I, 111.44 C-2, 111.97 C-5, 131 C-1, 120.73 C-6, 147.73 C-3, 149.01 C-4, 175.59 C-1-CO.

MS *m/z* (M^+) 253, 224, 165, 123.9 (124), 59.8 (60).

1-(1-Chloroethyl)-6,7-dimethoxy-1,2,3,4-tetrahydroisoquinoline. 1.550 mg (0.0061 mol) of *N*-(3,4-dimethoxyphenylethyl)-2-hydroxypropanamide were dissolved in 30 ml of absolute benzene, then 0.9404 mg (0.0061 mol) or 0.6–1 ml of POCl₃ were added. The reaction mixture was refluxed with a calcium chloride tube for 2 h. The progress of the reaction was monitored by TLC. After the reaction was complete after 2.5 h, benzene and residual POCl₃ were removed and the residue was dried. The residue was then dissolved in 50 mL of methanol. 0.6 g of NaBH₄ was added in small portions at 273–323 K for 3 h with constant stirring. This mixture was left overnight. The solvent was then removed and the residue was dissolved in distilled water. The aqueous layer was extracted several times with chloroform. The chloroform layer was combined and washed with water. After that, the chloroform layer was dried with Na₂SO₄. The residue was dissolved in methanol and precipitated as the hydrochloride using concentrated HCl solution. The precipitate was filtered, washed in acetone and dried. Yield 0.843 g (59%) (0.843 g), m.p. 476–477 K, $R_f = 0.57$ (chloroform–methanol 8:1.5).

Table 2

Experimental details.

Crystal data	
Chemical formula	C ₁₃ H ₁₉ ClNO ₂ ⁺ ·Cl ⁻
M_r	292.19
Crystal system, space group	Monoclinic, $P2_1/c$
Temperature (K)	293
a, b, c (Å)	16.1298 (3), 12.3736 (3), 7.46745 (16)
β (°)	100.190 (2)
V (Å ³)	1466.87 (6)
Z	4
Radiation type	Cu $K\alpha$
μ (mm ⁻¹)	3.94
Crystal size (mm)	0.25 × 0.10 × 0.05
Data collection	
Diffractometer	XtaLAB Synergy, Single source at home/near, HyPix3000
Absorption correction	Multi-scan (<i>CrysAlis PRO</i> ; Rigaku OD, 2018)
T_{\min}, T_{\max}	0.784, 1.000
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	12790, 4641, 4142
R_{int} ($\sin \theta/\lambda$) _{max} (Å ⁻¹)	0.061 0.602
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.050, 0.155, 1.09
No. of reflections	4641
No. of parameters	170
No. of restraints	2
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å ⁻³)	0.32, -0.26

Computer programs: *CrysAlis PRO* (Rigaku OD, 2018), *SHELXT2018/2* (Sheldrick, 2015a), *SHELXL2018/3* (Sheldrick, 2015b), *Mercury* (Macrae *et al.*, 2020), *SHELXL2014/7* (Sheldrick, 2015b).

¹H NMR (400 MHz, CDCl₃, δ , ppm, J / Hz): 1.87 (3H, *d*, $J = 7$, H-3'), 2.91 (2H, *m*, H-3a), 3.23 (2H, *m*, H-4), 3.84 (1H, *m*, H-3e), 3.85 (3H, *s*, OCH₃), 3.86 (3H, *s*, OCH₃), 4.59 (1H, *q*, $J = 3.5$, H-2'), 4.74 (1H, *d*, $J = 3.3$, H-1), 6.61 (1H, *s*, H-8), 6.69 (1H, *s*, H-5).

5.1. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The crystal under investigation exhibited twinning, which was identified during the initial analysis of the diffraction data. The twin law was determined based on the symmetry of the crystal and the diffraction analysis. In this case, a twofold rotation axis (along the *c* axis) related the two twin domains, with each domain contributing to the overall diffraction pattern. The twin fraction was estimated to be approximately 0.60 for component 1 and 0.40 for component 2, based on the refinement of the intensity data. Reflections in the HKLF 5 format were used for structure determination and refinement. The H atoms bonded to C atoms were placed geometrically (with C–H distances of 0.98 Å for CH, 0.97 Å for CH₂, 0.96 Å for CH₃ and 0.93 Å for C_{ar}) and included in the refinement in a riding-motion approximation with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ [$U_{\text{iso}} = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms]. The hydrogen atoms on the N1 were located in difference-Fourier maps and refined freely.

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References

- El Antri, A., Messouri, I., Bouktaib, M., El Alami, R., Bolte, M., El Bali, B. & Lachkar, M. (2004). *Molecules*, **9**, 650–657.
- Gitto, R., Barreca, M. L., De Luca, L., De Sarro, G., Ferreri, G., Quartarone, S., Russo, E., Constanti, A. & Chimirri, A. (2003). *J. Med. Chem.* **46**, 197–200.
- Groom, C. R., Bruno, I. J., Lightfoot, M. P. & Ward, S. C. (2016). *Acta Cryst.* **B72**, 171–179.
- Lundstom, J. (1983). *The Alkaloids*, Vol. 21 edited by A. Brossi, pp. 255–327. New York: Academic Press.
- Macrae, C. F., Sovago, I., Cottrell, S. J., Galek, P. T. A., McCabe, P., Pidcock, E., Platings, M., Shields, G. P., Stevens, J. S., Towler, M. & Wood, P. A. (2020). *J. Appl. Cryst.* **53**, 226–235.
- Naito, R., Yonetoku, Y., Okamoto, Y., Toyoshima, A., Ikeda, K. & Takeuchi, M. (2005). *J. Med. Chem.* **48**, 6597–6606.
- Niwa, T., Takeda, N., Kaneda, N., Hashizume, Y. & Nagatsu, T. (1987). *Biochem. Biophys. Res. Commun.* **144**, 1084–1089.
- Ohta, S., Kohno, M., Makino, Y., Tachikawa, O. & Hirobe, M. (1987). *Biomed. Res.* **8**, 453–456.
- Olszak, T. A., Stepień, A., Grabowski, M. J. & Brzezińska, E. (1996). *Acta Cryst.* **C52**, 1038–1040.
- Pashev, A., Burdzhiev, N. & Stanoeva, E. (2020). *Beilstein J. Org. Chem.* **16**, 1456–1464.
- Rigaku OD (2018). *CrysAlis PRO*. Rigaku OD, Yarnton, England.
- Sheldrick, G. M. (2015a). *Acta Cryst.* **A71**, 3–8.
- Sheldrick, G. M. (2015b). *Acta Cryst.* **C71**, 3–8.
- Shinohara, T., Toda, J. & Sano, T. (1997). *Chem. Pharm. Bull.* **45**, 813–819.
- Turgunov, K. K., Zhurakulov, Sh. N., Englert, U., Vinogradova, V. I. & Tashkhodjaev, B. (2016). *Acta Cryst.* **C72**, 607–611.

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Crystal structure of 1-(1-chloroethyl)-6,7-dimethoxy-1,2,3,4-tetrahydroisoquinolinium chloride

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Computing details

1-(1-Chloroethyl)-6,7-dimethoxy-1,2,3,4-tetrahydroisoquinolinium chloride

Crystal data

$C_{13}H_{19}ClNO_2^+ \cdot Cl^-$

$M_r = 292.19$

Monoclinic, $P2_1/c$

$a = 16.1298$ (3) Å

$b = 12.3736$ (3) Å

$c = 7.46745$ (16) Å

$\beta = 100.190$ (2)°

$V = 1466.87$ (6) Å³

$Z = 4$

$F(000) = 616$

$D_x = 1.323$ Mg m⁻³

Melting point: 476(2) K

Cu $K\alpha$ radiation, $\lambda = 1.54184$ Å

Cell parameters from 5593 reflections

$\theta = 2.8$ – 68.0 °

$\mu = 3.94$ mm⁻¹

$T = 293$ K

Prism, colourless

$0.25 \times 0.10 \times 0.05$ mm

Data collection

XtaLAB Synergy, Single source at home/near,

HyPix3000

diffractometer

Radiation source: micro-focus sealed X-ray

tube, PhotonJet (Cu) X-ray Source

Mirror monochromator

Detector resolution: 10.0000 pixels mm⁻¹

w σ cans

Absorption correction: multi-scan

(CrysAlisPro; Rigaku OD, 2018)

$T_{\min} = 0.784$, $T_{\max} = 1.000$

12790 measured reflections

4641 independent reflections

4142 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.061$

$\theta_{\max} = 68.2$ °, $\theta_{\min} = 7.0$ °

$h = -17 \rightarrow 19$

$k = -14 \rightarrow 14$

$l = -8 \rightarrow 7$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.050$

$wR(F^2) = 0.155$

$S = 1.09$

4641 reflections

170 parameters

2 restraints

Primary atom site location: iterative

Secondary atom site location: difference Fourier map

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.1122P)^2 + 0.0537P]$

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

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

$\Delta\rho_{\max} = 0.32$ e Å⁻³

$\Delta\rho_{\min} = -0.26$ 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. Refined as a 2-component twin

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	0.77537 (4)	0.66403 (6)	0.75105 (11)	0.0673 (3)
Cl2	0.91248 (3)	0.52447 (5)	1.18130 (7)	0.0506 (2)
O1	0.51533 (12)	0.3579 (2)	0.3763 (3)	0.0747 (6)
O2	0.58615 (12)	0.47214 (18)	0.1566 (3)	0.0644 (6)
C1	0.85038 (13)	0.49829 (17)	0.5981 (3)	0.0358 (5)
H1A	0.884823	0.477867	0.507792	0.043*
N2	0.89415 (11)	0.45706 (15)	0.7799 (2)	0.0374 (4)
H2A	0.9536 (13)	0.465 (2)	0.791 (4)	0.045*
H2B	0.8830 (17)	0.498 (2)	0.876 (3)	0.045*
C3	0.87520 (15)	0.34188 (19)	0.8137 (3)	0.0465 (5)
H3A	0.908683	0.318072	0.927730	0.056*
H3B	0.889133	0.296935	0.716790	0.056*
C4	0.78257 (15)	0.3314 (2)	0.8215 (3)	0.0496 (6)
H4A	0.770169	0.370253	0.926464	0.060*
H4B	0.768603	0.255903	0.834578	0.060*
C4A	0.72980 (14)	0.37668 (19)	0.6497 (3)	0.0416 (5)
C5	0.64592 (15)	0.3445 (2)	0.5957 (3)	0.0503 (6)
H5A	0.621781	0.297843	0.669519	0.060*
C6	0.59837 (15)	0.3808 (2)	0.4351 (4)	0.0512 (6)
C7	0.63650 (15)	0.4458 (2)	0.3182 (3)	0.0473 (6)
C8	0.71844 (14)	0.47887 (19)	0.3714 (3)	0.0414 (5)
H8A	0.743570	0.522290	0.294466	0.050*
C8A	0.76461 (13)	0.44768 (17)	0.5409 (3)	0.0369 (5)
C9	0.4704 (2)	0.3114 (4)	0.5026 (6)	0.1007 (14)
H9A	0.413028	0.299237	0.445017	0.151*
H9B	0.495855	0.243834	0.544775	0.151*
H9C	0.471544	0.359488	0.603867	0.151*
C10	0.6234 (2)	0.5306 (3)	0.0284 (4)	0.0711 (9)
H10A	0.581969	0.544019	-0.078361	0.107*
H10B	0.644673	0.598139	0.080729	0.107*
H10C	0.668884	0.489221	-0.004100	0.107*
C11	0.84659 (14)	0.62126 (19)	0.6022 (3)	0.0455 (5)
H11A	0.822722	0.645643	0.478997	0.055*
C12	0.93059 (18)	0.6771 (2)	0.6574 (4)	0.0650 (7)
H12A	0.922258	0.753986	0.656226	0.097*
H12B	0.955872	0.654494	0.777642	0.097*
H12C	0.966957	0.658458	0.573374	0.097*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0646 (4)	0.0535 (4)	0.0826 (5)	0.0072 (3)	0.0094 (4)	-0.0177 (3)
C12	0.0439 (3)	0.0715 (5)	0.0365 (3)	0.0006 (2)	0.0075 (2)	-0.0008 (2)
O1	0.0424 (9)	0.0996 (16)	0.0760 (12)	-0.0225 (10)	-0.0063 (9)	0.0100 (12)
O2	0.0483 (10)	0.0862 (15)	0.0512 (11)	-0.0048 (9)	-0.0120 (9)	0.0104 (9)
C1	0.0369 (10)	0.0382 (11)	0.0306 (10)	-0.0027 (8)	0.0019 (8)	0.0023 (8)
N2	0.0359 (9)	0.0418 (10)	0.0326 (9)	-0.0012 (7)	0.0008 (7)	0.0016 (7)
C3	0.0476 (12)	0.0411 (13)	0.0471 (12)	0.0041 (9)	-0.0015 (10)	0.0066 (9)
C4	0.0517 (13)	0.0465 (14)	0.0478 (12)	-0.0070 (10)	0.0009 (10)	0.0129 (10)
C4A	0.0426 (11)	0.0403 (12)	0.0405 (11)	-0.0027 (9)	0.0040 (9)	0.0012 (9)
C5	0.0455 (12)	0.0496 (14)	0.0552 (13)	-0.0112 (10)	0.0073 (10)	0.0035 (11)
C6	0.0390 (11)	0.0559 (15)	0.0555 (13)	-0.0083 (10)	0.0000 (10)	-0.0027 (11)
C7	0.0428 (11)	0.0532 (14)	0.0418 (12)	-0.0003 (10)	-0.0036 (9)	-0.0009 (10)
C8	0.0396 (11)	0.0455 (13)	0.0377 (11)	-0.0032 (9)	0.0032 (9)	0.0009 (9)
C8A	0.0371 (10)	0.0367 (11)	0.0353 (10)	-0.0010 (8)	0.0022 (8)	-0.0024 (8)
C9	0.0469 (16)	0.137 (4)	0.114 (3)	-0.030 (2)	0.0021 (18)	0.028 (3)
C10	0.0628 (17)	0.096 (2)	0.0488 (15)	0.0066 (16)	-0.0060 (13)	0.0178 (14)
C11	0.0465 (11)	0.0380 (12)	0.0482 (12)	-0.0029 (9)	-0.0019 (9)	0.0039 (9)
C12	0.0558 (15)	0.0462 (15)	0.0872 (19)	-0.0150 (12)	-0.0029 (14)	0.0039 (13)

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

C11—C11	1.813 (3)	C4A—C5	1.399 (3)
O1—C6	1.364 (3)	C5—C6	1.379 (4)
O1—C9	1.410 (4)	C5—H5A	0.9300
O2—C7	1.369 (3)	C6—C7	1.407 (4)
O2—C10	1.416 (4)	C7—C8	1.374 (3)
C1—N2	1.504 (3)	C8—C8A	1.405 (3)
C1—C8A	1.510 (3)	C8—H8A	0.9300
C1—C11	1.523 (3)	C9—H9A	0.9600
C1—H1A	0.9800	C9—H9B	0.9600
N2—C3	1.488 (3)	C9—H9C	0.9600
N2—H2A	0.95 (2)	C10—H10A	0.9600
N2—H2B	0.92 (2)	C10—H10B	0.9600
C3—C4	1.511 (3)	C10—H10C	0.9600
C3—H3A	0.9700	C11—C12	1.512 (3)
C3—H3B	0.9700	C11—H11A	0.9800
C4—C4A	1.516 (3)	C12—H12A	0.9600
C4—H4A	0.9700	C12—H12B	0.9600
C4—H4B	0.9700	C12—H12C	0.9600
C4A—C8A	1.382 (3)		
C6—O1—C9	117.5 (2)	C5—C6—C7	119.3 (2)
C7—O2—C10	117.4 (2)	O2—C7—C8	125.1 (2)
N2—C1—C8A	112.04 (17)	O2—C7—C6	115.3 (2)
N2—C1—C11	109.57 (17)	C8—C7—C6	119.7 (2)

C8A—C1—C11	112.46 (18)	C7—C8—C8A	120.6 (2)
N2—C1—H1A	107.5	C7—C8—H8A	119.7
C8A—C1—H1A	107.5	C8A—C8—H8A	119.7
C11—C1—H1A	107.5	C4A—C8A—C8	119.9 (2)
C3—N2—C1	113.60 (16)	C4A—C8A—C1	123.00 (18)
C3—N2—H2A	108.8 (15)	C8—C8A—C1	117.02 (19)
C1—N2—H2A	110.2 (16)	O1—C9—H9A	109.5
C3—N2—H2B	108.3 (17)	O1—C9—H9B	109.5
C1—N2—H2B	113.2 (17)	H9A—C9—H9B	109.5
H2A—N2—H2B	102 (2)	O1—C9—H9C	109.5
N2—C3—C4	108.83 (18)	H9A—C9—H9C	109.5
N2—C3—H3A	109.9	H9B—C9—H9C	109.5
C4—C3—H3A	109.9	O2—C10—H10A	109.5
N2—C3—H3B	109.9	O2—C10—H10B	109.5
C4—C3—H3B	109.9	H10A—C10—H10B	109.5
H3A—C3—H3B	108.3	O2—C10—H10C	109.5
C3—C4—C4A	110.28 (19)	H10A—C10—H10C	109.5
C3—C4—H4A	109.6	H10B—C10—H10C	109.5
C4A—C4—H4A	109.6	C12—C11—C1	115.2 (2)
C3—C4—H4B	109.6	C12—C11—C11	109.50 (19)
C4A—C4—H4B	109.6	C1—C11—C11	109.62 (16)
H4A—C4—H4B	108.1	C12—C11—H11A	107.4
C8A—C4A—C5	119.0 (2)	C1—C11—H11A	107.4
C8A—C4A—C4	120.46 (19)	C11—C11—H11A	107.4
C5—C4A—C4	120.6 (2)	C11—C12—H12A	109.5
C6—C5—C4A	121.2 (2)	C11—C12—H12B	109.5
C6—C5—H5A	119.4	H12A—C12—H12B	109.5
C4A—C5—H5A	119.4	C11—C12—H12C	109.5
O1—C6—C5	125.2 (2)	H12A—C12—H12C	109.5
O1—C6—C7	115.6 (2)	H12B—C12—H12C	109.5
C8A—C1—N2—C3	34.0 (2)	C5—C6—C7—C8	-4.8 (4)
C11—C1—N2—C3	159.49 (19)	O2—C7—C8—C8A	179.7 (2)
C1—N2—C3—C4	-64.4 (2)	C6—C7—C8—C8A	0.5 (4)
N2—C3—C4—C4A	55.1 (3)	C5—C4A—C8A—C8	-5.6 (3)
C3—C4—C4A—C8A	-19.8 (3)	C4—C4A—C8A—C8	172.3 (2)
C3—C4—C4A—C5	158.0 (2)	C5—C4A—C8A—C1	171.9 (2)
C8A—C4A—C5—C6	1.2 (4)	C4—C4A—C8A—C1	-10.1 (3)
C4—C4A—C5—C6	-176.7 (2)	C7—C8—C8A—C4A	4.8 (4)
C9—O1—C6—C5	12.7 (5)	C7—C8—C8A—C1	-172.9 (2)
C9—O1—C6—C7	-167.9 (3)	N2—C1—C8A—C4A	3.4 (3)
C4A—C5—C6—O1	-176.7 (3)	C11—C1—C8A—C4A	-120.5 (2)
C4A—C5—C6—C7	4.0 (4)	N2—C1—C8A—C8	-178.93 (18)
C10—O2—C7—C8	5.4 (4)	C11—C1—C8A—C8	57.1 (3)
C10—O2—C7—C6	-175.3 (3)	N2—C1—C11—C12	55.1 (3)
O1—C6—C7—O2	-3.5 (4)	C8A—C1—C11—C12	-179.6 (2)
C5—C6—C7—O2	175.9 (2)	N2—C1—C11—C11	-68.91 (19)
O1—C6—C7—C8	175.8 (2)	C8A—C1—C11—C11	56.4 (2)

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>
C1—H1A \cdots C12 ⁱ	0.98	2.62	3.450 (2)	143
N2—H2A \cdots C12 ⁱⁱ	0.95 (2)	2.14 (2)	3.0895 (19)	177 (2)
N2—H2B \cdots C11	0.92 (2)	2.75 (3)	3.1828 (19)	110 (2)
N2—H2B \cdots C12	0.92 (2)	2.27 (2)	3.0751 (19)	146 (2)
C11—H11A \cdots C11 ⁱⁱⁱ	0.98	2.93	3.767 (2)	144
C12—H12A \cdots C12 ⁱⁱⁱ	0.96	2.75	3.710 (3)	174

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