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Crystal structure of (*3E,5E*)-3,5-bis[4-(diethylazaniumyl)benzylidene]-1-methyl-4-oxopiperidin-1-ium trichloride dihydrate: a potential biophotonic material

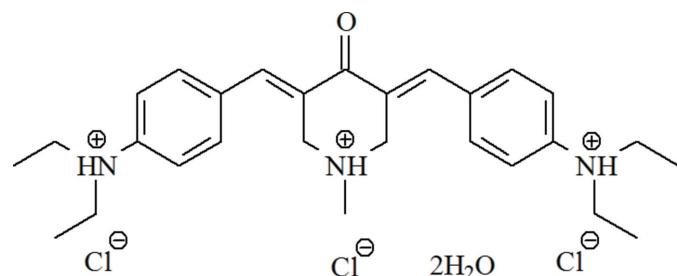
Volodymyr V. Nesterov,^a Lev N. Zakharov,^b Vladimir N. Nesterov^{a*} and Vladimir Shulaev^{c*}

^aDepartment of Chemistry, University of North Texas, Denton, TX 76203, USA, ^bCAMCOR Center for Advanced Materials Characterization in Oregon, University of Oregon, Eugene, Oregon 97403-1443, USA, and ^cDepartment of Biological Sciences, University of North Texas, Denton, TX 76203, USA. *Correspondence e-mail: vladimir.nesterov@unt.edu, shulaev@unt.edu

In the trication of the title hydrated molecular salt, $C_{28}H_{40}N_3O^{3+}\cdot 3Cl^- \cdot 2H_2O$, the central heterocyclic ring adopts a sofa conformation, with the exocyclic N—C bond in an equatorial orientation. The dihedral angles between the planar part of this heterocyclic ring and the two almost flat side-chain fragments, which include the aromatic ring and bridging atoms, are 28.8 (1) and 41.1 (1) $^\circ$. Both diethylazaniumyl substituents have a tetrahedral geometry, while the dihedral angles between the above-mentioned flat part of the aryl fragments and the imaginary planes drawn through atoms C—N—C of the diethylazaniumyl substituents are 86.3 (2) and 80.4 (1) $^\circ$, respectively. In the crystal, N—H···Cl hydrogen bonds link the cations and anions into [100] chains. The chains are cross-linked by numerous C—H···O and C—H···Cl interactions, generating a three-dimensional network. One of the chloride ions is disordered over two adjacent positions in a 0.895 (4):0.105 (4) ratio.

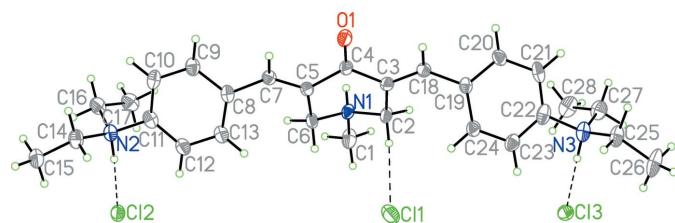
1. Chemical context

In a continuation of our work on the synthesis and structural investigations of non-linear optical organic compounds with two-photon absorption properties and potential biophotonic materials (Nesterov *et al.*, 2003, 2007; Nesterov *et al.*, 2011*a,b*; Sarkisov *et al.*, 2005), we determined the crystal structure of the title compound. This compound belongs to a group that has shown anticancer activity (Jia *et al.*, 1988; Dimmock *et al.*, 2001). It may also find application as an agent for locating cancer cells with two-photon excited fluorescence and as a potential agent for a photodynamic treatment of cancer (Nesterov *et al.*, 2003; Sarkisov *et al.*, 2005).



2. Structural commentary

The structure of the trication with chloride anions is illustrated in Fig. 1. There are also two water molecules of crystallization.

**Figure 1**

Perspective view of the trication and anions of (I), with hydrogen bonds shown as dashed lines. Displacement ellipsoids are drawn at the 30% probability level.

The central heterocycle adopts a sofa conformation: atom N1 lies $-0.732(3)$ Å out of the central C₅ plane [planar within $0.027(2)$ Å]. The dihedral angles between the flat part of the heterocycle (atoms C₂, C₃, C₄, C₅, and C₆) and the two almost planar fragments that include the phenyl-ring and the bridging atoms are $28.7(1)$ and $41.1(1)^\circ$ for (C₇–C₁₃) and (C₁₈–C₂₄), respectively. Such non-planarity might partly be caused by the presence of short intramolecular contacts H_{2AB}···H_{24A} and H_{6AB}···H_{13A} with distances 2.18 and 2.14 Å, respectively, which are shorter than the doubled van der Waals radius of the H atom (Rowland & Taylor, 1996). The mutual orientations of both aryl substituents relative to the flat part of the diethylazaniumyl groups (N₂, C₁₄, C₁₆ and N₃, C₂₅, C₂₇) are almost orthogonal [dihedral angles of $86.3(2)$ and $80.4(1)^\circ$, respectively]. This is in contrast to the starting material where such angles are close to zero and the substituents participate in conjugated systems with the respective aromatic rings (Nesterov *et al.*, 2003).

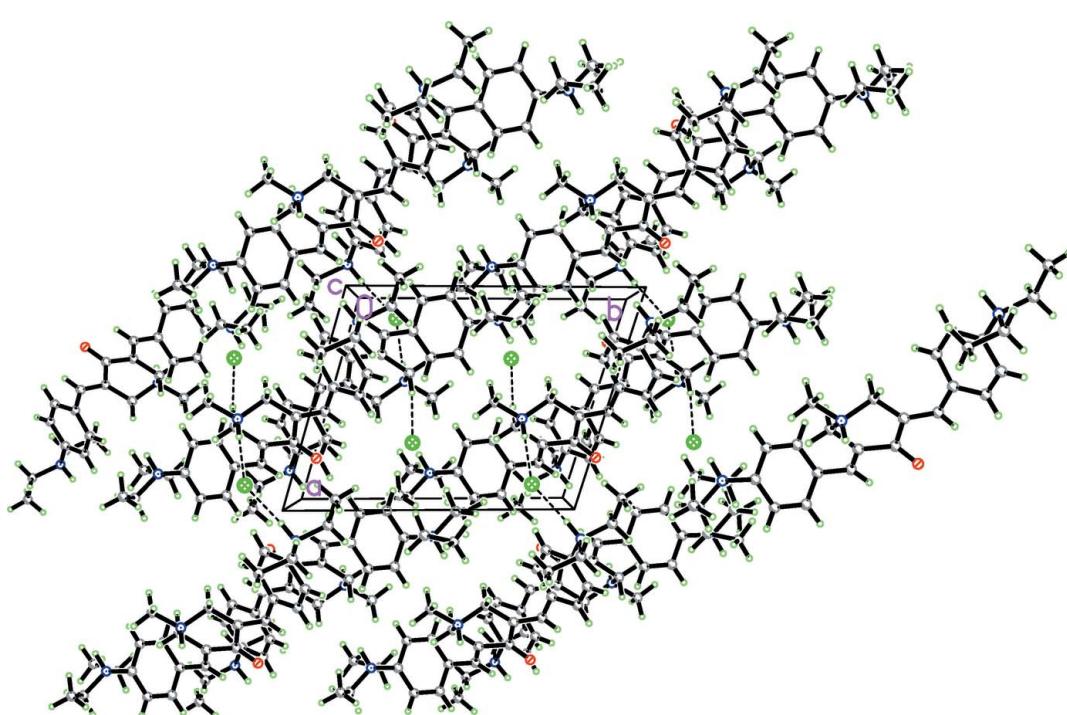
Table 1
Hydrogen-bond geometry (Å, °).

D–H···A	D–H	H···A	D···A	D–H···A
N2–H _{2C} ···Cl ₂	0.91(1)	2.27(1)	3.166(2)	169(3)
N3–H _{3A} ···Cl ₃	0.91(1)	2.14(1)	3.054(3)	175(5)
N1–H _{1D} ···Cl ₂ ⁱ	0.90(1)	2.15(1)	3.030(2)	167(2)
C1–H _{1B} ···Cl ₃ ⁱⁱ	0.98	2.81	3.717(4)	154
C2–H _{2B} ···Cl ₁	0.99	2.47	3.456(3)	174
C6–H _{6B} ···Cl ₃ ⁱⁱ	0.99	2.73	3.668(3)	158
C10–H _{10A} ···O _{1A} ⁱⁱⁱ	0.95	2.56	3.491(4)	166
C16–H _{16A} ···Cl ₃ ⁱⁱⁱ	0.99	2.73	3.576(3)	143
C20–H _{20A} ···O ₁ ^{iv}	0.95	2.49	3.224(3)	134
C21–H _{21A} ···Cl ₁ ⁱ	0.95	2.68	3.602(3)	164
C25–H _{25A} ···O _{1A} ^v	0.99	2.45	3.435(4)	171
C27–H _{27A} ···Cl ₂ ⁱⁱ	0.99	2.74	3.576(3)	143
C27–H _{27B} ···Cl ₁ ⁱ	0.99	2.66	3.621(3)	164

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

3. Supramolecular features

In the crystal, N–H···Cl hydrogen bonds (Table 1) link cations and anions (Fig. 2) into [100] chains. The chains are cross-linked by C–H···Cl and C–H···O interactions, forming a three-dimensional network. In addition, the existence of short (compared to the sum of the van der Waals radii of the corresponding pairs of atoms; Rowland & Taylor, 1996) intermolecular water-to-water O···O and water-to-chloride O···Cl contacts presumably correspond to O–H···X hydrogen bonds, although the water H atoms could not be located in the present study.

**Figure 2**

Projection of the crystal packing of the title compound along the c axis. Dashed lines denote strong intermolecular N–H···Cl hydrogen bonds. Water molecules have been omitted for clarity.

Table 2
Experimental details.

Crystal data	
Chemical formula	C ₂₈ H ₄₀ N ₃ O ³⁺ ·3Cl ⁻ ·2H ₂ O
M _r	577.01
Crystal system, space group	Triclinic, P <bar{1}< td=""></bar{1}<>
Temperature (K)	100
a, b, c (Å)	10.0933 (5), 12.0661 (6), 13.7576 (6)
α, β, γ (°)	97.759 (1), 110.795 (1), 102.733 (1)
V (Å ³)	1485.46 (12)
Z	2
Radiation type	Mo K α
μ (mm ⁻¹)	0.34
Crystal size (mm)	0.18 × 0.12 × 0.10
Data collection	
Diffractometer	Bruker APEXII CCD
Absorption correction	Multi-scan (<i>SADABS</i> ; Bruker, 2007)
T _{min} , T _{max}	0.941, 0.967
No. of measured, independent and observed [I > 2σ(I)] reflections	11768, 5787, 5056
R _{int}	0.016
(sin θ/λ) _{max} (Å ⁻¹)	0.617
Refinement	
R[F ² > 2σ(F ²)], wR(F ²), S	0.058, 0.164, 1.05
No. of reflections	5787
No. of parameters	356
No. of restraints	3
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
Δρ _{max} , Δρ _{min} (e Å ⁻³)	1.13, -0.62

Computer programs: *APEX2* and *SAINT* (Bruker, 2007), *SHELXS97*, *SHELXL97* and *SHELXTL* (Sheldrick, 2008).

4. Database survey

A search in the Cambridge Structural Database (Groom & Allen, 2014) for structures of piperidone with the amino substituents revealed eight hits with two salt structures of the oxopiperidinium iodide (Jia *et al.*, 1989; Nesterov *et al.*, 2007). Among these, there is a starting compound in which both diethylamino substituents participate in a conjugation with aromatic rings (Nesterov *et al.*, 2003).

5. Synthesis and crystallization

The starting compound (3*E*,5*E*)-3,5-bis[4-(diethylamino)-benzylidene]-1-methyl-4-piperidone was obtained according to a literature procedure (Nesterov *et al.*, 2003). The relatively stable colorless crystals of the investigated salt were obtained by slow evaporation of the solution of the above piperidone

from a mixture of ethanol and hydrochloric acid over several days.

6. Refinement

Crystal data, data collection, and structure refinement details are summarized in Table 2. All C-bound H-atoms were placed in idealized positions and allowed to ride on their parent atom: C—H = 0.95, 0.99 and 0.98 Å for CH, CH₂ and CH₃ H atoms, respectively, with U_{iso}(H) = k × U_{eq}(C), where k = 1.2 for CH and CH₂ and 1.5 for CH₃ H atoms. All N-bound H atoms were located using difference Fourier maps, but in the final refinement their distances were constrained at 0.90 Å (DFIX). H atoms of the two water molecules were not localized properly, since they appeared to be disordered over several positions. These H atoms were therefore removed from the refinement, but they were still included in the resulting chemical formula. Atom Cl3 is disordered over two positions in a 0.895 (4):0.105 (4) ratio.

Acknowledgements

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supporting information

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Crystal structure of (*3E,5E*)-3,5-bis[4-(diethylazaniumyl)benzylidene]-1-methyl-4-oxopiperidin-1-i^{um} trichloride dihydrate: a potential biophotonic material

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Computing details

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT* (Bruker, 2007); 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).

(*3E,5E*)-3,5-Bis[4-(diethylazaniumyl)benzylidene]-1-methyl- 4-oxopiperidin-1-i^{um}

Crystal data

$C_{28}H_{40}N_3O^{3+}\cdot 3Cl^- \cdot 2H_2O$	$Z = 2$
$M_r = 577.01$	$F(000) = 616$
Triclinic, $P\bar{1}$	$D_x = 1.290 \text{ Mg m}^{-3}$
Hall symbol: -P 1	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
$a = 10.0933 (5) \text{ \AA}$	Cell parameters from 4567 reflections
$b = 12.0661 (6) \text{ \AA}$	$\theta = 2.5\text{--}26.0^\circ$
$c = 13.7576 (6) \text{ \AA}$	$\mu = 0.34 \text{ mm}^{-1}$
$\alpha = 97.759 (1)^\circ$	$T = 100 \text{ K}$
$\beta = 110.795 (1)^\circ$	Block, colourless
$\gamma = 102.733 (1)^\circ$	$0.18 \times 0.12 \times 0.10 \text{ mm}$
$V = 1485.46 (12) \text{ \AA}^3$	

Data collection

Bruker APEXII CCD	11768 measured reflections
diffractometer	5787 independent reflections
Radiation source: fine-focus sealed tube	5056 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{\text{int}} = 0.016$
φ and ω scans	$\theta_{\text{max}} = 26.0^\circ, \theta_{\text{min}} = 1.6^\circ$
Absorption correction: multi-scan	$h = -12 \rightarrow 12$
(SADABS; Bruker, 2007)	$k = -14 \rightarrow 14$
$T_{\text{min}} = 0.941, T_{\text{max}} = 0.967$	$l = -16 \rightarrow 16$

Refinement

Refinement on F^2	356 parameters
Least-squares matrix: full	3 restraints
$R[F^2 > 2\sigma(F^2)] = 0.058$	Primary atom site location: structure-invariant direct methods
$wR(F^2) = 0.164$	Secondary atom site location: difference Fourier map
$S = 1.05$	
5787 reflections	

Hydrogen site location: inferred from neighbouring sites

H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.091P)^2 + 1.9P]$$

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

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

$$\Delta\rho_{\max} = 1.13 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.62 \text{ e \AA}^{-3}$$

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 $F^2 > 2\text{sigma}(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C11	0.69912 (7)	0.36862 (6)	0.70014 (6)	0.0379 (2)	
Cl2	1.10763 (7)	0.16702 (5)	0.19807 (5)	0.02863 (18)	
Cl3	0.41255 (10)	0.74605 (6)	0.84491 (6)	0.0319 (3)	0.895 (4)
Cl3A	0.4816 (15)	0.7207 (9)	0.8555 (8)	0.059 (3)*	0.105 (4)
O1	0.2253 (2)	-0.04179 (15)	0.41664 (15)	0.0288 (4)	
N1	0.4116 (2)	0.26939 (17)	0.37806 (17)	0.0229 (4)	
H1D	0.3294 (19)	0.239 (2)	0.3181 (14)	0.017 (6)*	
N2	0.8498 (2)	-0.06652 (19)	0.13943 (17)	0.0251 (5)	
H2C	0.924 (3)	0.0015 (16)	0.165 (2)	0.034 (8)*	
N3	0.1387 (3)	0.5404 (2)	0.78810 (19)	0.0308 (5)	
H3A	0.221 (3)	0.602 (3)	0.809 (4)	0.093 (17)*	
C1	0.4892 (4)	0.3863 (2)	0.3704 (3)	0.0365 (7)	
H1A	0.4236	0.4366	0.3621	0.055*	
H1B	0.5157	0.3772	0.3084	0.055*	
H1C	0.5792	0.4220	0.4356	0.055*	
C2	0.3736 (3)	0.2804 (2)	0.4734 (2)	0.0237 (5)	
H2A	0.3144	0.3362	0.4704	0.028*	
H2B	0.4656	0.3114	0.5392	0.028*	
C3	0.2870 (3)	0.1637 (2)	0.47709 (18)	0.0204 (5)	
C4	0.3029 (3)	0.0544 (2)	0.42288 (19)	0.0213 (5)	
C5	0.4212 (3)	0.0674 (2)	0.37980 (18)	0.0204 (5)	
C6	0.5053 (3)	0.1887 (2)	0.3839 (2)	0.0249 (5)	
H6A	0.5953	0.2179	0.4513	0.030*	
H6B	0.5368	0.1866	0.3235	0.030*	
C7	0.4468 (3)	-0.0299 (2)	0.33955 (18)	0.0204 (5)	
H7A	0.3874	-0.1019	0.3414	0.024*	
C8	0.5559 (3)	-0.0379 (2)	0.29304 (19)	0.0223 (5)	
C9	0.5226 (3)	-0.1372 (2)	0.2139 (2)	0.0261 (5)	
H9A	0.4346	-0.1988	0.1951	0.031*	
C10	0.6172 (3)	-0.1468 (2)	0.1624 (2)	0.0299 (6)	

H10A	0.5930	-0.2134	0.1070	0.036*
C11	0.7474 (3)	-0.0574 (2)	0.1932 (2)	0.0255 (5)
C12	0.7871 (3)	0.0384 (2)	0.2743 (2)	0.0273 (5)
H12A	0.8786	0.0970	0.2960	0.033*
C13	0.6914 (3)	0.0482 (2)	0.3240 (2)	0.0274 (5)
H13A	0.7179	0.1145	0.3803	0.033*
C14	0.9212 (3)	-0.1619 (2)	0.1686 (2)	0.0294 (6)
H14A	0.9628	-0.1503	0.2474	0.035*
H14B	0.8442	-0.2384	0.1379	0.035*
C15	1.0423 (3)	-0.1647 (3)	0.1296 (2)	0.0335 (6)
H15A	1.0940	-0.2198	0.1599	0.050*
H15B	1.1127	-0.0865	0.1521	0.050*
H15C	0.9992	-0.1897	0.0514	0.050*
C16	0.7779 (3)	-0.0775 (3)	0.0204 (2)	0.0314 (6)
H16A	0.6993	-0.1530	-0.0129	0.038*
H16B	0.8528	-0.0778	-0.0104	0.038*
C17	0.7110 (3)	0.0207 (3)	-0.0065 (2)	0.0357 (6)
H17A	0.6672	0.0103	-0.0844	0.054*
H17B	0.7883	0.0957	0.0258	0.054*
H17C	0.6340	0.0196	0.0214	0.054*
C18	0.1975 (3)	0.1549 (2)	0.53023 (19)	0.0205 (5)
H18A	0.1401	0.0785	0.5243	0.025*
C19	0.1804 (3)	0.2522 (2)	0.59652 (19)	0.0203 (5)
C20	0.0408 (3)	0.2538 (2)	0.5932 (2)	0.0247 (5)
H20A	-0.0434	0.1902	0.5488	0.030*
C21	0.0235 (3)	0.3478 (3)	0.6542 (2)	0.0300 (6)
H21A	-0.0719	0.3494	0.6503	0.036*
C22	0.1474 (3)	0.4382 (2)	0.7200 (2)	0.0285 (6)
C23	0.2872 (3)	0.4361 (2)	0.7293 (2)	0.0276 (5)
H23A	0.3716	0.4972	0.7780	0.033*
C24	0.3036 (3)	0.3444 (2)	0.6671 (2)	0.0245 (5)
H24A	0.3998	0.3437	0.6723	0.029*
C25	0.1258 (3)	0.5097 (2)	0.8880 (2)	0.0312 (6)
H25A	0.1994	0.4683	0.9191	0.037*
H25B	0.0260	0.4562	0.8689	0.037*
C26	0.1511 (5)	0.6178 (3)	0.9711 (3)	0.0505 (9)
H26A	0.1669	0.5976	1.0402	0.076*
H26B	0.0643	0.6469	0.9486	0.076*
H26C	0.2385	0.6784	0.9779	0.076*
C27	0.0234 (3)	0.5949 (2)	0.7310 (2)	0.0336 (6)
H27A	0.0316	0.6662	0.7805	0.040*
H27B	-0.0757	0.5397	0.7110	0.040*
C28	0.0356 (4)	0.6268 (3)	0.6323 (2)	0.0416 (7)
H28A	-0.0141	0.6870	0.6145	0.062*
H28B	-0.0115	0.5574	0.5728	0.062*
H28C	0.1403	0.6569	0.6451	0.062*
O1A	0.5901 (3)	0.6022 (2)	0.9885 (3)	0.0693 (8)
O2A	0.7711 (5)	0.5914 (3)	0.8786 (3)	0.0984 (12)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0277 (3)	0.0416 (4)	0.0486 (4)	0.0113 (3)	0.0192 (3)	0.0105 (3)
Cl2	0.0306 (3)	0.0305 (3)	0.0273 (3)	0.0124 (3)	0.0130 (3)	0.0049 (2)
Cl3	0.0304 (5)	0.0243 (4)	0.0365 (5)	0.0025 (3)	0.0142 (3)	-0.0012 (3)
O1	0.0339 (10)	0.0195 (9)	0.0368 (10)	0.0005 (7)	0.0247 (8)	0.0012 (7)
N1	0.0286 (11)	0.0174 (10)	0.0253 (11)	0.0029 (8)	0.0169 (9)	0.0023 (8)
N2	0.0276 (11)	0.0259 (11)	0.0264 (11)	0.0107 (9)	0.0140 (9)	0.0071 (9)
N3	0.0442 (14)	0.0291 (12)	0.0295 (12)	0.0167 (11)	0.0221 (11)	0.0090 (10)
C1	0.0512 (18)	0.0193 (12)	0.0517 (18)	0.0051 (12)	0.0380 (15)	0.0080 (12)
C2	0.0297 (13)	0.0205 (11)	0.0241 (12)	0.0045 (10)	0.0174 (10)	0.0016 (9)
C3	0.0209 (11)	0.0209 (11)	0.0190 (11)	0.0048 (9)	0.0089 (9)	0.0033 (9)
C4	0.0226 (11)	0.0223 (12)	0.0198 (11)	0.0040 (9)	0.0114 (10)	0.0029 (9)
C5	0.0200 (11)	0.0221 (11)	0.0202 (11)	0.0042 (9)	0.0109 (9)	0.0039 (9)
C6	0.0248 (12)	0.0208 (12)	0.0336 (13)	0.0047 (10)	0.0193 (11)	0.0020 (10)
C7	0.0213 (11)	0.0222 (11)	0.0190 (11)	0.0061 (9)	0.0091 (9)	0.0059 (9)
C8	0.0281 (12)	0.0247 (12)	0.0232 (12)	0.0141 (10)	0.0146 (10)	0.0114 (10)
C9	0.0306 (13)	0.0242 (12)	0.0304 (13)	0.0115 (10)	0.0168 (11)	0.0094 (10)
C10	0.0390 (15)	0.0258 (13)	0.0312 (14)	0.0159 (11)	0.0181 (12)	0.0049 (11)
C11	0.0277 (13)	0.0311 (13)	0.0272 (13)	0.0132 (11)	0.0160 (11)	0.0154 (11)
C12	0.0255 (12)	0.0276 (13)	0.0340 (14)	0.0094 (10)	0.0161 (11)	0.0090 (11)
C13	0.0268 (13)	0.0293 (13)	0.0314 (14)	0.0112 (11)	0.0156 (11)	0.0081 (11)
C14	0.0357 (14)	0.0277 (13)	0.0323 (14)	0.0166 (11)	0.0163 (12)	0.0106 (11)
C15	0.0347 (15)	0.0333 (14)	0.0391 (15)	0.0163 (12)	0.0185 (13)	0.0077 (12)
C16	0.0291 (13)	0.0477 (16)	0.0205 (13)	0.0132 (12)	0.0122 (11)	0.0078 (11)
C17	0.0347 (15)	0.0444 (16)	0.0331 (15)	0.0120 (13)	0.0170 (12)	0.0150 (13)
C18	0.0218 (11)	0.0189 (11)	0.0215 (11)	0.0045 (9)	0.0098 (9)	0.0052 (9)
C19	0.0261 (12)	0.0207 (11)	0.0208 (11)	0.0081 (9)	0.0147 (10)	0.0092 (9)
C20	0.0229 (12)	0.0305 (13)	0.0242 (12)	0.0071 (10)	0.0131 (10)	0.0087 (10)
C21	0.0350 (14)	0.0423 (15)	0.0314 (14)	0.0229 (12)	0.0241 (12)	0.0185 (12)
C22	0.0463 (16)	0.0261 (13)	0.0242 (12)	0.0179 (12)	0.0204 (12)	0.0115 (10)
C23	0.0367 (14)	0.0238 (12)	0.0247 (12)	0.0056 (11)	0.0170 (11)	0.0049 (10)
C24	0.0265 (12)	0.0253 (12)	0.0240 (12)	0.0051 (10)	0.0145 (10)	0.0050 (10)
C25	0.0481 (16)	0.0274 (13)	0.0302 (14)	0.0151 (12)	0.0256 (13)	0.0104 (11)
C26	0.093 (3)	0.0444 (18)	0.0347 (16)	0.0295 (19)	0.0425 (18)	0.0120 (14)
C27	0.0432 (16)	0.0266 (13)	0.0366 (15)	0.0180 (12)	0.0171 (13)	0.0087 (11)
C28	0.057 (2)	0.0377 (16)	0.0338 (16)	0.0221 (15)	0.0162 (14)	0.0130 (13)
O1A	0.0472 (14)	0.0392 (13)	0.104 (2)	0.0153 (11)	0.0153 (15)	-0.0023 (14)
O2A	0.142 (4)	0.075 (2)	0.076 (2)	0.027 (2)	0.044 (2)	0.0198 (18)

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

O1—C4	1.221 (3)	C13—H13A	0.9500
N1—C2	1.489 (3)	C14—C15	1.504 (4)
N1—C6	1.491 (3)	C14—H14A	0.9900
N1—C1	1.492 (3)	C14—H14B	0.9900
N1—H1D	0.895 (10)	C15—H15A	0.9800

N2—C11	1.483 (3)	C15—H15B	0.9800
N2—C16	1.510 (3)	C15—H15C	0.9800
N2—C14	1.513 (3)	C16—C17	1.513 (4)
N2—H2C	0.907 (10)	C16—H16A	0.9900
N3—C22	1.483 (3)	C16—H16B	0.9900
N3—C27	1.494 (4)	C17—H17A	0.9800
N3—C25	1.512 (3)	C17—H17B	0.9800
N3—H3A	0.912 (10)	C17—H17C	0.9800
C1—H1A	0.9800	C18—C19	1.468 (3)
C1—H1B	0.9800	C18—H18A	0.9500
C1—H1C	0.9800	C19—C20	1.398 (3)
C2—C3	1.503 (3)	C19—C24	1.402 (4)
C2—H2A	0.9900	C20—C21	1.399 (4)
C2—H2B	0.9900	C20—H20A	0.9500
C3—C18	1.345 (3)	C21—C22	1.379 (4)
C3—C4	1.495 (3)	C21—H21A	0.9500
C4—C5	1.499 (3)	C22—C23	1.377 (4)
C5—C7	1.346 (3)	C23—C24	1.382 (4)
C5—C6	1.503 (3)	C23—H23A	0.9500
C6—H6A	0.9900	C24—H24A	0.9500
C6—H6B	0.9900	C25—C26	1.521 (4)
C7—C8	1.470 (3)	C25—H25A	0.9900
C7—H7A	0.9500	C25—H25B	0.9900
C8—C9	1.394 (4)	C26—H26A	0.9800
C8—C13	1.404 (4)	C26—H26B	0.9800
C9—C10	1.391 (4)	C26—H26C	0.9800
C9—H9A	0.9500	C27—C28	1.499 (4)
C10—C11	1.388 (4)	C27—H27A	0.9900
C10—H10A	0.9500	C27—H27B	0.9900
C11—C12	1.367 (4)	C28—H28A	0.9800
C12—C13	1.383 (4)	C28—H28B	0.9800
C12—H12A	0.9500	C28—H28C	0.9800
C2—N1—C6	109.88 (19)	N2—C14—H14A	108.9
C2—N1—C1	110.58 (19)	C15—C14—H14B	108.9
C6—N1—C1	110.6 (2)	N2—C14—H14B	108.9
C2—N1—H1D	110.5 (18)	H14A—C14—H14B	107.7
C6—N1—H1D	107.7 (17)	C14—C15—H15A	109.5
C1—N1—H1D	107.5 (17)	C14—C15—H15B	109.5
C11—N2—C16	112.58 (19)	H15A—C15—H15B	109.5
C11—N2—C14	110.55 (19)	C14—C15—H15C	109.5
C16—N2—C14	113.6 (2)	H15A—C15—H15C	109.5
C11—N2—H2C	108 (2)	H15B—C15—H15C	109.5
C16—N2—H2C	105 (2)	N2—C16—C17	112.6 (2)
C14—N2—H2C	107 (2)	N2—C16—H16A	109.1
C22—N3—C27	114.4 (2)	C17—C16—H16A	109.1
C22—N3—C25	110.3 (2)	N2—C16—H16B	109.1
C27—N3—C25	112.8 (2)	C17—C16—H16B	109.1

C22—N3—H3A	112 (3)	H16A—C16—H16B	107.8
C27—N3—H3A	99 (3)	C16—C17—H17A	109.5
C25—N3—H3A	108 (3)	C16—C17—H17B	109.5
N1—C1—H1A	109.5	H17A—C17—H17B	109.5
N1—C1—H1B	109.5	C16—C17—H17C	109.5
H1A—C1—H1B	109.5	H17A—C17—H17C	109.5
N1—C1—H1C	109.5	H17B—C17—H17C	109.5
H1A—C1—H1C	109.5	C3—C18—C19	126.0 (2)
H1B—C1—H1C	109.5	C3—C18—H18A	117.0
N1—C2—C3	110.49 (19)	C19—C18—H18A	117.0
N1—C2—H2A	109.6	C20—C19—C24	118.1 (2)
C3—C2—H2A	109.6	C20—C19—C18	120.7 (2)
N1—C2—H2B	109.6	C24—C19—C18	121.1 (2)
C3—C2—H2B	109.6	C19—C20—C21	120.9 (2)
H2A—C2—H2B	108.1	C19—C20—H20A	119.5
C18—C3—C4	118.8 (2)	C21—C20—H20A	119.5
C18—C3—C2	121.6 (2)	C22—C21—C20	118.7 (2)
C4—C3—C2	119.6 (2)	C22—C21—H21A	120.6
O1—C4—C3	121.3 (2)	C20—C21—H21A	120.6
O1—C4—C5	121.4 (2)	C23—C22—C21	121.6 (2)
C3—C4—C5	117.3 (2)	C23—C22—N3	116.2 (2)
C7—C5—C4	118.3 (2)	C21—C22—N3	122.1 (2)
C7—C5—C6	123.4 (2)	C22—C23—C24	119.4 (3)
C4—C5—C6	118.2 (2)	C22—C23—H23A	120.3
N1—C6—C5	110.69 (19)	C24—C23—H23A	120.3
N1—C6—H6A	109.5	C23—C24—C19	121.0 (2)
C5—C6—H6A	109.5	C23—C24—H24A	119.5
N1—C6—H6B	109.5	C19—C24—H24A	119.5
C5—C6—H6B	109.5	N3—C25—C26	111.9 (2)
H6A—C6—H6B	108.1	N3—C25—H25A	109.2
C5—C7—C8	127.6 (2)	C26—C25—H25A	109.2
C5—C7—H7A	116.2	N3—C25—H25B	109.2
C8—C7—H7A	116.2	C26—C25—H25B	109.2
C9—C8—C13	118.3 (2)	H25A—C25—H25B	107.9
C9—C8—C7	117.8 (2)	C25—C26—H26A	109.5
C13—C8—C7	123.8 (2)	C25—C26—H26B	109.5
C10—C9—C8	120.6 (2)	H26A—C26—H26B	109.5
C10—C9—H9A	119.7	C25—C26—H26C	109.5
C8—C9—H9A	119.7	H26A—C26—H26C	109.5
C11—C10—C9	118.9 (2)	H26B—C26—H26C	109.5
C11—C10—H10A	120.6	N3—C27—C28	113.5 (2)
C9—C10—H10A	120.6	N3—C27—H27A	108.9
C12—C11—C10	122.0 (2)	C28—C27—H27A	108.9
C12—C11—N2	118.4 (2)	N3—C27—H27B	108.9
C10—C11—N2	119.6 (2)	C28—C27—H27B	108.9
C11—C12—C13	118.7 (2)	H27A—C27—H27B	107.7
C11—C12—H12A	120.6	C27—C28—H28A	109.5
C13—C12—H12A	120.6	C27—C28—H28B	109.5

C12—C13—C8	121.3 (2)	H28A—C28—H28B	109.5
C12—C13—H13A	119.3	C27—C28—H28C	109.5
C8—C13—H13A	119.3	H28A—C28—H28C	109.5
C15—C14—N2	113.3 (2)	H28B—C28—H28C	109.5
C15—C14—H14A	108.9		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N2—H2C···Cl2	0.91 (1)	2.27 (1)	3.166 (2)	169 (3)
N3—H3A···Cl3	0.91 (1)	2.14 (1)	3.054 (3)	175 (5)
N1—H1D···Cl2 ⁱ	0.90 (1)	2.15 (1)	3.030 (2)	167 (2)
C1—H1B···Cl3 ⁱⁱ	0.98	2.81	3.717 (4)	154
C2—H2B···Cl1	0.99	2.47	3.456 (3)	174
C6—H6B···Cl3 ⁱⁱ	0.99	2.73	3.668 (3)	158
C10—H10A···O1A ⁱⁱⁱ	0.95	2.56	3.491 (4)	166
C16—H16A···Cl3 ⁱⁱⁱ	0.99	2.73	3.576 (3)	143
C20—H20A···O1 ^{iv}	0.95	2.49	3.224 (3)	134
C21—H21A···Cl1 ⁱ	0.95	2.68	3.602 (3)	164
C25—H25A···O1A ^v	0.99	2.45	3.435 (4)	171
C27—H27A···Cl2 ⁱⁱ	0.99	2.74	3.576 (3)	143
C27—H27B···Cl1 ⁱ	0.99	2.66	3.621 (3)	164

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