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Bis(4-hydroxy-*N*-isopropyl-*N*-methyltryptammonium) fumarate: a new crystalline form of miprocin

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The title compound, bis(4-hydroxy-*N*-isopropyl-*N*-methyltryptammonium) (4-HO-MiPT) fumarate (systematic name: bis[[2-(4-hydroxy-1*H*-indol-3-yl)ethyl](methyl)propan-2-ylazanium] but-2-enedioate), $2C_{14}H_{21}N_2O^+ \cdot C_4H_2O_4^{2-}$, has a singly protonated tryptammonium cation and one half of a fumarate dianion in the asymmetric unit. The tryptammonium and fumarate ions are held together in one-dimensional chains by $N-H \cdots O$ and $O-H \cdots O$ hydrogen bonds. These chains are a combination of $R_4^2(20)$ rings, and $C_2^2(15)$ and $C_4^4(30)$ parallel chains along (110). They are further consolidated by $N-H \cdots \pi$ interactions. There are two two-component types of disorder impacting the tryptammonium fragment with a 0.753 (7):0.247 (7) occupancy ratio and one of the fumarate oxygen atoms with a 0.73 (8):0.27 (8) ratio.

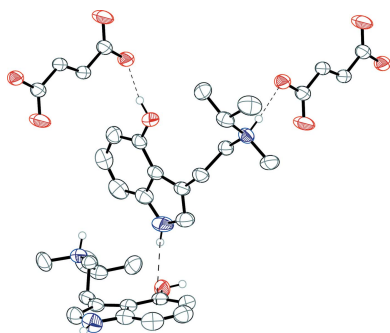
1. Chemical context

A wide variety of naturally occurring organisms, including over 200 species of 'magic' mushrooms, contain psychoactive tryptamine compounds (Stamets, 1996). Of these compounds, psilocybin has received the most scientific and commercial attention because of recent studies demonstrating its potential for treating mood disorders including addiction, anxiety, depression and PTSD (Johnson & Griffiths, 2017; Carhart-Harris & Goodwin, 2017).

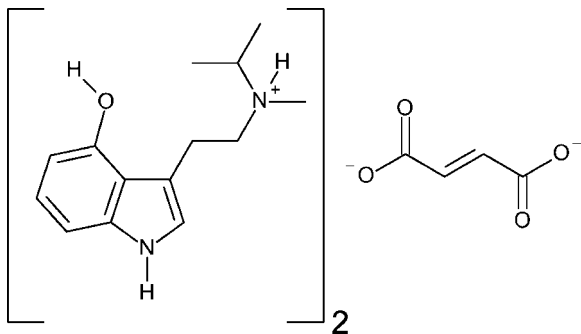
Although psilocybin is currently classified as a schedule I drug, the US Food and Drug Administration recently designated treatment using psilocybin a 'breakthrough therapy'. This status has allowed psilocybin to be administered in clinical trials to treat major depressive disorder and treatment-resistant depression (Feltman, 2019). Recent reports also suggest that psychedelic microdosing can improve memory, attention and sociability (Cameron, *et al.* 2020).

Psilocybin is one of at least ten psychoactive tryptamines present in 'magic' mushrooms, with natural psilocybin analogs being identified as recently as 2019 (Lenz *et al.*, 2017; Blei *et al.*, 2020). Variations in the three-dimensional structure of these natural analogs (as well as synthetic analogs) correlate with differences in their cellular and clinical pharmacology through their structure–activity relationship (SAR) (Nichols, 2018). Understanding the SAR for psilocybin analogs requires the attainment of accurate information about each compound's 3D structure, best provided through single crystal X-ray diffraction.

Last year, we reported the structure of 4-acetoxy-*N,N*-dimethyl tryptamine (4-AcO-DMT) fumarate, which is a synthetic analogue of psilocybin. The compound crystallized as a



one-to-one tryptammonium/hydrofumarate salt (Chadeayne *et al.*, 2019c). We later synthesized bis(4-acetoxy-*N,N*-dimethyltryprammonium)fumarate by treating 4-AcO-DMT fumarate with one half equivalent of lead(II) acetate, precipitating half of the fumarate dianions as lead(II) fumarate (Chadeayne, Golen & Manke, 2019a).



4-Hydroxy-*N*-methyl-*N*-isopropyltryptamine (4-HO-MiPT), aka 'miprocin', is a psilocybin analogue. Its synthesis was first reported in 1981 by Repke and co-workers (Repke *et al.*, 1981); its psychedelic effects were later described in collaboration with Alexander Shulgin (Repke *et al.*, 1985). Miprocin is reported to produce an experience that is both relaxing, stoning and mildly sedating with a marked physical stimulation that distinguishes it from related substances such as psilocybin mushrooms. In a report last year, we presented the first structure of 4-HO-MiPT (Chadeayne, Pham *et al.*, 2019a), which crystallizes as the hydrofumarate monohydrate. Herein we report the reaction of this salt with lead(II) acetate to generate the 4-hydroxy-*N*-isopropyl-*N*-methyltryptammonium/fumarate compound in a 2:1 ratio. The solid state structure of the new salt is presented here.

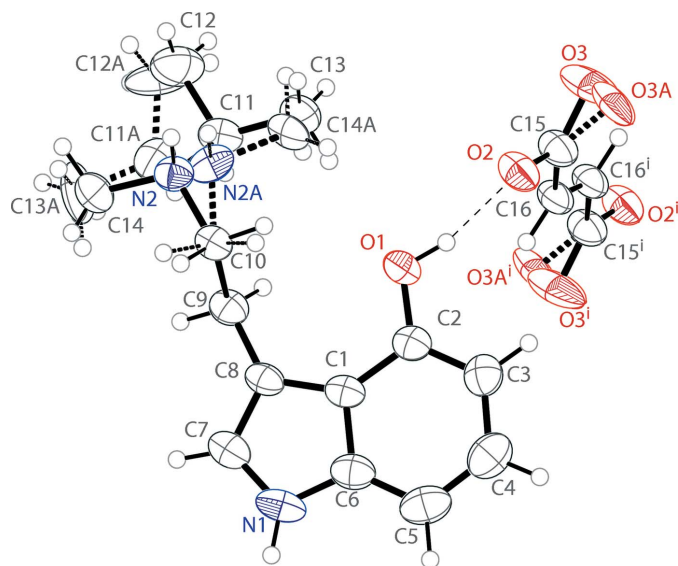


Figure 1

The molecular structure of bis(4-hydroxy-*N*-isopropyl-*N*-methyltryptammonium)fumarate, showing the atom labeling. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen bonds are shown as dashed lines. Symmetry code: (i) $1 - x, 1 - y, 1 - z$.

2. Structural commentary

The asymmetric unit of bis(4-hydroxy-*N*-isopropyl-*N*-methyltryptammonium) fumarate contains one tryptammonium cation and one half of a fumarate dianion (Fig. 1). The cation possesses a near planar indole, with mean deviation from planarity of 0.014 Å. The methylamino group is turned away from this plane, with a C1–C8–C9–C10 torsion angle of $-74.2(2)^\circ$. The *N*-isopropyl-*N*-methyltryptammonium group is disordered over two orientations in a 0.753(7):0.247(7) ratio, with the two moieties related to each other by a pseudo-mirror operation. In solution, the two conformations are most likely interconverting into each other by rapid de- and reprotonation. One oxygen atom of the half fumarate anion is also disordered over two positions in a 0.73(8):0.27(8) ratio. Half of the fumarate dianion is present in the asymmetric unit, with the other half generated by inversion; it is slightly distorted from planarity with r.m.s. deviations of 0.020 and 0.070 Å for the two components. The carboxylate unit is fairly delocalized, with C–O distances ranging from 1.251(10) to 1.284(2) Å.

3. Supramolecular features

There are N2–H2···O2 and N2A–H2A···O2 hydrogen bonds between the two configurations of the ammonium cations and one fumarate oxygen. These two different N–H···O hydrogen bonds, resulting from the disorder, are also likely to be what produces the fumarate disorder. There is an O1–H1···O2 hydrogen bond between the phenol hydroxy group and one fumarate oxygen atom. Two tryptammonium

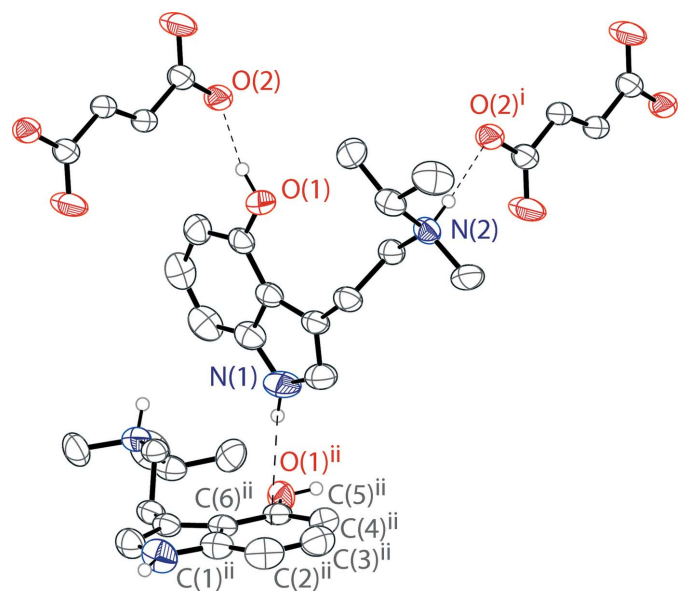


Figure 2

The hydrogen bonding of the tryptammonium cation in the structure of the title compound (Table 1), with hydrogen bonds shown as dashed lines. There is also an N–H··· π interaction shown. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen atoms not involved in hydrogen bonds are omitted for clarity. Symmetry codes: (i) $\frac{1}{2} - x, \frac{3}{2} - y, 1 - z$, (ii) $\frac{1}{2} - x, -\frac{1}{2} + y, \frac{3}{2} - z$.

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

C_g2 is the centroid of the C1–C6 ring.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1–H1 \cdots O2	0.89 (1)	1.75 (1)	2.618 (2)	165 (2)
N2–H2 \cdots O2 ⁱ	0.88 (1)	1.85 (1)	2.730 (5)	175 (3)
N2A–H2A \cdots O2 ⁱ	0.87 (1)	1.89 (4)	2.727 (12)	160 (11)
N1–H1A \cdots Cg2 ⁱⁱ	0.87 (1)	2.78 (2)	3.552 (3)	148 (2)

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

cations and two fumarate anions are joined together through the N–H \cdots O and O–H \cdots O hydrogen bonds (Fig. 2), forming rings with graph-set notation $R_4^2(20)$ (Etter *et al.*, 1990). The rings are joined together by two parallel chains along (110). These chains have graph-set notation $C_2^2(15)$ and

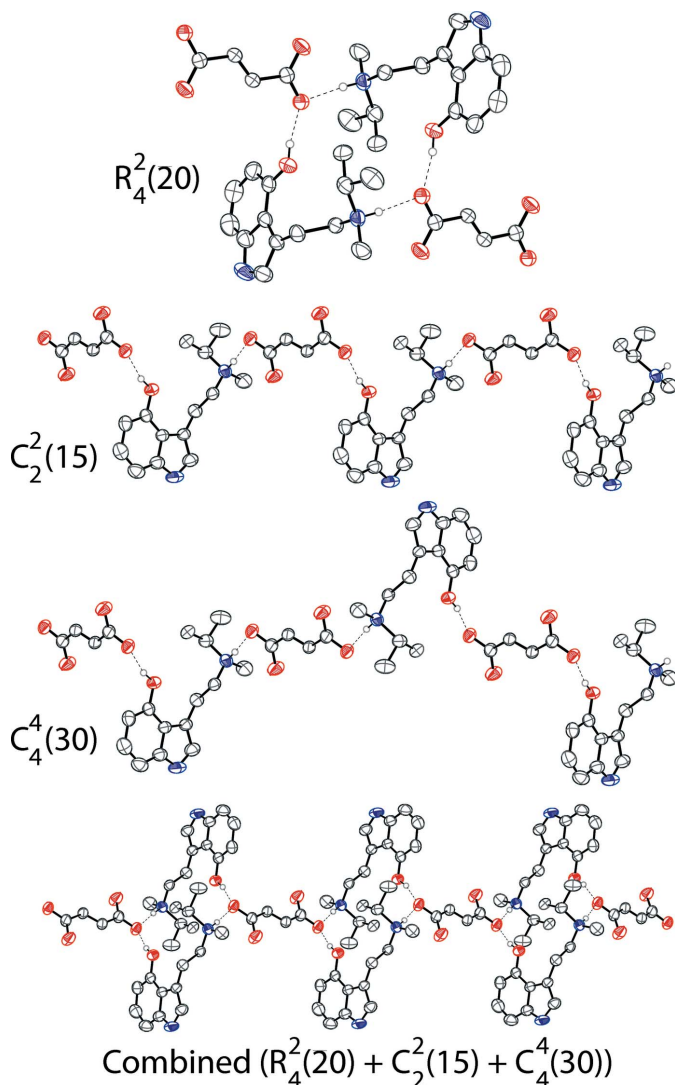


Figure 3
The hydrogen-bonding network along (110), which consists of $R_4^2(20)$ rings that are joined together by two parallel $C_2^2(15)$ and $C_4^4(30)$ chains. The three components described in graph-set notation and the combined chain are shown. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen atoms not involved in hydrogen bonding are omitted for clarity. Hydrogen bonds are shown as dashed lines.

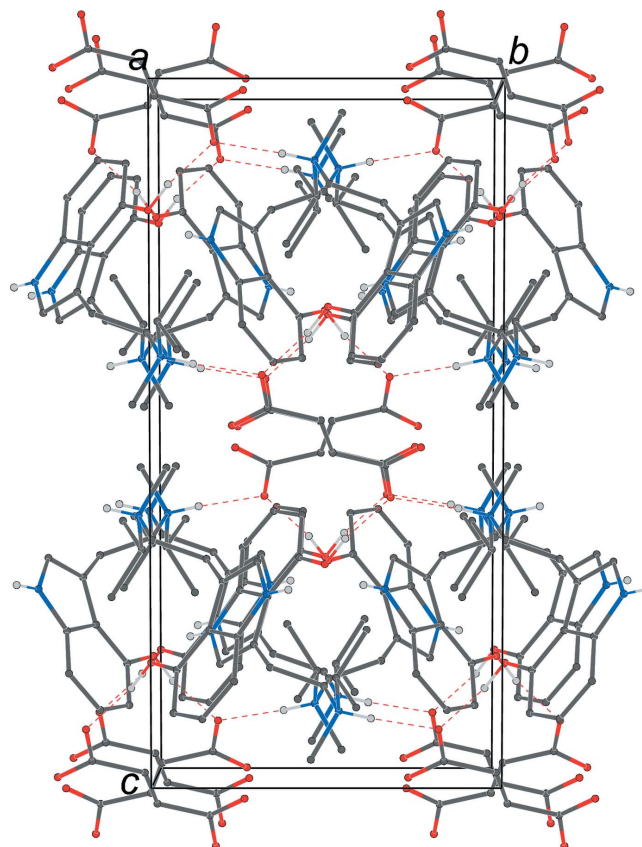


Figure 4
The crystal packing of the title compound, viewed along the a axis. The N–H \cdots O and O–H \cdots O hydrogen bonds (Table 1) are shown as dashed lines. Hydrogen atoms not involved in hydrogen bonding are omitted for clarity.

$C_4^4(30)$. The chains and rings are shown in Fig. 3. The ions are further linked through N–H \cdots π interactions between the indole N–H and the aromatic ring of the indole of another tryptammonium ion (Fig. 2). The hydrogen bonds in the system are outlined in Table 1. The packing of the compound is shown in Fig. 4.

4. Database survey

The structure of a number of neutral tryptamines have been reported, including psilocin (Petcher & Weber, 1974), psilocybin (Weber & Petcher, 1974), bufotenine (Falkenberg, 1972*b*), DMT (Falkenberg, 1972*a*) and MPT (Chadeayne, Golen & Manke, 2019*b*). A series of one-to-one tryptammonium hydrofumarate salts have been structurally characterized, including psilacetin (Chadeayne *et al.*, 2019*c*), miprocain and MiPT (Chadeayne, Pham *et al.*, 2019*a*). As discussed above, the two-to-one tryptammonium/fumarate salt of 4-AcO-DMT was previously prepared and its structure reported (Chadeayne, Golen & Manke, 2019*a*). The only other reported two-to-one tryptammonium fumarate salt was that of 4-HO-DPT, or procin (Chadeayne, Pham *et al.*, 2019*b*). The metrical parameters of the tryptammonium cations of 4-HO-MiPT are comparable to those observed for the other reported tryptamine structures.

Table 2
Experimental details.

Crystal data	
Chemical formula	C ₁₄ H ₂₁ N ₂ O ⁺ ·C ₂ H ₂ O ₂ [−]
<i>M_r</i>	290.35
Crystal system, space group	Monoclinic, <i>C2/c</i>
Temperature (K)	200
<i>a</i> , <i>b</i> , <i>c</i> (Å)	19.770 (13), 9.477 (6), 17.620 (12)
β (°)	105.78 (2)
<i>V</i> (Å ³)	3177 (4)
<i>Z</i>	8
Radiation type	Mo <i>K</i> α
μ (mm ^{−1})	0.08
Crystal size (mm)	0.25 × 0.2 × 0.1
Data collection	
Diffractometer	Bruker D8 Venture CMOS
Absorption correction	Multi-scan (<i>SADABS</i> ; Bruker, 2018)
<i>T_{min}</i> , <i>T_{max}</i>	0.692, 0.745
No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections	39417, 2890, 2007
<i>R_{int}</i>	0.086
(<i>sin</i> θ / λ) _{max} (Å ^{−1})	0.606
Refinement	
<i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.047, 0.105, 1.06
No. of reflections	2890
No. of parameters	265
No. of restraints	11
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})	0.15, −0.13

Computer programs: *APEX3* and *SAINT* (Bruker, 2018), *SHELXT2014* (Sheldrick, 2015a), *SHELXL2018* (Sheldrick, 2015b), *OLEX2* (Dolomanov *et al.*, 2009), and *PUBLICIF* (Westrip, 2010).

5. Synthesis and crystallization

61.2 mg of 4-HO-MiPT fumarate were dissolved in 10 mL of deionized water. 29.3 mg of lead(II) acetate was dissolved in 2 mL of deionized water and then added to the tryptamine solution. After sonication, a white precipitate formed. The powder was removed *via* vacuum filtration. The solvent was removed from the resulting solution *in vacuo* to yield a sticky powder. The powder was recrystallized from methanol to yield single crystals suitable for X-ray diffraction.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. Hydrogen atoms H1, H1A, H2 and H2A were found from a difference-Fourier map and were refined isotropically, using *DFIX* restraints with N–H distances of 0.87 (1) Å and an O–H distance of 0.88 (1) Å. Isotropic displacement parameters were set to 1.2*U*_{eq} of the parent indolic nitrogen atom and 1.5*U*_{eq} of the parent oxygen atom and the parent ammonium nitrogen atoms. All other hydrogen atoms were placed in calculated positions with appropriate carbon–hydrogen bond lengths: (*sp*²) 0.95 Å, (CH₃) 0.98 Å, (CH₂) 0.99 Å and (CH) 1.00 Å. Isotropic displacement parameters were set to 1.2*U*_{eq}(C) for *sp*², CH and CH₂ parent carbon atoms and 1.5*U*_{eq}(C-methyl). Atoms

N2 and C11–C14 were modeled as being disordered over two sets of sites [0.753 (7):0.247 (7)] and refined with *SADI* (0.03) restraints on C–C(methyl) and N–C(methyl) bonds to maintain consistent bond lengths in the disorder. Oxygen atom O3 was also modeled as disordered over two sites [0.73 (8):0.27 (8)].

Acknowledgements

We would like to thank Jerry Jasinski for useful advise in analyzing the supramolecular features. Financial statements and conflict of interest: This study was funded by CaaMTech, Inc. ARC reports an ownership interest in CaaMTech, Inc., which owns US and worldwide patent applications, covering new tryptamine compounds, compositions, formulations, novel crystalline forms, and methods of making and using the same.

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supporting information

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Computing details

Data collection: *APEX3* (Bruker, 2018); cell refinement: *SAINTE* (Bruker, 2018); data reduction: *SAINTE* (Bruker, 2018); program(s) used to solve structure: *SHELXT2014* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2018* (Sheldrick, 2015b); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *publCIF* (Westrip, 2010).

Bis[[2-(4-hydroxy-1*H*-indol-3-yl)ethyl](methyl)propan-2-ylazanium} but-2-enedioate

Crystal data

$C_{14}H_{21}N_2O^+ \cdot C_2HO_2^-$
 $M_r = 290.35$
 Monoclinic, *C2/c*
 $a = 19.770$ (13) Å
 $b = 9.477$ (6) Å
 $c = 17.620$ (12) Å
 $\beta = 105.78$ (2)°
 $V = 3177$ (4) Å³
 $Z = 8$

$F(000) = 1248$
 $D_x = 1.214$ Mg m⁻³
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 5659 reflections
 $\theta = 2.6$ – 23.3 °
 $\mu = 0.08$ mm⁻¹
 $T = 200$ K
 Block, colourless
 $0.25 \times 0.2 \times 0.1$ mm

Data collection

Bruker D8 Venture CMOS
 diffractometer
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Bruker, 2018)
 $T_{\min} = 0.692$, $T_{\max} = 0.745$
 39417 measured reflections

2890 independent reflections
 2007 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.086$
 $\theta_{\max} = 25.5$ °, $\theta_{\min} = 2.6$ °
 $h = -23 \rightarrow 23$
 $k = -11 \rightarrow 11$
 $l = -21 \rightarrow 21$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.105$
 $S = 1.06$
 2890 reflections
 265 parameters
 11 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
 $w = 1/[\sigma^2(F_o^2) + (0.0307P)^2 + 2.6574P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.15$ e Å⁻³
 $\Delta\rho_{\min} = -0.13$ e Å⁻³

Extinction correction: SHELXL2018
 (Sheldrick, 2015b),
 $F_c^* = kF_c [1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.0036 (5)

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.

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

	<i>x</i>	<i>y</i>	<i>z</i>	U_{iso}^*/U_{eq}	Occ. (<1)
O1	0.31856 (7)	0.49585 (15)	0.58406 (9)	0.0493 (4)	
H1	0.3549 (9)	0.553 (2)	0.5887 (14)	0.074*	
N1	0.21977 (11)	0.1682 (2)	0.71648 (12)	0.0605 (5)	
H1A	0.2128 (12)	0.107 (2)	0.7503 (11)	0.073*	
C1	0.27162 (10)	0.33085 (19)	0.65757 (11)	0.0381 (5)	
C2	0.32760 (11)	0.42144 (19)	0.65257 (12)	0.0400 (5)	
C3	0.38761 (11)	0.4290 (2)	0.71573 (13)	0.0505 (6)	
H3	0.425086	0.489562	0.712691	0.061*	
C4	0.39340 (13)	0.3476 (3)	0.78447 (14)	0.0609 (6)	
H4	0.434890	0.354722	0.826879	0.073*	
C5	0.34054 (14)	0.2583 (3)	0.79158 (14)	0.0620 (7)	
H5	0.344946	0.203862	0.837966	0.074*	
C6	0.27957 (12)	0.2504 (2)	0.72753 (13)	0.0487 (6)	
C7	0.17506 (12)	0.1943 (2)	0.64280 (13)	0.0522 (6)	
H7	0.130641	0.150538	0.622100	0.063*	
C8	0.20396 (10)	0.29280 (19)	0.60364 (12)	0.0400 (5)	
C9	0.17292 (11)	0.33949 (19)	0.51916 (12)	0.0431 (5)	
H9A	0.133346	0.276143	0.494062	0.052*	
H9B	0.209080	0.329163	0.490173	0.052*	
C10	0.14642 (10)	0.49194 (19)	0.51102 (11)	0.0392 (5)	
H10A	0.112079	0.503834	0.542269	0.047*	0.753 (7)
H10B	0.186480	0.555914	0.533343	0.047*	0.753 (7)
H10C	0.098438	0.494556	0.518071	0.047*	0.247 (7)
H10D	0.177245	0.549775	0.553315	0.047*	0.247 (7)
N2	0.1120 (2)	0.5345 (4)	0.4263 (3)	0.0424 (10)	0.753 (7)
H2	0.1020 (16)	0.6251 (14)	0.427 (2)	0.064*	0.753 (7)
C11	0.16090 (17)	0.5281 (3)	0.37193 (17)	0.0460 (11)	0.753 (7)
H11	0.174245	0.427448	0.366898	0.055*	0.753 (7)
C12	0.1208 (6)	0.5835 (11)	0.2891 (5)	0.075 (2)	0.753 (7)
H12A	0.081724	0.519686	0.265521	0.113*	0.753 (7)
H12B	0.152896	0.587641	0.255495	0.113*	0.753 (7)
H12C	0.102418	0.678046	0.294075	0.113*	0.753 (7)
C13	0.2280 (5)	0.6136 (11)	0.4063 (5)	0.0550 (16)	0.753 (7)
H13A	0.254848	0.570569	0.455889	0.082*	0.753 (7)
H13B	0.215425	0.710610	0.416231	0.082*	0.753 (7)

H13C	0.256526	0.614481	0.368717	0.082*	0.753 (7)
C14	0.0444 (4)	0.4557 (8)	0.3919 (4)	0.0602 (16)	0.753 (7)
H14A	0.012269	0.515034	0.352607	0.090*	0.753 (7)
H14B	0.022643	0.432096	0.434091	0.090*	0.753 (7)
H14C	0.054337	0.368833	0.366817	0.090*	0.753 (7)
N2A	0.1450 (7)	0.5560 (14)	0.4311 (9)	0.050 (3)	0.247 (7)
H2A	0.134 (5)	0.643 (4)	0.439 (7)	0.075*	0.247 (7)
C11A	0.1027 (5)	0.4723 (9)	0.3612 (6)	0.056 (4)	0.247 (7)
H11A	0.129490	0.383119	0.360557	0.067*	0.247 (7)
C12A	0.1043 (18)	0.553 (3)	0.2865 (15)	0.072 (9)	0.247 (7)
H12D	0.152027	0.587736	0.291618	0.108*	0.247 (7)
H12E	0.071726	0.632858	0.279034	0.108*	0.247 (7)
H12F	0.090242	0.489724	0.240840	0.108*	0.247 (7)
C13A	0.0317 (13)	0.428 (3)	0.3716 (17)	0.083 (7)	0.247 (7)
H13D	0.038819	0.359379	0.414764	0.124*	0.247 (7)
H13E	0.003480	0.384461	0.322671	0.124*	0.247 (7)
H13F	0.007061	0.510726	0.383946	0.124*	0.247 (7)
C14A	0.2162 (13)	0.601 (3)	0.4231 (16)	0.048 (5)	0.247 (7)
H14D	0.210287	0.677296	0.384573	0.072*	0.247 (7)
H14E	0.238906	0.520050	0.405164	0.072*	0.247 (7)
H14F	0.245452	0.633016	0.474299	0.072*	0.247 (7)
O2	0.41362 (7)	0.68230 (14)	0.57453 (9)	0.0518 (4)	
C15	0.45985 (11)	0.6662 (2)	0.53625 (14)	0.0489 (5)	
C16	0.47710 (10)	0.5170 (2)	0.51944 (13)	0.0438 (5)	
H16	0.453512	0.442783	0.538017	0.053*	
O3	0.4846 (16)	0.7662 (12)	0.5058 (18)	0.079 (4)	0.73 (8)
O3A	0.5040 (17)	0.762 (4)	0.537 (4)	0.066 (7)	0.27 (8)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0463 (9)	0.0440 (8)	0.0593 (9)	-0.0065 (7)	0.0175 (7)	0.0114 (7)
N1	0.0821 (14)	0.0515 (12)	0.0562 (13)	-0.0066 (11)	0.0332 (11)	0.0146 (10)
C1	0.0498 (12)	0.0287 (10)	0.0412 (11)	0.0039 (9)	0.0215 (10)	-0.0007 (8)
C2	0.0464 (12)	0.0298 (10)	0.0474 (12)	0.0064 (9)	0.0189 (10)	-0.0015 (9)
C3	0.0466 (13)	0.0467 (12)	0.0581 (14)	0.0043 (10)	0.0145 (12)	-0.0041 (11)
C4	0.0633 (15)	0.0632 (15)	0.0520 (14)	0.0116 (13)	0.0085 (12)	-0.0056 (12)
C5	0.0827 (18)	0.0618 (15)	0.0435 (14)	0.0132 (14)	0.0206 (14)	0.0069 (11)
C6	0.0628 (15)	0.0419 (12)	0.0475 (13)	0.0043 (11)	0.0251 (12)	0.0032 (10)
C7	0.0583 (14)	0.0436 (12)	0.0600 (15)	-0.0069 (10)	0.0253 (12)	0.0032 (11)
C8	0.0485 (12)	0.0290 (10)	0.0485 (12)	0.0010 (8)	0.0234 (10)	0.0018 (9)
C9	0.0504 (12)	0.0305 (10)	0.0503 (12)	-0.0006 (9)	0.0168 (10)	-0.0035 (9)
C10	0.0436 (11)	0.0336 (10)	0.0454 (12)	0.0016 (9)	0.0204 (10)	-0.0011 (9)
N2	0.048 (2)	0.0347 (16)	0.0467 (19)	0.0054 (18)	0.016 (2)	0.0011 (13)
C11	0.063 (3)	0.0365 (16)	0.0449 (19)	0.0035 (15)	0.0246 (17)	-0.0003 (13)
C12	0.102 (4)	0.074 (6)	0.049 (3)	-0.009 (4)	0.019 (3)	0.009 (3)
C13	0.051 (4)	0.063 (3)	0.059 (3)	0.001 (3)	0.028 (2)	0.010 (3)
C14	0.056 (4)	0.044 (3)	0.074 (3)	-0.016 (2)	0.007 (3)	0.003 (3)

N2A	0.060 (8)	0.034 (6)	0.056 (6)	0.022 (6)	0.017 (7)	-0.001 (5)
C11A	0.059 (8)	0.034 (5)	0.064 (7)	0.004 (5)	0.000 (5)	-0.007 (5)
C12A	0.13 (2)	0.032 (8)	0.054 (10)	0.029 (12)	0.032 (12)	0.011 (6)
C13A	0.048 (10)	0.061 (11)	0.130 (18)	-0.024 (7)	0.007 (11)	-0.035 (10)
C14A	0.049 (9)	0.039 (7)	0.074 (12)	0.011 (6)	0.046 (7)	0.010 (7)
O2	0.0507 (9)	0.0341 (7)	0.0808 (11)	0.0017 (6)	0.0353 (8)	0.0008 (7)
C15	0.0470 (12)	0.0326 (11)	0.0739 (16)	0.0036 (10)	0.0278 (12)	0.0051 (11)
C16	0.0402 (11)	0.0306 (10)	0.0651 (14)	-0.0004 (9)	0.0221 (10)	0.0069 (9)
O3	0.100 (8)	0.0310 (18)	0.137 (9)	0.008 (3)	0.083 (7)	0.017 (4)
O3A	0.057 (10)	0.031 (6)	0.124 (19)	-0.012 (6)	0.048 (11)	0.001 (9)

Geometric parameters (Å, °)

O1—H1	0.885 (10)	C11—C13	1.531 (10)
O1—C2	1.367 (2)	C12—H12A	0.9800
N1—H1A	0.870 (10)	C12—H12B	0.9800
N1—C6	1.385 (3)	C12—H12C	0.9800
N1—C7	1.380 (3)	C13—H13A	0.9800
C1—C2	1.422 (3)	C13—H13B	0.9800
C1—C6	1.421 (3)	C13—H13C	0.9800
C1—C8	1.460 (3)	C14—H14A	0.9800
C2—C3	1.390 (3)	C14—H14B	0.9800
C3—H3	0.9500	C14—H14C	0.9800
C3—C4	1.414 (3)	N2A—H2A	0.874 (11)
C4—H4	0.9500	N2A—C11A	1.511 (18)
C4—C5	1.377 (3)	N2A—C14A	1.51 (2)
C5—H5	0.9500	C11A—H11A	1.0000
C5—C6	1.412 (3)	C11A—C12A	1.53 (2)
C7—H7	0.9500	C11A—C13A	1.52 (2)
C7—C8	1.374 (3)	C12A—H12D	0.9800
C8—C9	1.514 (3)	C12A—H12E	0.9800
C9—H9A	0.9900	C12A—H12F	0.9800
C9—H9B	0.9900	C13A—H13D	0.9800
C9—C10	1.530 (3)	C13A—H13E	0.9800
C10—H10A	0.9900	C13A—H13F	0.9800
C10—H10B	0.9900	C14A—H14D	0.9800
C10—H10C	0.9900	C14A—H14E	0.9800
C10—H10D	0.9900	C14A—H14F	0.9800
C10—N2	1.518 (5)	O2—C15	1.284 (2)
C10—N2A	1.527 (15)	C15—C16	1.503 (3)
N2—H2	0.882 (10)	C15—O3	1.251 (10)
N2—C11	1.536 (5)	C15—O3A	1.26 (3)
N2—C14	1.506 (7)	C16—C16 ⁱ	1.315 (4)
C11—H11	1.0000	C16—H16	0.9500
C11—C12	1.550 (9)		
C2—O1—H1	109.0 (16)	C13—C11—C12	111.0 (5)
C6—N1—H1A	124.6 (17)	C11—C12—H12A	109.5

C7—N1—H1A	125.8 (17)	C11—C12—H12B	109.5
C7—N1—C6	109.58 (18)	C11—C12—H12C	109.5
C2—C1—C8	134.38 (18)	H12A—C12—H12B	109.5
C6—C1—C2	118.24 (19)	H12A—C12—H12C	109.5
C6—C1—C8	107.32 (18)	H12B—C12—H12C	109.5
O1—C2—C1	116.68 (18)	C11—C13—H13A	109.5
O1—C2—C3	123.93 (19)	C11—C13—H13B	109.5
C3—C2—C1	119.38 (19)	C11—C13—H13C	109.5
C2—C3—H3	119.7	H13A—C13—H13B	109.5
C2—C3—C4	120.7 (2)	H13A—C13—H13C	109.5
C4—C3—H3	119.7	H13B—C13—H13C	109.5
C3—C4—H4	119.1	N2—C14—H14A	109.5
C5—C4—C3	121.8 (2)	N2—C14—H14B	109.5
C5—C4—H4	119.1	N2—C14—H14C	109.5
C4—C5—H5	121.2	H14A—C14—H14B	109.5
C4—C5—C6	117.6 (2)	H14A—C14—H14C	109.5
C6—C5—H5	121.2	H14B—C14—H14C	109.5
N1—C6—C1	106.9 (2)	C10—N2A—H2A	100 (8)
N1—C6—C5	130.8 (2)	C11A—N2A—C10	114.2 (10)
C5—C6—C1	122.3 (2)	C11A—N2A—H2A	122 (8)
N1—C7—H7	124.8	C11A—N2A—C14A	113.2 (16)
C8—C7—N1	110.3 (2)	C14A—N2A—C10	114.4 (13)
C8—C7—H7	124.8	C14A—N2A—H2A	92 (7)
C1—C8—C9	128.51 (17)	N2A—C11A—H11A	106.0
C7—C8—C1	105.86 (18)	N2A—C11A—C12A	107.6 (15)
C7—C8—C9	125.42 (19)	N2A—C11A—C13A	111.8 (12)
C8—C9—H9A	108.8	C12A—C11A—H11A	106.0
C8—C9—H9B	108.8	C13A—C11A—H11A	106.0
C8—C9—C10	113.90 (16)	C13A—C11A—C12A	118.5 (19)
H9A—C9—H9B	107.7	C11A—C12A—H12D	109.5
C10—C9—H9A	108.8	C11A—C12A—H12E	109.5
C10—C9—H9B	108.8	C11A—C12A—H12F	109.5
C9—C10—H10A	108.9	H12D—C12A—H12E	109.5
C9—C10—H10B	108.9	H12D—C12A—H12F	109.5
C9—C10—H10C	109.1	H12E—C12A—H12F	109.5
C9—C10—H10D	109.1	C11A—C13A—H13D	109.5
H10A—C10—H10B	107.8	C11A—C13A—H13E	109.5
H10C—C10—H10D	107.8	C11A—C13A—H13F	109.5
N2—C10—C9	113.2 (2)	H13D—C13A—H13E	109.5
N2—C10—H10A	108.9	H13D—C13A—H13F	109.5
N2—C10—H10B	108.9	H13E—C13A—H13F	109.5
N2A—C10—C9	112.5 (5)	N2A—C14A—H14D	109.5
N2A—C10—H10C	109.1	N2A—C14A—H14E	109.5
N2A—C10—H10D	109.1	N2A—C14A—H14F	109.5
C10—N2—H2	106 (2)	H14D—C14A—H14E	109.5
C10—N2—C11	114.3 (3)	H14D—C14A—H14F	109.5
C11—N2—H2	104 (2)	H14E—C14A—H14F	109.5
C14—N2—C10	112.0 (4)	O2—C15—C16	116.58 (17)

C14—N2—H2	108 (2)	O3—C15—O2	123.5 (5)
C14—N2—C11	111.7 (4)	O3—C15—C16	119.5 (5)
N2—C11—H11	108.6	O3A—C15—O2	120.2 (17)
N2—C11—C12	109.0 (5)	O3A—C15—C16	119.1 (14)
C12—C11—H11	108.6	C15—C16—H16	118.0
C13—C11—N2	110.9 (4)	C16 ⁱ —C16—C15	123.9 (2)
C13—C11—H11	108.6	C16 ⁱ —C16—H16	118.0
O1—C2—C3—C4	-179.34 (19)	C8—C1—C2—O1	2.2 (3)
N1—C7—C8—C1	-0.3 (2)	C8—C1—C2—C3	-177.32 (19)
N1—C7—C8—C9	174.91 (18)	C8—C1—C6—N1	-0.1 (2)
C1—C2—C3—C4	0.1 (3)	C8—C1—C6—C5	178.10 (19)
C1—C8—C9—C10	-74.2 (2)	C8—C9—C10—N2	-176.9 (2)
C2—C1—C6—N1	-177.84 (17)	C8—C9—C10—N2A	155.6 (6)
C2—C1—C6—C5	0.4 (3)	C9—C10—N2—C11	-61.5 (3)
C2—C1—C8—C7	177.5 (2)	C9—C10—N2—C14	66.9 (5)
C2—C1—C8—C9	2.4 (3)	C9—C10—N2A—C11A	55.6 (9)
C2—C3—C4—C5	0.1 (3)	C9—C10—N2A—C14A	-77.0 (14)
C3—C4—C5—C6	-0.1 (3)	C10—N2—C11—C12	-176.2 (5)
C4—C5—C6—N1	177.6 (2)	C10—N2—C11—C13	-53.7 (5)
C4—C5—C6—C1	-0.1 (3)	C10—N2A—C11A—C12A	178.0 (14)
C6—N1—C7—C8	0.3 (3)	C10—N2A—C11A—C13A	46.2 (17)
C6—C1—C2—O1	179.16 (17)	C14—N2—C11—C12	55.2 (6)
C6—C1—C2—C3	-0.4 (3)	C14—N2—C11—C13	177.7 (6)
C6—C1—C8—C7	0.3 (2)	C14A—N2A—C11A—C12A	-49 (2)
C6—C1—C8—C9	-174.77 (18)	C14A—N2A—C11A—C13A	179.3 (19)
C7—N1—C6—C1	-0.1 (2)	O2—C15—C16—C16 ⁱ	179.6 (3)
C7—N1—C6—C5	-178.1 (2)	O3—C15—C16—C16 ⁱ	7 (2)
C7—C8—C9—C10	111.7 (2)	O3A—C15—C16—C16 ⁱ	-23 (4)

Symmetry code: (i) $-x+1, -y+1, -z+1$.

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

*Cg*2 is the centroid of the C1—C6 ring.

<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>
O1—H1 \cdots O2	0.89 (1)	1.75 (1)	2.618 (2)	165 (2)
N2—H2 \cdots O2 ⁱⁱ	0.88 (1)	1.85 (1)	2.730 (5)	175 (3)
N2A—H2A \cdots O2 ⁱⁱ	0.87 (1)	1.89 (4)	2.727 (12)	160 (11)
N1—H1A \cdots <i>Cg</i> 2 ⁱⁱⁱ	0.87 (1)	2.78 (2)	3.552 (3)	148 (2)

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