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# The fumarate salts of the *N*-isopropyl-*N*-methyl derivatives of DMT and psilocin

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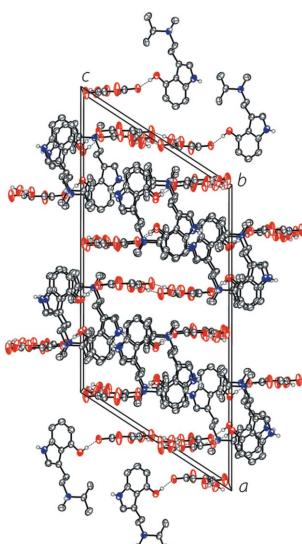
The solid-state structures of the salts of two substituted tryptamines, namely *N*-isopropyl-*N*-methyltryptaminium (MiPT) fumarate [systematic name: [2-(1*H*-indol-3-yl)ethyl](methyl)propan-2-ylazanium 3-carboxyprop-2-enoate],  $C_{14}H_{21}N_2^+ \cdot C_4H_3O_4^-$ , and 4-hydroxy-*N*-isopropyl-*N*-methyltryptaminium (4-HO-MiPT) fumarate monohydrate [systematic name: [2-(4-hydroxy-1*H*-indol-3-yl)ethyl](methyl)propan-2-ylazanium 3-carboxyprop-2-enoate monohydrate],  $C_{14}H_{21}N_2O^+ \cdot C_4H_3O_4^- \cdot H_2O$ , are reported. Both salts possess a protonated trypt ammonium cation and a 3-carboxyacrylate (hydrogen fumarate) anion in the asymmetric unit; the 4-HO-MiPT structure also contains a water molecule of crystallization. Both cations feature disorder of the side chain over two orientations, in a 0.630 (3):0.370 (3) ratio for MiPT and a 0.775 (5):0.225 (5) ratio for 4-HO-MiPT. In both extended structures, N—H···O and O—H···O hydrogen bonds generate infinite two-dimensional networks.

## 1. Chemical context

*N,N*-dimethyltryptamine (DMT) and its derivatives have been used by humans for centuries because of their psychoactive, entheogenic, or hallucinogenic effects, or combinations thereof (Cameron & Olson, 2018). Psilocybin, the 4-phosphate variant of DMT, is arguably its most studied derivative. Psilocybin is one of several naturally occurring psychoactive tryptamines found in ‘magic’ mushrooms. When consumed by humans, psilocybin serves as a prodrug of psilocin. Upon digestion, psilocybin hydrolyses to generate psilocin, the 4-hydroxy derivative of DMT. Psilocin is a potent serotonin 2a-agonist, which is responsible for its psychoactive properties (Dinis-Oliveira, 2017; Nichols, 2012). Psychoactive tryptamines like DMT and psilocin have garnered significant interest recently because of their potential for treating mood disorders, including depression, anxiety, addiction, and post-traumatic stress disorder (PTSD) (Johnson & Griffiths, 2017; Carhart-Harris & Goodwin, 2017).

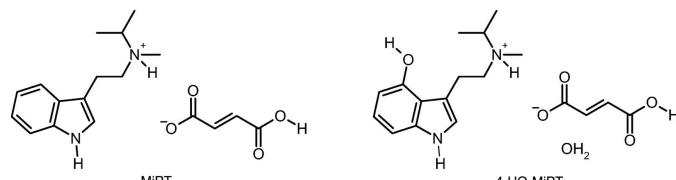
Altering the chemical structure within this class of compounds can dramatically influence the potency and action of the drugs. For example, merely changing the *N,N*-dialkyl groups on DMT can modify its psychoactive properties: increasing the chain length of the two alkyl groups of the tryptamine to larger than *n*-butyl dramatically reduces or eliminates the psychoactive effects (Bradley & Johnston, 1970).

The synthesis of *N*-methyl-*N*-isopropyltryptamine (MiPT) was reported in 1981 (Repke *et al.*, 1981). In 1985, Repke and co-workers reported that of the compounds in the series of



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*N,N*-dialkyl-4-hydroxytryptamines, the *N*-methyl-*N*-isopropyl derivative (4-HO-MiPT) is the most potent based upon qualitative effects on humans (Repke *et al.*, 1985). Later quantitative studies showed the *N*-methyl-*N*-isopropyl derivatives of DMT and psilocin to be more potent as serotonin-1A, -2A and -2B receptors compared to the analogous dimethyl compounds (McKenna *et al.*, 1990).



Improving our understanding of how these drugs interact with particular biological receptors requires a complete understanding of their chemical structures. Given their therapeutic potential and the significant structure–activity relationship between them, further studies would benefit from better understanding of their chemical structures. Responding to this unmet need, we report the crystal structures of the fumarate salts of MiPT and 4-HO-MiPT herein.

## 2. Structural commentary

The molecular structure of MiPT fumarate is shown on the left of Fig. 1. The asymmetric unit contains one *N*-methyl-*N*-isopropyltryptammonium ( $C_{14}H_{21}N_2^+$ ) cation and one 3-carboxyacrylate ( $C_4H_3O_4^-$ ) anion. The indole ring system of the cation is near planar with an r.m.s. deviation from planarity of 0.006 Å. The singly protonated fumarate anion is in the *trans* configuration and is slightly distorted from planarity with an r.m.s. deviation of 0.133 Å and a carboxylate twist angle of 18.370 (5)°. The *N*-methyl-*N*-isopropylammonium group is disordered over two orientations in a 0.630 (3):0.370 (3) ratio.

The molecular structure of 4-HO-MiPT fumarate monohydrate is shown on the right of Fig. 1. The asymmetric unit contains one 4-hydroxy-*N*-methyl-*N*-isopropyltryptammonium cation, one 4-hydroxy-3-carboxyacrylate anion, and one water molecule of crystallization.

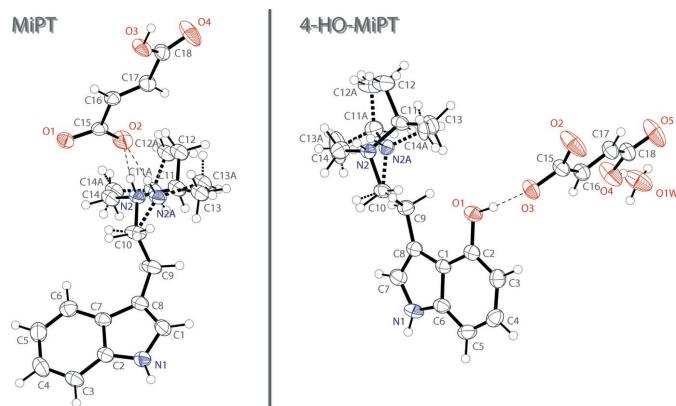


Figure 1

The molecular structure of MiPT fumarate (left) and HO-MiPT fumarate monohydrate (right), showing the atomic labeling. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen bonds are shown as dashed lines.

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

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O3—H3A $\cdots$ O1 <sup>i</sup>	0.87 (1)	1.66 (1)	2.5316 (18)	176 (3)
N1—H1 $\cdots$ O1 <sup>ii</sup>	0.87 (1)	2.04 (1)	2.874 (2)	160 (2)
N2—H2 $\cdots$ O2	0.88 (1)	1.79 (1)	2.667 (3)	173 (3)
N2A—H2A $\cdots$ O2	0.88 (1)	1.81 (2)	2.670 (5)	167 (6)

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

nium ( $C_{14}H_{21}N_2O^+$ ) cation, one 3-carboxyacrylate anion and one water molecule of crystallization. The indole ring system of the cation is close to planar with an r.m.s. deviation of 0.021 Å. The singly protonated fumarate anion is also near planar with an r.m.s. deviation of 0.049 Å. The *N*-methyl-*N*-isopropylammonium group shows a similar disorder to the MiPT structure over two orientations in a 0.775 (5):0.225 (5) ratio.

## 3. Supramolecular features

In the extended structure of MiPT fumarate, the *N*-methyl-*N*-isopropylamine and fumarate ions are linked into infinite two-dimensional networks lying parallel to the (010) plane through  $N-H \cdots O$  and  $O-H \cdots O$  hydrogen bonds (Table 1). The proton of the ammonium cation forms a hydrogen bond with one of the oxygen atoms of the deprotonated  $-CO_2^-$  group of the 3-carboxyacrylate ion. The carboxylic acid proton forms a hydrogen bond with an oxygen atom of an adjacent 3-carboxyacrylate anion. The  $N-H$  grouping of the indole ring also forms hydrogen bonds to one of the oxygen atoms of the 3-carboxyacrylate anion. The hydrogen bonding is shown on the left in Fig. 2, and the packing of MiPT fumarate is shown on the left in Fig. 3.

In the structure of 4-HO-MiPT fumarate, there are  $N-H \cdots O$  and  $O-H \cdots O$  hydrogen bonds that link together the cations and anions as well as the water molecules of crystallization (Table 2). The result is a two-dimensional network lying parallel to the (201) plane. The proton of the ammonium cation forms a bifurcated  $N-H \cdots (O,O)$  hydrogen bond with

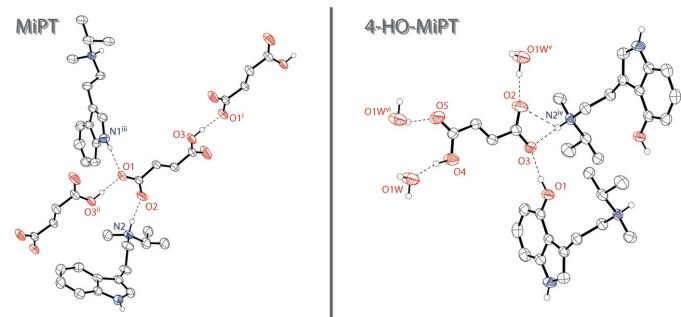


Figure 2

The hydrogen bonding of the fumarate ion in the structure of MiPT (left) and HO-MiPT (right). Displacement ellipsoids are drawn at the 50% probability level. Hydrogen atoms not involved in hydrogen bonds are omitted for clarity. Only one component of the amine disorder is shown. Symmetry codes: (i)  $x, \frac{3}{2} - y, -\frac{1}{2} + z$  (ii)  $x, \frac{3}{2} - y, \frac{1}{2} + z$  (iii)  $-1 + x, \frac{3}{2} - y, -\frac{1}{2} + z$  (iv)  $\frac{3}{2} - x, \frac{1}{2} - y, 1 - z$  (v)  $x, -1 + y, z$  (vi)  $1 - x, 2 - y, -z$ .

**Table 2**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ) for 4-HO-MiPT.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N2—H2 $\cdots$ O2 <sup>i</sup>	0.88 (1)	2.51 (2)	3.085 (2)	124 (2)
N2—H2 $\cdots$ O3 <sup>i</sup>	0.88 (1)	1.89 (1)	2.775 (2)	178 (2)
N2A—H2A $\cdots$ O3 <sup>i</sup>	0.87 (1)	1.85 (2)	2.717 (6)	172 (8)
O1—H1 $\cdots$ O3	0.87 (2)	1.79 (3)	2.6512 (17)	172 (2)
O4—H4A $\cdots$ O1W	0.93 (3)	1.66 (3)	2.579 (2)	167 (3)
O1W—H1WA $\cdots$ O5 <sup>ii</sup>	0.84 (3)	1.98 (3)	2.779 (2)	160 (2)
O1W—H1WB $\cdots$ O2 <sup>iii</sup>	0.87 (3)	1.74 (3)	2.599 (2)	170 (2)

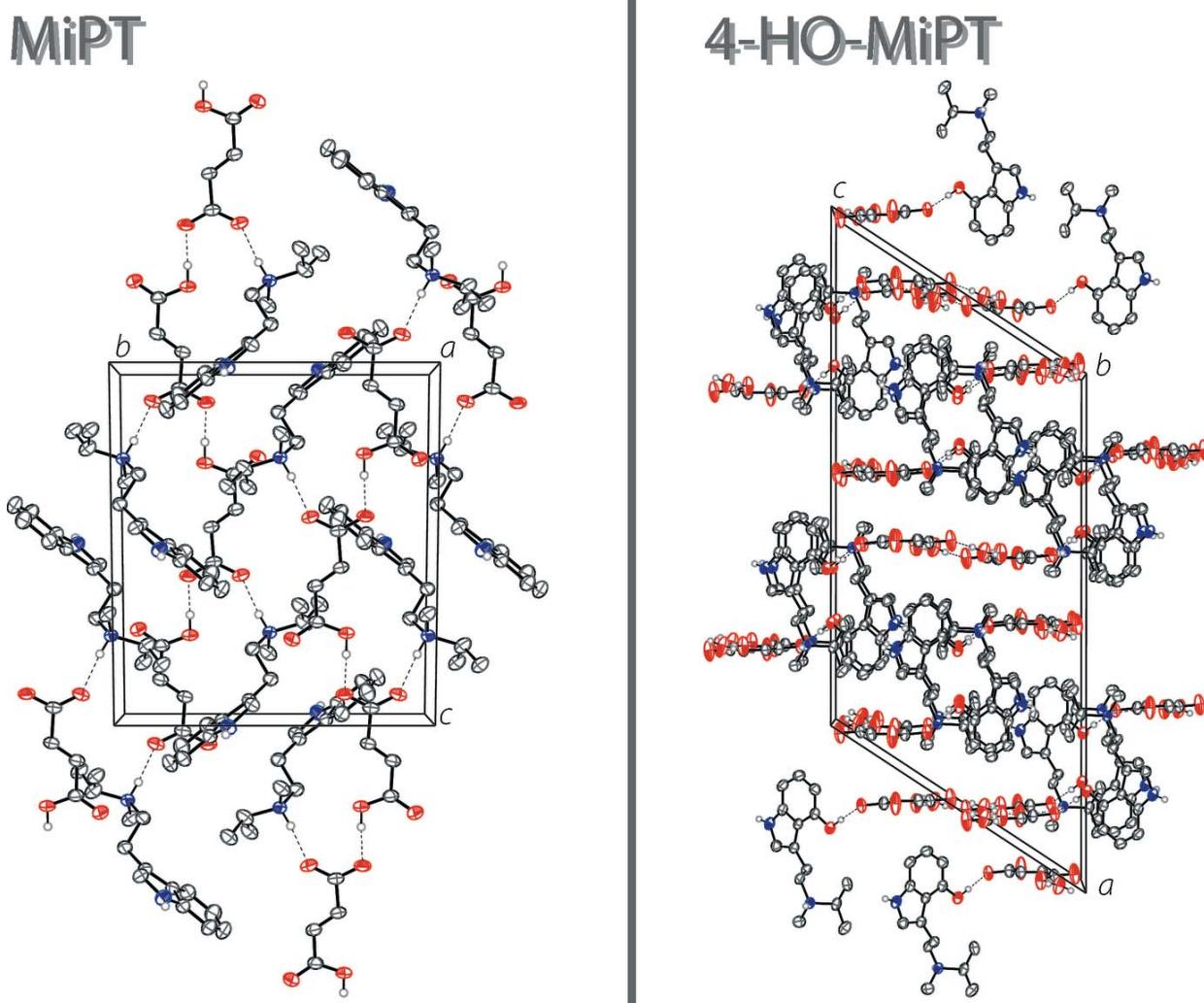
Symmetry codes: (i)  $-x + \frac{3}{2}, -y + \frac{1}{2}, -z + 1$ ; (ii)  $-x + 1, -y + 2, -z$ ; (iii)  $x, y + 1, z$ .

the deprotonated  $-\text{CO}_2^-$  group of the 3-carboxyacrylate ion. The hydrogen of the hydroxy group also hydrogen bonds to the same oxygen atom of the anion. The carboxylic acid proton hydrogen bonds with a water molecule in the structure. Two other water molecules form hydrogen bonds with two different oxygen atoms of the anion. The hydrogen bonding is

shown on the right in Fig. 2, and the packing of 4-HO-MiPT fumarate is shown on the right in Fig. 3.

#### 4. Database survey

The MiPT structure described above is a derivative of DMT (*N,N*-dimethyltryptamine), which has been structurally characterized (Falkenberg, 1972), as well as its close derivative MPT, *N*-methyl-*N*-propyltryptamine (Chadeayne *et al.* 2019*b*). In both cases, these were crystallized as free bases, while MiPT is the fumarate salt. In the case of 4-HO-MiPT, the most closely related molecule is psilocin, which has been structurally characterized (Petcher & Weber, 1974), as well as psilocybin (Weber & Petcher, 1974). Psilocin was reported as the free base and psilocybin was reported as a zwitterionic molecule, while the structure of 4-HO-MiPT reported here is the hydrated fumarate salt. Two different ionic structures of the 4-acetoxy derivative of DMT have been reported as



**Figure 3**

The crystal packing of MiPT fumarate (left), viewed along the  $a$  axis, and the crystal packing of HO-MiPT fumarate (right), viewed along the  $b$  axis. The hydrogen bonds (Tables 1 and 2) are shown as dashed lines. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen atoms not involved in hydrogen bonds are omitted for clarity. Only one component of the amine disorder is shown.

**Table 3**  
Experimental details.

	MiPT	4-HO-MiPT
Crystal data		
Chemical formula	$C_{14}H_{21}N_2^+ \cdot C_4H_3O_4^-$	$C_{14}H_{21}N_2O^+ \cdot C_4H_3O_4^- \cdot H_2O$
$M_r$	332.39	366.41
Crystal system, space group	Monoclinic, $P2_1/c$	Monoclinic, $C2/c$
Temperature (K)	200	200
$a, b, c$ (Å)	9.852 (2), 12.789 (2), 14.875 (3)	29.507 (3), 8.7445 (8), 17.3659 (18)
$\beta$ (°)	106.932 (7)	123.389 (3)
$V$ (Å <sup>3</sup> )	1793.0 (6)	3741.2 (7)
$Z$	4	8
Radiation type	Mo $K\alpha$	Mo $K\alpha$
$\mu$ (mm <sup>-1</sup> )	0.09	0.10
Crystal size (mm)	0.20 × 0.18 × 0.05	0.30 × 0.25 × 0.20
Data collection		
Diffractometer	Bruker D8 Venture CMOS	Bruker D8 Venture CMOS
Absorption correction	Multi-scan ( <i>SADABS</i> ; Bruker, 2016)	Multi-scan ( <i>SADABS</i> ; Bruker, 2016)
$T_{min}, T_{max}$	0.687, 0.745	0.719, 0.745
No. of measured, independent and observed [ $I > 2\sigma(I)$ ] reflections	36899, 3297, 2605	70395, 3458, 2978
$R_{int}$	0.052	0.041
(sin $\theta/\lambda$ ) <sub>max</sub> (Å <sup>-1</sup> )	0.604	0.604
Refinement		
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.049, 0.127, 1.06	0.041, 0.096, 1.08
No. of reflections	3297	3458
No. of parameters	240	320
No. of restraints	8	12
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å <sup>-3</sup> )	0.26, -0.26	0.22, -0.20

Computer programs: *APEX3* and *SAINT* (Bruker, 2016), *SHELXS97* and *SHELXL97* (Sheldrick, 2008) and *OLEX2* (Dolomanov *et al.*, 2009).

fumarate salts (Chadeayne *et al.* 2019a,c). The metrical parameters of the tryptammonium cations for MiPT and 4-HO-MiPT are consistent with those of the other tryptammonium structures reported.

## 5. Synthesis and crystallization

Single crystals suitable for X-ray analysis were obtained from the slow evaporation of aqueous solutions of commercial samples of *N*-methyl-*N*-isopropyltryptammonium fumarate and 4-hydroxy-*N*-methyl-*N*-isopropyltryptammonium fumarate (The Indole Shop).

## 6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. C-bound H atoms were placed in calculated positions (C—H = 0.95–1.00 Å) and refined as riding with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{C-methyl})$ . The following restraints were applied: C—N = 1.54±0.01, N—H = 0.87±0.01, O—H = 0.86±0.01 Å. N- and O-bound H atoms were refined with  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{N,O})$ . In the MiPT fumarate, the *N*-methyl-*N*-isopropylaminium group is disordered. It is modeled as two components: N2 and C11–C14 with an occupancy of 0.630 (3) and N2A and C11A–C14A with an occupancy of 0.370 (3). 4-HO MiPT fumarate exhibits a similar disorder of the *N*-methyl-*N*-isopropylaminium group that is modeled as two components: N2, C11–C14 with an

occupancy of 0.775 (5) and N2A, C11A–C14A with an occupancy of 0.225 (5).

## Acknowledgements

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## References

- Bradley, R. J. & Johnston, V. S. (1970). *Origin and Mechanism of Hallucinations*, edited by W. Keup, pp. 333–344. New York: Plenum Press.
- Bruker (2016). *APEX3*, *SAINT*, and *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Cameron, L. P. & Olson, D. E. (2018). *ACS Chem. Neurosci.* **9**, 2344–2357.
- Carhart-Harris, R. L. & Goodwin, G. M. (2017). *Neuropsychopharmacology*, **42**, 2105–2113.
- Chadeayne, A. R., Golen, J. A. & Manke, D. R. (2019a). *Acta Cryst. E* **75**, 900–902.
- Chadeayne, A. R., Golen, J. A. & Manke, D. R. (2019b). *IUCrData*, **4**, x190962.

- Chadeayne, A. R., Golen, J. A. & Manke, D. R. (2019c). *Psychedelic Science Review*, <https://psychedelicreview.com/the-crystal-structure-of-4-aco-dmt-fumarate/>
- Dinis-Oliveira, R. J. (2017). *Drug Metab. Rev.* **49**, 84–91.
- Dolomanov, O. V., Bourhis, L. J., Gildea, R. J., Howard, J. A. K. & Puschmann, H. (2009). *J. Appl. Cryst.* **42**, 339–341.
- Falkenberg, G. (1972). *Acta Cryst. B* **28**, 3075–3083.
- Johnson, M. W. & Griffiths, R. R. (2017). *Neurotherapeutics* **14**, 734–740.
- McKenna, D. J., Repke, D. B., Lo, L. & Peroutka, S. J. (1990). *Neuropharmacology*, **29**, 193–198.
- Nichols, D. E. (2012). *WIREs Membr. Transp. Signal.* **1**, 559–579.
- Petcher, T. J. & Weber, H. P. (1974). *J. Chem. Soc. Perkin Trans. 2*, pp. 946–948.
- Repke, D. B., Ferguson, W. J. & Bates, D. K. (1981). *J. Heterocycl. Chem.* **18**, 175–179.
- Repke, D. B., Grotjahn, D. B. & Shulgin, A. T. (1985). *J. Med. Chem.* **28**, 892–896.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Weber, H. P. & Petcher, T. J. (1974). *J. Chem. Soc. Perkin Trans. 2*, pp. 942–946.

# supporting information

*Acta Cryst.* (2019). E75, 1316-1320 [https://doi.org/10.1107/S2056989019011253]

## The fumarate salts of the *N*-isopropyl-*N*-methyl derivatives of DMT and psilocin

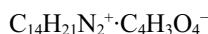
Andrew R. Chadeayne, Duyen N. K. Pham, James A. Golen and David R. Manke

### Computing details

For both structures, data collection: *APEX3* (Bruker, 2016); cell refinement: *SAINT* (Bruker, 2016); data reduction: *SAINT* (Bruker, 2016); program(s) used to solve structure: *SHELXS97* (Sheldrick 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2* (Dolomanov *et al.*, 2009).

### [2-(1*H*-Indol-3-yl)ethyl](methyl)propan-2-ylazanium 3-carboxyprop-2-enoate (MiPT)

#### Crystal data



$$M_r = 332.39$$

Monoclinic,  $P2_1/c$

$$a = 9.852 (2) \text{ \AA}$$

$$b = 12.789 (2) \text{ \AA}$$

$$c = 14.875 (3) \text{ \AA}$$

$$\beta = 106.932 (7)^\circ$$

$$V = 1793.0 (6) \text{ \AA}^3$$

$$Z = 4$$

$$F(000) = 712$$

$$D_x = 1.231 \text{ Mg m}^{-3}$$

$$\text{Mo } K\alpha \text{ radiation, } \lambda = 0.71073 \text{ \AA}$$

Cell parameters from 9752 reflections

$$\theta = 3.0\text{--}25.3^\circ$$

$$\mu = 0.09 \text{ mm}^{-1}$$

$$T = 200 \text{ K}$$

BLOCK, colourless

$$0.20 \times 0.18 \times 0.05 \text{ mm}$$

#### Data collection

Bruker D8 Venture CMOS

diffractometer

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan  
(SADABS; Bruker, 2016)

$$T_{\min} = 0.687, T_{\max} = 0.745$$

36899 measured reflections

3297 independent reflections

2605 reflections with  $I > 2\sigma(I)$

$$R_{\text{int}} = 0.052$$

$$\theta_{\max} = 25.4^\circ, \theta_{\min} = 3.0^\circ$$

$$h = -11 \rightarrow 11$$

$$k = -15 \rightarrow 15$$

$$l = -17 \rightarrow 17$$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.049$$

$$wR(F^2) = 0.127$$

$$S = 1.06$$

3297 reflections

240 parameters

8 restraints

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0488P)^2 + 1.0971P] \quad \text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.26 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.25 \text{ e \AA}^{-3}$$

Extinction correction: SHELXL,

$$Fc^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$$

Extinction coefficient: 0.039 (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.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1	0.17442 (15)	0.78060 (11)	0.58471 (8)	0.0395 (4)	
O2	0.21577 (18)	0.61007 (12)	0.58250 (9)	0.0521 (4)	
O3	0.19484 (17)	0.72434 (11)	0.25829 (9)	0.0442 (4)	
H3A	0.184 (3)	0.724 (2)	0.1982 (8)	0.066*	
O4	0.1592 (2)	0.55393 (12)	0.24156 (11)	0.0731 (6)	
N1	0.88917 (17)	0.64452 (13)	1.00719 (12)	0.0382 (4)	
H1	0.9809 (11)	0.6526 (17)	1.0259 (15)	0.046*	
N2	0.3185 (3)	0.5163 (2)	0.74864 (18)	0.0303 (7)	0.630 (3)
H2	0.281 (3)	0.551 (2)	0.6962 (14)	0.036*	0.630 (3)
C11	0.3410 (4)	0.4055 (3)	0.7190 (2)	0.0443 (4)	0.630 (3)
H11	0.3829	0.3640	0.7775	0.053*	0.630 (3)
C12	0.2020 (4)	0.3543 (3)	0.6666 (3)	0.0443 (4)	0.630 (3)
H12A	0.2210	0.2905	0.6359	0.066*	0.630 (3)
H12B	0.1459	0.4027	0.6191	0.066*	0.630 (3)
H12C	0.1490	0.3365	0.7110	0.066*	0.630 (3)
C13	0.4553 (5)	0.4179 (4)	0.6694 (3)	0.0443 (4)	0.630 (3)
H13A	0.5401	0.4493	0.7127	0.066*	0.630 (3)
H13B	0.4200	0.4631	0.6143	0.066*	0.630 (3)
H13C	0.4794	0.3491	0.6493	0.066*	0.630 (3)
C14	0.2176 (6)	0.5221 (4)	0.8107 (4)	0.0443 (4)	0.630 (3)
H14A	0.2112	0.5946	0.8304	0.066*	0.630 (3)
H14B	0.2545	0.4779	0.8664	0.066*	0.630 (3)
H14C	0.1232	0.4974	0.7749	0.066*	0.630 (3)
N2A	0.3757 (5)	0.4914 (4)	0.7207 (3)	0.0342 (11)	0.370 (3)
H2A	0.336 (6)	0.533 (4)	0.674 (3)	0.041*	0.370 (3)
C11A	0.2757 (6)	0.4302 (5)	0.7616 (4)	0.0443 (4)	0.370 (3)
H11A	0.3302	0.3822	0.8124	0.053*	0.370 (3)
C12A	0.1685 (8)	0.3698 (6)	0.6867 (5)	0.0443 (4)	0.370 (3)
H12D	0.2131	0.3066	0.6710	0.066*	0.370 (3)
H12E	0.1341	0.4134	0.6305	0.066*	0.370 (3)
H12F	0.0887	0.3500	0.7099	0.066*	0.370 (3)
C13A	0.4250 (8)	0.4028 (6)	0.6507 (5)	0.0443 (4)	0.370 (3)
H13D	0.5166	0.4234	0.6426	0.066*	0.370 (3)
H13E	0.3532	0.4006	0.5892	0.066*	0.370 (3)
H13F	0.4336	0.3335	0.6800	0.066*	0.370 (3)
C14A	0.1952 (10)	0.5037 (7)	0.7980 (7)	0.0443 (4)	0.370 (3)
H14D	0.1334	0.4659	0.8279	0.066*	0.370 (3)
H14E	0.1373	0.5467	0.7465	0.066*	0.370 (3)
H14F	0.2601	0.5487	0.8445	0.066*	0.370 (3)

C1	0.8135 (2)	0.57049 (15)	0.94651 (13)	0.0380 (5)	
H1A	0.8536	0.5193	0.9156	0.046*	
C2	0.7970 (2)	0.70501 (14)	1.03799 (13)	0.0333 (4)	
C3	0.8232 (2)	0.78851 (15)	1.10072 (14)	0.0411 (5)	
H3	0.9169	0.8130	1.1293	0.049*	
C4	0.7083 (2)	0.83418 (16)	1.11972 (16)	0.0475 (5)	
H4	0.7230	0.8911	1.1625	0.057*	
C5	0.5702 (2)	0.79874 (17)	1.07743 (16)	0.0468 (5)	
H5	0.4929	0.8323	1.0916	0.056*	
C6	0.5444 (2)	0.71618 (15)	1.01564 (14)	0.0392 (5)	
H6	0.4501	0.6926	0.9874	0.047*	
C7	0.6584 (2)	0.66727 (14)	0.99481 (12)	0.0317 (4)	
C8	0.6725 (2)	0.58089 (14)	0.93698 (12)	0.0343 (4)	
C9	0.5561 (2)	0.51227 (15)	0.87944 (13)	0.0410 (5)	
H9A	0.4972	0.4877	0.9190	0.049*	
H9B	0.5982	0.4502	0.8581	0.049*	
C10	0.4622 (2)	0.56961 (15)	0.79425 (13)	0.0413 (5)	
H10A	0.5141	0.5759	0.7466	0.050*	0.630 (3)
H10B	0.4446	0.6412	0.8135	0.050*	0.630 (3)
H10C	0.5221	0.6129	0.7657	0.050*	0.370 (3)
H10D	0.3967	0.6167	0.8143	0.050*	0.370 (3)
C15	0.19062 (19)	0.69693 (15)	0.54455 (12)	0.0311 (4)	
C16	0.17850 (19)	0.70575 (15)	0.44231 (12)	0.0314 (4)	
H16	0.1595	0.7725	0.4133	0.038*	
C17	0.1928 (2)	0.62619 (16)	0.39115 (13)	0.0379 (5)	
H17	0.2124	0.5600	0.4212	0.045*	
C18	0.1807 (2)	0.63118 (15)	0.28967 (13)	0.0366 (5)	

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0506 (8)	0.0474 (8)	0.0207 (6)	0.0070 (6)	0.0106 (6)	-0.0012 (6)
O2	0.0820 (12)	0.0477 (9)	0.0241 (7)	0.0098 (8)	0.0117 (7)	0.0083 (6)
O3	0.0714 (10)	0.0420 (8)	0.0197 (7)	-0.0031 (7)	0.0143 (7)	-0.0007 (6)
O4	0.1495 (19)	0.0411 (9)	0.0324 (8)	-0.0018 (10)	0.0323 (10)	-0.0061 (7)
N1	0.0326 (9)	0.0421 (9)	0.0379 (9)	-0.0025 (7)	0.0068 (7)	-0.0066 (7)
N2	0.0357 (16)	0.0298 (14)	0.0220 (14)	-0.0031 (12)	0.0034 (12)	-0.0004 (11)
C11	0.0445 (12)	0.0460 (10)	0.0386 (10)	-0.0045 (7)	0.0064 (7)	-0.0135 (8)
C12	0.0445 (12)	0.0460 (10)	0.0386 (10)	-0.0045 (7)	0.0064 (7)	-0.0135 (8)
C13	0.0445 (12)	0.0460 (10)	0.0386 (10)	-0.0045 (7)	0.0064 (7)	-0.0135 (8)
C14	0.0445 (12)	0.0460 (10)	0.0386 (10)	-0.0045 (7)	0.0064 (7)	-0.0135 (8)
N2A	0.040 (3)	0.033 (3)	0.025 (2)	-0.002 (2)	0.002 (2)	0.000 (2)
C11A	0.0445 (12)	0.0460 (10)	0.0386 (10)	-0.0045 (7)	0.0064 (7)	-0.0135 (8)
C12A	0.0445 (12)	0.0460 (10)	0.0386 (10)	-0.0045 (7)	0.0064 (7)	-0.0135 (8)
C13A	0.0445 (12)	0.0460 (10)	0.0386 (10)	-0.0045 (7)	0.0064 (7)	-0.0135 (8)
C14A	0.0445 (12)	0.0460 (10)	0.0386 (10)	-0.0045 (7)	0.0064 (7)	-0.0135 (8)
C1	0.0453 (12)	0.0365 (10)	0.0305 (10)	0.0001 (9)	0.0084 (9)	-0.0056 (8)
C2	0.0369 (10)	0.0314 (9)	0.0296 (9)	-0.0035 (8)	0.0066 (8)	-0.0005 (7)

C3	0.0431 (11)	0.0376 (11)	0.0385 (11)	-0.0104 (9)	0.0055 (9)	-0.0081 (9)
C4	0.0602 (14)	0.0360 (11)	0.0467 (12)	-0.0029 (10)	0.0159 (11)	-0.0117 (9)
C5	0.0477 (13)	0.0433 (12)	0.0514 (13)	0.0043 (10)	0.0175 (10)	-0.0052 (10)
C6	0.0346 (10)	0.0393 (11)	0.0409 (11)	-0.0028 (8)	0.0069 (9)	0.0015 (9)
C7	0.0360 (10)	0.0288 (9)	0.0264 (9)	-0.0031 (7)	0.0029 (8)	0.0020 (7)
C8	0.0419 (11)	0.0313 (9)	0.0252 (9)	-0.0041 (8)	0.0025 (8)	-0.0006 (7)
C9	0.0505 (12)	0.0342 (10)	0.0288 (10)	-0.0102 (9)	-0.0033 (9)	0.0004 (8)
C10	0.0500 (12)	0.0360 (10)	0.0291 (10)	-0.0150 (9)	-0.0025 (9)	0.0034 (8)
C15	0.0283 (9)	0.0436 (11)	0.0201 (9)	0.0028 (8)	0.0049 (7)	0.0025 (8)
C16	0.0340 (10)	0.0384 (10)	0.0208 (9)	0.0026 (8)	0.0063 (7)	0.0036 (7)
C17	0.0505 (12)	0.0392 (10)	0.0237 (9)	0.0066 (9)	0.0104 (8)	0.0038 (8)
C18	0.0478 (12)	0.0392 (11)	0.0232 (9)	0.0067 (9)	0.0108 (8)	-0.0008 (8)

*Geometric parameters (Å, °)*

O1—C15	1.258 (2)	C12A—H12F	0.9800
O2—C15	1.238 (2)	C13A—H13D	0.9800
O3—H3A	0.869 (10)	C13A—H13E	0.9800
O3—C18	1.302 (2)	C13A—H13F	0.9800
O4—C18	1.202 (2)	C14A—H14D	0.9800
N1—H1	0.870 (10)	C14A—H14E	0.9800
N1—C1	1.369 (2)	C14A—H14F	0.9800
N1—C2	1.369 (3)	C1—H1A	0.9500
N2—H2	0.880 (10)	C1—C8	1.361 (3)
N2—C11	1.518 (4)	C2—C3	1.392 (3)
N2—C14	1.543 (7)	C2—C7	1.414 (3)
N2—C10	1.540 (3)	C3—H3	0.9500
C11—H11	1.0000	C3—C4	1.375 (3)
C11—C12	1.513 (5)	C4—H4	0.9500
C11—C13	1.523 (6)	C4—C5	1.397 (3)
C12—H12A	0.9800	C5—H5	0.9500
C12—H12B	0.9800	C5—C6	1.374 (3)
C12—H12C	0.9800	C6—H6	0.9500
C13—H13A	0.9800	C6—C7	1.397 (3)
C13—H13B	0.9800	C7—C8	1.432 (3)
C13—H13C	0.9800	C8—C9	1.499 (3)
C14—H14A	0.9800	C9—H9A	0.9900
C14—H14B	0.9800	C9—H9B	0.9900
C14—H14C	0.9800	C9—C10	1.522 (3)
N2A—H2A	0.876 (10)	C10—H10A	0.9900
N2A—C11A	1.518 (6)	C10—H10B	0.9900
N2A—C13A	1.702 (10)	C10—H10C	0.9900
N2A—C10	1.543 (4)	C10—H10D	0.9900
C11A—H11A	1.0000	C15—C16	1.495 (2)
C11A—C12A	1.506 (9)	C16—H16	0.9500
C11A—C14A	1.433 (12)	C16—C17	1.303 (3)
C12A—H12D	0.9800	C17—H17	0.9500
C12A—H12E	0.9800	C17—C18	1.481 (3)

C18—O3—H3A	111.8 (17)	C11A—C14A—H14E	109.5
C1—N1—H1	127.4 (15)	C11A—C14A—H14F	109.5
C2—N1—H1	123.6 (15)	H14D—C14A—H14E	109.5
C2—N1—C1	108.93 (16)	H14D—C14A—H14F	109.5
C11—N2—H2	106 (2)	H14E—C14A—H14F	109.5
C11—N2—C14	113.1 (3)	N1—C1—H1A	124.9
C11—N2—C10	110.3 (3)	C8—C1—N1	110.14 (17)
C14—N2—H2	109 (2)	C8—C1—H1A	124.9
C10—N2—H2	105 (2)	N1—C2—C3	130.27 (18)
C10—N2—C14	112.6 (2)	N1—C2—C7	107.58 (16)
N2—C11—H11	107.5	C3—C2—C7	122.15 (18)
N2—C11—C13	103.6 (3)	C2—C3—H3	121.3
C12—C11—N2	111.5 (3)	C4—C3—C2	117.45 (19)
C12—C11—H11	107.5	C4—C3—H3	121.3
C12—C11—C13	118.8 (3)	C3—C4—H4	119.3
C13—C11—H11	107.5	C3—C4—C5	121.44 (19)
C11—C12—H12A	109.5	C5—C4—H4	119.3
C11—C12—H12B	109.5	C4—C5—H5	119.4
C11—C12—H12C	109.5	C6—C5—C4	121.1 (2)
H12A—C12—H12B	109.5	C6—C5—H5	119.4
H12A—C12—H12C	109.5	C5—C6—H6	120.4
H12B—C12—H12C	109.5	C5—C6—C7	119.22 (19)
C11—C13—H13A	109.5	C7—C6—H6	120.4
C11—C13—H13B	109.5	C2—C7—C8	106.62 (17)
C11—C13—H13C	109.5	C6—C7—C2	118.61 (17)
H13A—C13—H13B	109.5	C6—C7—C8	134.76 (18)
H13A—C13—H13C	109.5	C1—C8—C7	106.73 (16)
H13B—C13—H13C	109.5	C1—C8—C9	126.19 (18)
N2—C14—H14A	109.5	C7—C8—C9	127.05 (18)
N2—C14—H14B	109.5	C8—C9—H9A	109.3
N2—C14—H14C	109.5	C8—C9—H9B	109.3
H14A—C14—H14B	109.5	C8—C9—C10	111.81 (16)
H14A—C14—H14C	109.5	H9A—C9—H9B	107.9
H14B—C14—H14C	109.5	C10—C9—H9A	109.3
C11A—N2A—H2A	116 (4)	C10—C9—H9B	109.3
C11A—N2A—C13A	103.7 (4)	N2—C10—H10A	108.7
C11A—N2A—C10	109.8 (4)	N2—C10—H10B	108.7
C13A—N2A—H2A	94 (4)	N2A—C10—H10C	109.5
C10—N2A—H2A	101 (4)	N2A—C10—H10D	109.5
C10—N2A—C13A	131.9 (5)	C9—C10—N2	114.26 (17)
N2A—C11A—H11A	110.5	C9—C10—N2A	110.7 (2)
C12A—C11A—N2A	111.4 (5)	C9—C10—H10A	108.7
C12A—C11A—H11A	110.5	C9—C10—H10B	108.7
C14A—C11A—N2A	107.9 (6)	C9—C10—H10C	109.5
C14A—C11A—H11A	110.5	C9—C10—H10D	109.5
C14A—C11A—C12A	105.7 (5)	H10A—C10—H10B	107.6
C11A—C12A—H12D	109.5	H10C—C10—H10D	108.1

C11A—C12A—H12E	109.5	O1—C15—C16	115.80 (16)
C11A—C12A—H12F	109.5	O2—C15—O1	125.66 (16)
H12D—C12A—H12E	109.5	O2—C15—C16	118.53 (17)
H12D—C12A—H12F	109.5	C15—C16—H16	118.5
H12E—C12A—H12F	109.5	C17—C16—C15	123.04 (17)
N2A—C13A—H13D	109.5	C17—C16—H16	118.5
N2A—C13A—H13E	109.5	C16—C17—H17	117.6
N2A—C13A—H13F	109.5	C16—C17—C18	124.86 (18)
H13D—C13A—H13E	109.5	C18—C17—H17	117.6
H13D—C13A—H13F	109.5	O3—C18—C17	114.82 (16)
H13E—C13A—H13F	109.5	O4—C18—O3	123.88 (17)
C11A—C14A—H14D	109.5	O4—C18—C17	121.30 (18)
O1—C15—C16—C17	179.92 (19)	C2—C7—C8—C9	177.68 (18)
O2—C15—C16—C17	0.2 (3)	C3—C2—C7—C6	0.4 (3)
N1—C1—C8—C7	0.4 (2)	C3—C2—C7—C8	-178.93 (18)
N1—C1—C8—C9	-177.70 (17)	C3—C4—C5—C6	0.5 (3)
N1—C2—C3—C4	-179.1 (2)	C4—C5—C6—C7	-0.2 (3)
N1—C2—C7—C6	179.60 (16)	C5—C6—C7—C2	-0.3 (3)
N1—C2—C7—C8	0.2 (2)	C5—C6—C7—C8	178.9 (2)
C11—N2—C10—C9	-57.8 (3)	C6—C7—C8—C1	-179.6 (2)
C14—N2—C11—C12	57.6 (4)	C6—C7—C8—C9	-1.5 (3)
C14—N2—C11—C13	-173.6 (3)	C7—C2—C3—C4	-0.1 (3)
C14—N2—C10—C9	69.7 (3)	C7—C8—C9—C10	71.3 (3)
C11A—N2A—C10—C9	63.3 (5)	C8—C9—C10—N2	-163.3 (2)
C13A—N2A—C11A—C12A	-47.0 (6)	C8—C9—C10—N2A	162.0 (3)
C13A—N2A—C11A—C14A	-162.6 (6)	C10—N2—C11—C12	-175.2 (3)
C13A—N2A—C10—C9	-68.0 (6)	C10—N2—C11—C13	-46.4 (3)
C1—N1—C2—C3	179.1 (2)	C10—N2A—C11A—C12A	168.1 (4)
C1—N1—C2—C7	0.0 (2)	C10—N2A—C11A—C14A	52.4 (6)
C1—C8—C9—C10	-111.0 (2)	C15—C16—C17—C18	179.57 (18)
C2—N1—C1—C8	-0.2 (2)	C16—C17—C18—O3	18.9 (3)
C2—C3—C4—C5	-0.3 (3)	C16—C17—C18—O4	-160.9 (2)
C2—C7—C8—C1	-0.4 (2)		

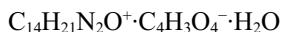
*Hydrogen-bond geometry (Å, °)*

D—H···A	D—H	H···A	D···A	D—H···A
O3—H3A···O1 <sup>i</sup>	0.87 (1)	1.66 (1)	2.5316 (18)	176 (3)
N1—H1···O1 <sup>ii</sup>	0.87 (1)	2.04 (1)	2.874 (2)	160 (2)
N2—H2···O2	0.88 (1)	1.79 (1)	2.667 (3)	173 (3)
N2A—H2A···O2	0.88 (1)	1.81 (2)	2.670 (5)	167 (6)

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

[2-(4-Hydroxy-1*H*-indol-3-yl)ethyl](methyl)propan-2-ylazanium 3-carboxyprop-2-enoate monohydrate (4-HO-MiPT)

*Crystal data*



$M_r = 366.41$

Monoclinic,  $C2/c$

$a = 29.507 (3) \text{ \AA}$

$b = 8.7445 (8) \text{ \AA}$

$c = 17.3659 (18) \text{ \AA}$

$\beta = 123.389 (3)^\circ$

$V = 3741.2 (7) \text{ \AA}^3$

$Z = 8$

$F(000) = 1568$

$D_x = 1.301 \text{ Mg m}^{-3}$

Mo  $K\alpha$  radiation,  $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 9718 reflections

$\theta = 2.9\text{--}25.3^\circ$

$\mu = 0.10 \text{ mm}^{-1}$

$T = 200 \text{ K}$

BLOCK, colourless

$0.30 \times 0.25 \times 0.20 \text{ mm}$

*Data collection*

Bruker D8 Venture CMOS

diffractometer

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan  
(SADABS; Bruker, 2016)

$T_{\min} = 0.719$ ,  $T_{\max} = 0.745$

70395 measured reflections

3458 independent reflections

2978 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.041$

$\theta_{\max} = 25.4^\circ$ ,  $\theta_{\min} = 3.1^\circ$

$h = -35 \rightarrow 35$

$k = -10 \rightarrow 10$

$l = -20 \rightarrow 20$

*Refinement*

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.041$

$wR(F^2) = 0.096$

$S = 1.08$

3458 reflections

320 parameters

12 restraints

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0299P)^2 + 3.9177P]$   
where  $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.22 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.20 \text{ e \AA}^{-3}$

Extinction correction: SHELXL,

$F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0081 (4)

*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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1	0.69858 (5)	0.51214 (13)	0.50306 (8)	0.0369 (3)	
O1W	0.52724 (8)	1.07404 (17)	0.09877 (11)	0.0664 (5)	
O2	0.56158 (8)	0.24517 (15)	0.24314 (9)	0.0786 (6)	
O3	0.61162 (5)	0.35608 (13)	0.37801 (7)	0.0421 (3)	
O4	0.54759 (7)	0.80247 (15)	0.16809 (9)	0.0621 (4)	
O5	0.49277 (7)	0.70299 (16)	0.03041 (9)	0.0718 (5)	
N1	0.77812 (6)	0.83838 (17)	0.75125 (10)	0.0411 (4)	
C1	0.73684 (6)	0.67519 (16)	0.63265 (10)	0.0277 (3)	

C2	0.69254 (6)	0.58832 (17)	0.56519 (10)	0.0304 (3)
C3	0.64541 (7)	0.58566 (19)	0.56435 (11)	0.0376 (4)
H3	0.6153	0.5269	0.5191	0.045*
C4	0.64171 (7)	0.6688 (2)	0.62958 (12)	0.0420 (4)
H4	0.6088	0.6650	0.6274	0.050*
C5	0.68368 (7)	0.7552 (2)	0.69635 (12)	0.0410 (4)
H5	0.6807	0.8105	0.7405	0.049*
C6	0.73117 (7)	0.75834 (18)	0.69665 (10)	0.0337 (4)
C7	0.81267 (7)	0.81061 (19)	0.72352 (11)	0.0376 (4)
H7	0.8477	0.8547	0.7507	0.045*
C8	0.78980 (6)	0.71090 (17)	0.65144 (10)	0.0303 (3)
C9	0.81461 (7)	0.65950 (17)	0.60004 (11)	0.0327 (4)
H9A	0.8449	0.7289	0.6150	0.039*
H9B	0.7871	0.6673	0.5330	0.039*
C10	0.83576 (7)	0.49632 (18)	0.62290 (11)	0.0319 (4)
H10A	0.8586	0.4839	0.6907	0.038*
H10B	0.8046	0.4253	0.5984	0.038*
H10C	0.810 (3)	0.427 (9)	0.621 (5)	0.038*
H10D	0.868 (3)	0.486 (9)	0.684 (5)	0.038*
N2	0.86881 (10)	0.4538 (2)	0.58322 (14)	0.0299 (6)
H2	0.8759 (9)	0.3557 (13)	0.5963 (16)	0.036*
C11	0.83717 (10)	0.4648 (2)	0.47872 (14)	0.0347 (7)
H11	0.8309	0.5755	0.4615	0.042*
C12	0.8709 (2)	0.3979 (9)	0.4448 (4)	0.0500 (13)
H12A	0.9040	0.4584	0.4692	0.075*
H12B	0.8497	0.4003	0.3772	0.075*
H12C	0.8806	0.2919	0.4661	0.075*
C13	0.7823 (3)	0.3875 (9)	0.4349 (4)	0.0421 (12)
H13A	0.7633	0.3916	0.3676	0.063*
H13B	0.7607	0.4403	0.4539	0.063*
H13C	0.7874	0.2805	0.4549	0.063*
C14	0.92160 (18)	0.5366 (7)	0.6263 (3)	0.0410 (10)
H14A	0.9364	0.5538	0.6918	0.062*
H14B	0.9158	0.6352	0.5954	0.062*
H14C	0.9472	0.4752	0.6201	0.062*
N2A	0.8395 (3)	0.4167 (7)	0.5478 (5)	0.032 (2)
H2A	0.855 (3)	0.327 (4)	0.567 (5)	0.038*
C11A	0.8776 (3)	0.5046 (8)	0.5319 (6)	0.038 (2)
H11A	0.8584	0.5998	0.4974	0.046*
C12A	0.8893 (9)	0.410 (3)	0.4699 (16)	0.063 (6)
H12D	0.9164	0.4626	0.4635	0.094*
H12E	0.8558	0.3970	0.4089	0.094*
H12F	0.9033	0.3089	0.4978	0.094*
C13A	0.9288 (7)	0.554 (3)	0.6248 (13)	0.080 (8)
H13D	0.9558	0.5949	0.6140	0.120*
H13E	0.9440	0.4655	0.6660	0.120*
H13F	0.9190	0.6331	0.6532	0.120*
C14A	0.7888 (8)	0.371 (3)	0.4588 (10)	0.041 (4)
				0.225 (5)

H14D	0.7606	0.3451	0.4699	0.061*	0.225 (5)
H14E	0.7963	0.2819	0.4332	0.061*	0.225 (5)
H14F	0.7764	0.4561	0.4150	0.061*	0.225 (5)
C15	0.57935 (7)	0.36013 (18)	0.29127 (11)	0.0373 (4)	
C16	0.56391 (6)	0.51453 (18)	0.24727 (11)	0.0348 (4)	
H16	0.5762	0.6017	0.2864	0.042*	
C17	0.53429 (7)	0.53709 (19)	0.15761 (11)	0.0380 (4)	
H17	0.5193	0.4499	0.1188	0.046*	
C18	0.52272 (7)	0.68853 (19)	0.11306 (11)	0.0378 (4)	
H1A	0.7846 (8)	0.902 (3)	0.7946 (15)	0.061 (6)*	
H1WA	0.5221 (10)	1.123 (3)	0.0534 (18)	0.075 (8)*	
H1WB	0.5371 (10)	1.140 (3)	0.1427 (18)	0.073 (7)*	
H1	0.6684 (10)	0.469 (3)	0.4607 (17)	0.076 (8)*	
H4A	0.5378 (13)	0.894 (4)	0.135 (2)	0.125 (11)*	

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0456 (7)	0.0314 (6)	0.0333 (6)	-0.0030 (5)	0.0214 (6)	-0.0072 (5)
O1W	0.1121 (14)	0.0383 (8)	0.0472 (9)	-0.0134 (8)	0.0428 (9)	-0.0065 (7)
O2	0.1293 (14)	0.0301 (7)	0.0322 (7)	0.0008 (8)	0.0165 (8)	-0.0033 (6)
O3	0.0499 (7)	0.0332 (6)	0.0275 (6)	0.0026 (5)	0.0113 (5)	0.0022 (5)
O4	0.0913 (11)	0.0326 (7)	0.0362 (7)	-0.0040 (7)	0.0184 (7)	0.0013 (6)
O5	0.0944 (11)	0.0447 (8)	0.0316 (7)	-0.0044 (8)	0.0063 (7)	0.0075 (6)
N1	0.0512 (9)	0.0411 (8)	0.0336 (7)	-0.0029 (7)	0.0251 (7)	-0.0114 (6)
C1	0.0364 (8)	0.0237 (7)	0.0244 (7)	0.0042 (6)	0.0176 (6)	0.0041 (6)
C2	0.0403 (9)	0.0228 (7)	0.0280 (7)	0.0045 (6)	0.0188 (7)	0.0062 (6)
C3	0.0365 (9)	0.0355 (9)	0.0369 (9)	0.0011 (7)	0.0178 (7)	0.0061 (7)
C4	0.0406 (9)	0.0463 (10)	0.0458 (10)	0.0100 (8)	0.0281 (8)	0.0130 (8)
C5	0.0519 (10)	0.0441 (10)	0.0381 (9)	0.0115 (8)	0.0318 (9)	0.0051 (8)
C6	0.0442 (9)	0.0309 (8)	0.0281 (8)	0.0057 (7)	0.0212 (7)	0.0028 (6)
C7	0.0390 (9)	0.0385 (9)	0.0351 (8)	-0.0027 (7)	0.0202 (7)	-0.0034 (7)
C8	0.0371 (8)	0.0264 (8)	0.0282 (7)	0.0030 (6)	0.0183 (7)	0.0022 (6)
C9	0.0404 (9)	0.0301 (8)	0.0328 (8)	0.0045 (7)	0.0234 (7)	0.0043 (6)
C10	0.0418 (9)	0.0312 (8)	0.0288 (8)	0.0047 (7)	0.0233 (7)	0.0045 (6)
N2	0.0369 (13)	0.0259 (10)	0.0279 (11)	0.0045 (9)	0.0186 (10)	0.0037 (8)
C11	0.0512 (17)	0.0285 (11)	0.0276 (12)	0.0075 (10)	0.0237 (12)	0.0049 (9)
C12	0.070 (3)	0.048 (2)	0.048 (3)	0.005 (2)	0.043 (3)	0.0000 (18)
C13	0.045 (2)	0.037 (2)	0.029 (2)	0.006 (2)	0.0113 (18)	0.001 (2)
C14	0.043 (2)	0.0354 (18)	0.048 (2)	-0.0057 (17)	0.0271 (19)	-0.0075 (14)
N2A	0.037 (4)	0.026 (4)	0.030 (4)	0.010 (3)	0.018 (4)	0.005 (3)
C11A	0.045 (5)	0.035 (4)	0.039 (5)	0.004 (4)	0.027 (5)	0.004 (3)
C12A	0.090 (15)	0.071 (10)	0.056 (12)	0.002 (11)	0.058 (12)	-0.009 (9)
C13A	0.039 (8)	0.090 (14)	0.068 (12)	-0.036 (7)	0.002 (8)	0.003 (10)
C14A	0.033 (6)	0.046 (8)	0.025 (8)	0.016 (5)	0.005 (5)	0.014 (7)
C15	0.0455 (9)	0.0306 (8)	0.0286 (8)	0.0031 (7)	0.0159 (7)	-0.0008 (7)
C16	0.0399 (9)	0.0279 (8)	0.0311 (8)	-0.0002 (7)	0.0161 (7)	-0.0021 (6)
C17	0.0415 (9)	0.0301 (8)	0.0303 (8)	0.0006 (7)	0.0121 (7)	-0.0014 (7)

C18	0.0395 (9)	0.0344 (9)	0.0293 (8)	0.0016 (7)	0.0125 (7)	0.0008 (7)
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*Geometric parameters ( $\text{\AA}$ ,  $\text{^{\circ}}$ )*

O1—C2	1.3606 (18)	N2—C11	1.520 (3)
O1—H1	0.87 (2)	N2—C14	1.493 (5)
O1W—H1WA	0.84 (3)	C11—H11	1.0000
O1W—H1WB	0.87 (3)	C11—C12	1.524 (5)
O2—C15	1.225 (2)	C11—C13	1.518 (6)
O3—C15	1.2642 (19)	C12—H12A	0.9800
O4—C18	1.293 (2)	C12—H12B	0.9800
O4—H4A	0.93 (3)	C12—H12C	0.9800
O5—C18	1.209 (2)	C13—H13A	0.9800
N1—C6	1.365 (2)	C13—H13B	0.9800
N1—C7	1.368 (2)	C13—H13C	0.9800
N1—H1A	0.87 (2)	C14—H14A	0.9800
C1—C2	1.404 (2)	C14—H14B	0.9800
C1—C6	1.413 (2)	C14—H14C	0.9800
C1—C8	1.444 (2)	N2A—H2A	0.872 (10)
C2—C3	1.383 (2)	N2A—C11A	1.507 (8)
C3—H3	0.9500	N2A—C14A	1.497 (10)
C3—C4	1.400 (2)	C11A—H11A	1.0000
C4—H4	0.9500	C11A—C12A	1.543 (10)
C4—C5	1.367 (3)	C11A—C13A	1.545 (10)
C5—H5	0.9500	C12A—H12D	0.9800
C5—C6	1.398 (2)	C12A—H12E	0.9800
C7—H7	0.9500	C12A—H12F	0.9800
C7—C8	1.361 (2)	C13A—H13D	0.9800
C8—C9	1.501 (2)	C13A—H13E	0.9800
C9—H9A	0.9900	C13A—H13F	0.9800
C9—H9B	0.9900	C14A—H14D	0.9800
C9—C10	1.520 (2)	C14A—H14E	0.9800
C10—H10A	0.9900	C14A—H14F	0.9800
C10—H10B	0.9900	C15—C16	1.494 (2)
C10—H10C	0.97 (8)	C16—H16	0.9500
C10—H10D	0.96 (8)	C16—C17	1.315 (2)
C10—N2	1.518 (2)	C17—H17	0.9500
C10—N2A	1.536 (6)	C17—C18	1.476 (2)
N2—H2	0.882 (10)		
C2—O1—H1	111.3 (16)	C11—C12—H12A	109.5
H1WA—O1W—H1WB	106 (2)	C11—C12—H12B	109.5
C18—O4—H4A	110 (2)	C11—C12—H12C	109.5
C6—N1—C7	109.43 (14)	H12A—C12—H12B	109.5
C6—N1—H1A	125.7 (14)	H12A—C12—H12C	109.5
C7—N1—H1A	124.7 (14)	H12B—C12—H12C	109.5
C2—C1—C6	118.41 (14)	C11—C13—H13A	109.5
C2—C1—C8	134.44 (14)	C11—C13—H13B	109.5

C6—C1—C8	107.03 (13)	C11—C13—H13C	109.5
O1—C2—C1	117.29 (14)	H13A—C13—H13B	109.5
O1—C2—C3	123.58 (15)	H13A—C13—H13C	109.5
C3—C2—C1	119.13 (14)	H13B—C13—H13C	109.5
C2—C3—H3	119.7	N2—C14—H14A	109.5
C2—C3—C4	120.57 (16)	N2—C14—H14B	109.5
C4—C3—H3	119.7	N2—C14—H14C	109.5
C3—C4—H4	118.8	H14A—C14—H14B	109.5
C5—C4—C3	122.39 (16)	H14A—C14—H14C	109.5
C5—C4—H4	118.8	H14B—C14—H14C	109.5
C4—C5—H5	121.6	C10—N2A—H2A	110 (6)
C4—C5—C6	116.89 (15)	C11A—N2A—C10	109.5 (6)
C6—C5—H5	121.6	C11A—N2A—H2A	104 (6)
N1—C6—C1	107.22 (14)	C14A—N2A—C10	120.0 (13)
N1—C6—C5	130.14 (15)	C14A—N2A—H2A	100 (5)
C5—C6—C1	122.61 (15)	C14A—N2A—C11A	111.7 (11)
N1—C7—H7	124.8	N2A—C11A—H11A	107.5
C8—C7—N1	110.45 (15)	N2A—C11A—C12A	109.2 (12)
C8—C7—H7	124.8	N2A—C11A—C13A	110.6 (13)
C1—C8—C9	128.47 (14)	C12A—C11A—H11A	107.5
C7—C8—C1	105.86 (13)	C12A—C11A—C13A	114.4 (14)
C7—C8—C9	125.56 (15)	C13A—C11A—H11A	107.5
C8—C9—H9A	109.0	C11A—C12A—H12D	109.5
C8—C9—H9B	109.0	C11A—C12A—H12E	109.5
C8—C9—C10	112.98 (12)	C11A—C12A—H12F	109.5
H9A—C9—H9B	107.8	H12D—C12A—H12E	109.5
C10—C9—H9A	109.0	H12D—C12A—H12F	109.5
C10—C9—H9B	109.0	H12E—C12A—H12F	109.5
C9—C10—H10A	109.1	C11A—C13A—H13D	109.5
C9—C10—H10B	109.1	C11A—C13A—H13E	109.5
C9—C10—H10C	112 (4)	C11A—C13A—H13F	109.5
C9—C10—H10D	113 (5)	H13D—C13A—H13E	109.5
C9—C10—N2A	114.5 (3)	H13D—C13A—H13F	109.5
H10A—C10—H10B	107.8	H13E—C13A—H13F	109.5
H10C—C10—H10D	105 (6)	N2A—C14A—H14D	109.5
N2—C10—C9	112.69 (13)	N2A—C14A—H14E	109.5
N2—C10—H10A	109.1	N2A—C14A—H14F	109.5
N2—C10—H10B	109.1	H14D—C14A—H14E	109.5
N2A—C10—H10C	97 (4)	H14D—C14A—H14F	109.5
N2A—C10—H10D	114 (5)	H14E—C14A—H14F	109.5
C10—N2—H2	104.1 (17)	O2—C15—O3	123.28 (15)
C10—N2—C11	113.62 (18)	O2—C15—C16	119.80 (14)
C11—N2—H2	105.8 (16)	O3—C15—C16	116.91 (14)
C14—N2—C10	114.0 (2)	C15—C16—H16	118.0
C14—N2—H2	107.9 (16)	C17—C16—C15	123.93 (15)
C14—N2—C11	110.7 (3)	C17—C16—H16	118.0
N2—C11—H11	108.0	C16—C17—H17	117.7
N2—C11—C12	109.5 (3)	C16—C17—C18	124.51 (15)

C12—C11—H11	108.0	C18—C17—H17	117.7
C13—C11—N2	110.8 (3)	O4—C18—C17	115.49 (14)
C13—C11—H11	108.0	O5—C18—O4	122.93 (16)
C13—C11—C12	112.2 (4)	O5—C18—C17	121.56 (15)
O1—C2—C3—C4	-178.98 (14)	C7—C8—C9—C10	106.26 (18)
O2—C15—C16—C17	-4.2 (3)	C8—C1—C2—O1	3.2 (2)
O3—C15—C16—C17	174.39 (17)	C8—C1—C2—C3	-176.09 (16)
N1—C7—C8—C1	0.49 (18)	C8—C1—C6—N1	-0.47 (17)
N1—C7—C8—C9	176.96 (14)	C8—C1—C6—C5	177.70 (14)
C1—C2—C3—C4	0.2 (2)	C8—C9—C10—N2	-170.62 (16)
C1—C8—C9—C10	-78.1 (2)	C8—C9—C10—N2A	156.1 (4)
C2—C1—C6—N1	-177.00 (13)	C9—C10—N2—C11	-60.2 (2)
C2—C1—C6—C5	1.2 (2)	C9—C10—N2—C14	67.9 (3)
C2—C1—C8—C7	175.72 (16)	C9—C10—N2A—C11A	60.3 (6)
C2—C1—C8—C9	-0.6 (3)	C9—C10—N2A—C14A	-70.7 (12)
C2—C3—C4—C5	-0.1 (3)	C10—N2—C11—C12	-171.7 (4)
C3—C4—C5—C6	0.4 (2)	C10—N2—C11—C13	-47.3 (4)
C4—C5—C6—N1	176.71 (17)	C10—N2A—C11A—C12A	169.9 (12)
C4—C5—C6—C1	-1.0 (2)	C10—N2A—C11A—C13A	43.1 (13)
C6—N1—C7—C8	-0.81 (19)	C14—N2—C11—C12	58.5 (4)
C6—C1—C2—O1	178.51 (13)	C14—N2—C11—C13	-177.2 (4)
C6—C1—C2—C3	-0.7 (2)	C14A—N2A—C11A—C12A	-54.8 (18)
C6—C1—C8—C7	-0.01 (17)	C14A—N2A—C11A—C13A	178.4 (18)
C6—C1—C8—C9	-176.34 (14)	C15—C16—C17—C18	-174.40 (16)
C7—N1—C6—C1	0.78 (18)	C16—C17—C18—O4	5.5 (3)
C7—N1—C6—C5	-177.20 (16)	C16—C17—C18—O5	-175.90 (19)

*Hydrogen-bond geometry (Å, °)*

D—H···A	D—H	H···A	D···A	D—H···A
N2—H2···O2 <sup>i</sup>	0.88 (1)	2.51 (2)	3.085 (2)	124 (2)
N2—H2···O3 <sup>i</sup>	0.88 (1)	1.89 (1)	2.775 (2)	178 (2)
N2A—H2A···O3 <sup>i</sup>	0.87 (1)	1.85 (2)	2.717 (6)	172 (8)
O1—H1···O3	0.87 (2)	1.79 (3)	2.6512 (17)	172 (2)
O4—H4A···O1W	0.93 (3)	1.66 (3)	2.579 (2)	167 (3)
O1W—H1WA···O5 <sup>ii</sup>	0.84 (3)	1.98 (3)	2.779 (2)	160 (2)
O1W—H1WB···O2 <sup>iii</sup>	0.87 (3)	1.74 (3)	2.599 (2)	170 (2)

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