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Synthesis, crystal structure and Hirshfeld surface analysis of (*E*)-benzo[*d*][1,3]dioxole-5-carbaldehyde oxime

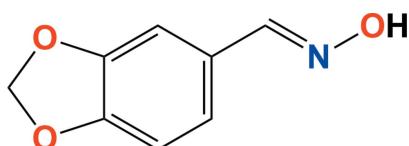
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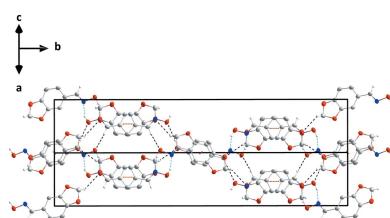
The asymmetric unit of the title molecule, $C_8H_7NO_3$, consists of two molecules differing slightly in conformation and in their intermolecular interactions in the solid. The dihedral angle between the benzene and dioxolane rings is 0.20 (7) $^\circ$ in one molecule and 0.31 (7) $^\circ$ in the other. In the crystal, the two molecules are linked into dimers through pairwise O—H···N hydrogen bonds, with these units being formed into stacks by two different sets of aromatic π -stacking interactions. The stacks are connected by C—H···O hydrogen bonds. A Hirshfeld surface analysis indicates that the most significant contacts in the crystal packing are H···O/O···H (36.7%), H···H (32.2%) and C···H/H···C (12.7%).

1. Chemical context

Oxime compounds containing an $R_2C\equiv N—OH$ functional group have been studied for many years because of their important role as acetylcholinesterase reactivators and their utility as therapeutic agents for various diseases (Musilek *et al.*, 2011; Canario *et al.*, 2018). Various oximes have been identified in plants as biosynthetic intermediates and can facilitate a range of processes associated with plant growth and development (Sørensen *et al.*, 2018). Oximes also have a wide range of biological activities, such as human immunodeficiency virus (HIV) agents that can inhibit HIV protease (Komai *et al.*, 1997) and can act as anti-inflammatories (Li *et al.*, 2018; Kwon *et al.*, 2014). The introduction of an oxime group into an appropriate chemical backbone is a reasonable approach for the preparation of cytotoxic agents and many oxime derivatives have been reported to have therapeutic activity for cancer (Canario *et al.*, 2018; Shen *et al.*, 2015) and neurodegenerative disorders (Avrahami *et al.*, 2013; Yuskaitis *et al.*, 2009).

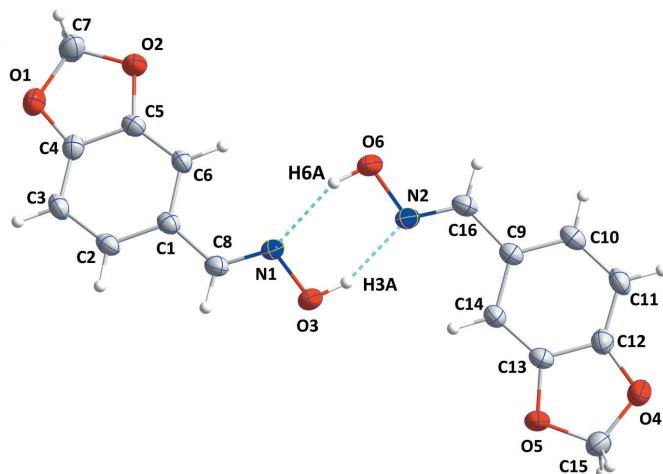


As part of our studies in this area, we now describe the synthesis, structure and Hirshfeld surface analysis of the title compound (I).



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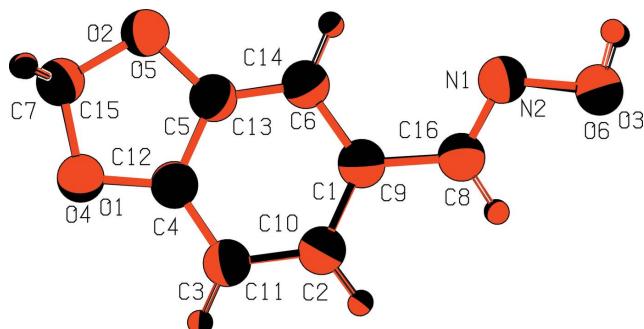
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**Figure 1**

The asymmetric unit with 50% probability ellipsoids. The $\text{O}-\text{H}\cdots\text{N}$ hydrogen bonds are depicted by dashed lines.

2. Structural commentary

The asymmetric unit (Fig. 1) consists of two independent molecules differing slightly in the orientation of some hydrogen atoms. The benzodioxolane portion of the molecule containing O1 is planar to within 0.0171 (12) Å (r.m.s. deviation of the fitted atoms = 0.0091 Å) with C7 deviating by 0.0171 (12) Å from one side of the mean plane and O1 by 0.0170 (10) Å from the other, indicating a slight twist in the dioxolane ring. The corresponding portion of the second molecule containing O4 is planar to within 0.0041 (11) Å (r.m.s. deviation of the fitted atoms = 0.0030 Å), indicating a conformational difference, albeit small, between the two molecules. The overlay fit of inverted molecule 2 on molecule 1 is shown in Fig. 2 with the weighted r.m.s. fit of the 12 non-H atoms being 0.036 Å and showing the major differences to be in the hydrogen-atom positions. The C6–C1–C8–N1 and C1–C8–N1–O3 torsion angles are, respectively, 3.9 (2) and –179.96 (11)°, indicating the side chain to be nearly coplanar with the benzodioxolane unit. The corresponding torsion angles in the second molecule are virtually the same as above. The two molecules are connected into dimers through O3–H3A···N2 and O6–H6A···N1 hydrogen bonds (Table 1 and Fig. 1), generating $R_2^2(6)$ loops.

**Figure 2**

A least-squares overlay of the two independent molecules [inverted O4 molecule (red) on O1 molecule (black)].

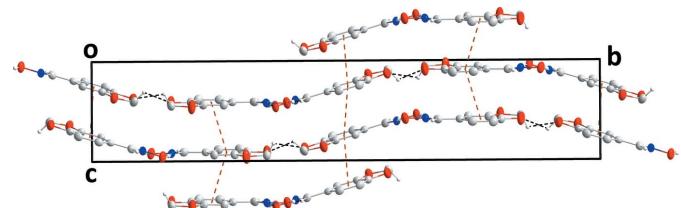
Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O3–H3A···N2	0.87	1.93	2.7549 (16)	158
C7–H7B···O4 ⁱ	0.99	2.58	3.239 (2)	124
C8–H8···O6 ⁱⁱ	0.95	2.43	3.3754 (18)	173
O6–H6A···N1	0.87	1.97	2.7989 (17)	158
C15–H15A···O1 ⁱⁱⁱ	0.99	2.54	3.1775 (19)	122
C16–H16···O3 ^{iv}	0.95	2.59	3.5173 (18)	167

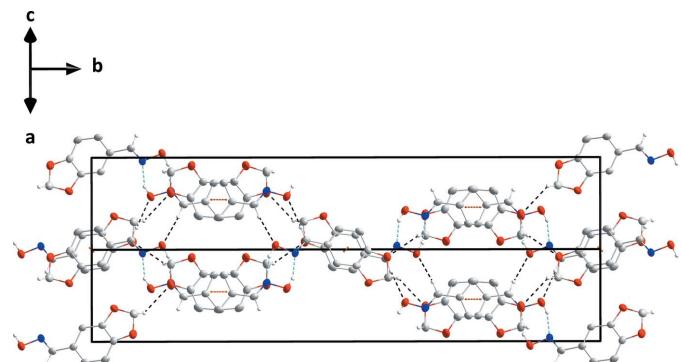
Symmetry codes: (i) $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$; (ii) $x - 1, y, z$; (iii) $-x + \frac{1}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$; (iv) $x + 1, y, z$.

3. Supramolecular features

In the crystal, the dimers are connected into stacks extending along the [101] direction through slipped π -stacking interactions between the six-membered ($Cg2$: C1–C6 and $Cg5$: C9–C14) rings. For the C1–C6 rings, the centroid–centroid distance is 3.6024 (11) Å with a slippage of 1.185 Å between molecules at x, y, z and $-x, -y + 1, -z$. These paired molecules make weak, slipped π -stacking interactions with corresponding pairs at $-x + 1, -y + 1, -z + 1$ with a centroid–centroid distance of 3.8479 (11) Å and a slippage of 1.947 Å. The C9–C14 ring has slipped π -stacking interactions with its counterparts in molecules at $x - \frac{1}{2}, -y + \frac{1}{2}, z - \frac{1}{2}$ and at $x + \frac{1}{2}, -y + \frac{1}{2}, z + \frac{1}{2}$ with centroid–centroid distances of 3.8380 (11) Å and dihedral angles of 2.41 (6)° for both. The slippages for these interactions (Fig. 3) are 1.572 and 1.662 Å, respectively. These differences in the π -stacking interactions also support the independence of the two molecules in the asymmetric unit.

**Figure 3**

View of the packing seen along the a -axis direction with $\text{O}-\text{H}\cdots\text{N}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds and π -stacking interactions depicted, respectively, by light blue, black and orange dashed lines.

**Figure 4**

View of the packing seen along the [101] direction. Intermolecular interactions are depicted as in Fig. 3.

The stacks are associated through C7—H7B···O4, C8—H8···O6, C15—H15A···O1 and C16—H16···O3 hydrogen bonds (Table 1 and Fig. 4).

4. Database survey

A search using CCDC ConQuest of the Cambridge Structural Database (CSD, Version 5.44, updated to April 2023; Groom *et al.*, 2016) using the title molecule with all hydrogen atoms deleted gave 26 hits. Most of these contain the search fragment as part of a larger, often polycyclic molecule, but three are considered similar to (I). These are *N*-[1-(2,2-dimethyl-2*H*-1,3-benzodioxol-5-yl)-2-(1*H*-imidazol-1-yl)ethylidene]hydroxylamine (CSD refcode: GAVWUZ; Ren *et al.*, 2022), in which the benzo[d][1,3]dioxole unit is similar to that in (I), 1-(1,3-benzodioxol-5-yl)-*N*-hydroxy-3-(1*H*-imidazol-1-yl)propan-1-imine isopropanol solvate (QEKMAX; Al-Wabli *et al.*, 2017), in which the benzo[d][1,3]dioxole-5-carbaldehydeoxime takes a (Z) form and (Z)-3,4-methylenedioxybenzaldehyde oxinium 4-toluenesulfonate (VADDIN; Jerslev *et al.*, 1988), in which the benzo[d][1,3]dioxole unit is similar to that in (I).

5. Hirshfeld surface analysis

The Hirshfeld surface analysis was performed with *Crystal Explorer* (Version 21.5; Spackman *et al.*, 2021). Fig. 5 shows views of the d_{norm} surfaces for the two molecules in the asymmetric unit plotted over the limits from −0.63 to 1.18 a.u for molecule 1 and −0.63 to 1.07 a.u for molecule 2. The O···N hydrogen bonds, which generate the dimers are indicated by the bright-red spots in Fig. 5(a) and 5(b), respectively. Fig. 6 presents the two-dimensional fingerprint plots involving all intermolecular interactions [Fig. 6(a)] and delineated into C···H/H···C [Fig. 6(c)], and H···O/O···H [Fig. 6(h)] interactions. For completeness, the H···H interactions constitute 32.2% of the surface [Fig. 6(b)]. The other interactions contribute small amounts, *viz.*, C···N/N···C (1.0%), C···O/O···C (2.4%), C···C (9.5%), H···N/N···H (4.1%), N···O/O···N (1.1%), N···N (0.0%) and O···O (0.4%).

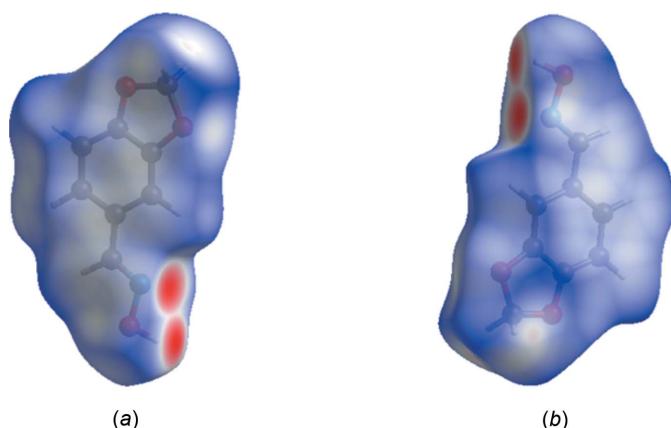


Figure 5

The Hirshfeld surface plots for (I): (a) d_{norm} for the O1-containing molecule; (b) d_{norm} for the O4-containing molecule.

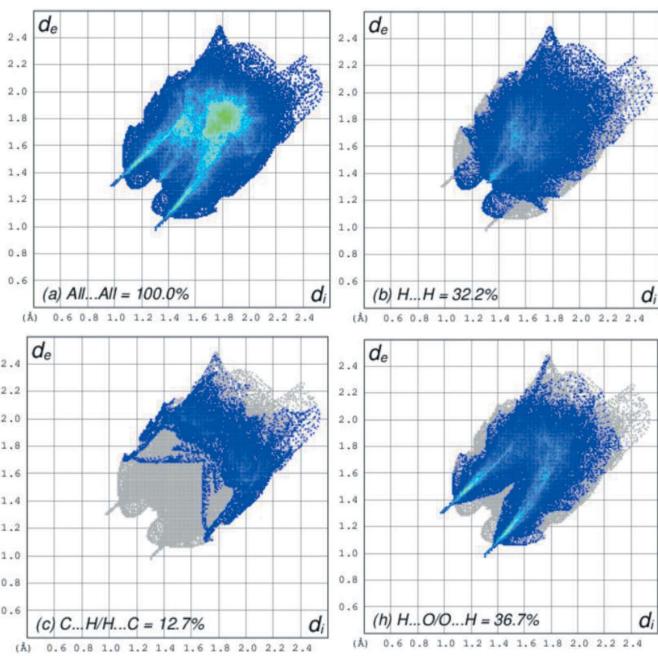


Figure 6

Fingerprint plots for (I) (both molecules): (a) all interactions; (b) H···H; (c) C···H/H···C and (d) H···O/O···H.

6. Synthesis and crystallization

A solution of 5.0 g of sodium hydroxide dissolved in 20 ml of water was mixed with 8.0 g of hydroxylamine hydrochloride

Table 2
Experimental details.

Crystal data	
Chemical formula	$\text{C}_8\text{H}_7\text{NO}_3$
M_r	165.15
Crystal system, space group	Monoclinic, $P2_1/n$
Temperature (K)	150
a, b, c (Å)	6.8724 (14), 33.502 (7), 7.3449 (15)
β (°)	117.238 (3)
V (Å ³)	1503.6 (5)
Z	8
Radiation type	Mo $K\alpha$
μ (mm ^{−1})	0.11
Crystal size (mm)	0.36 × 0.17 × 0.10
Data collection	
Diffractometer	Bruker SMART APEX CCD
Absorption correction	Multi-scan (<i>SADABS</i> ; Krause <i>et al.</i> , 2015)
T_{\min}, T_{\max}	0.82, 0.99
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	28032, 3858, 2836
R_{int}	0.048
(sin θ/λ) _{max} (Å ^{−1})	0.675
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.046, 0.127, 1.05
No. of reflections	3858
No. of parameters	217
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å ^{−3})	0.31, −0.19

Computer programs: *APEX3* and *SAINT* (Bruker, 2016), *SHELXT* (Sheldrick, 2015a), *SHELXL2018/3* (Sheldrick, 2015b), *DIAMOND* (Brandenburg & Putz, 2012), *PLATON* (Spek, 2020) and *publCIF* (Westrip, 2010).

dissolved in 15 ml of water, then 8.0 g of benzo[*d*][1,3]dioxole-5-carbaldehyde dissolved in 50 ml of ethanol was added to the mixture. After 5 h of stirring at 273 K, the product was allowed to precipitate and then filtered with a yield of 90%. Single crystals were recrystallized from ethanol solution.

7. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. H atoms attached to carbon were placed in calculated positions ($C-H = 0.95\text{--}0.99 \text{\AA}$) while those attached to oxygen were placed in locations derived from a difference map and their coordinates adjusted to give $O-H = 0.87 \text{\AA}$. All were included as riding contributions with isotropic displacement parameters 1.2–1.5 times those of the attached atoms.

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References

- Al-Wabli, R., Al-Ghamdi, A., Ghabbour, H., Al-Agamy, M., Monicka, J., Joe, I. & Attia, M. (2017). *Molecules*, **22**, 373. <https://doi.org/10.3390/molecules22030373>
- Avrahami, L., Farfara, D., Shaham-Kol, M., Vassar, R., Frenkel, D. & Eldar-Finkelman, H. (2013). *J. Biol. Chem.* **288**, 1295–1306.
- Brandenburg, K. & Putz, H. (2012). *DIAMOND*. Crystal Impact GbR, Bonn, Germany.
- Bruker (2016). *APEX3* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Canario, C., Silvestre, S., Falcao, A. & Alves, G. (2018). *Curr. Med. Chem.* **25**, 660–686.
- Groom, C. R., Bruno, I. J., Lightfoot, M. P. & Ward, S. C. (2016). *Acta Cryst. B* **72**, 171–179.
- Jerslev, B., Larsen, S. & Hansen, F. (1988). *Acta Chem. Scand.* **42b**, 646–649.
- Komai, T., Yagi, R., Suzuki-Sunagawa, H., Ishikawa, Y., Kasuya, A., Miyamoto, S., Handa, H. & Nishigaki, T. (1997). *Biochem. Biophys. Res. Commun.* **230**, 557–561.
- Krause, L., Herbst-Irmer, R., Sheldrick, G. M. & Stalke, D. (2015). *J. Appl. Cryst.* **48**, 3–10.
- Kwon, Y. J., Yoon, C. H., Lee, S. W., Park, Y. B., Lee, S. K. & Park, M. C. (2014). *Joint Bone Spine*, **81**, 240–246.
- Li, Q., Zhang, J., Chen, L. Z., Wang, J. Q., Zhou, H. P., Tang, W. J., Xue, W. & Liu, X. H. (2018). *J. Enzyme Inhib. Med. Chem.* **33**, 130–138.
- Musilek, K., Dolezal, M., Gunn-Moore, F. & Kuca, K. (2011). *Med. Res. Rev.* **31**, 548–575.
- Ren, B., Guo, C., Liu, R., Bian, Z., Liu, R., Huang, L. & Tang, J. (2022). *Eur. J. Med. Chem.* **228**, 114031.
- Sheldrick, G. M. (2015a). *Acta Cryst. A* **71**, 3–8.
- Sheldrick, G. M. (2015b). *Acta Cryst. C* **71**, 3–8.
- Shen, S., Xu, N., Klamer, G., Ko, K. H., Khoo, M., Ma, D., Moore, J., O'Brien, T. A. & Dolnikov, A. (2015). *Stem Cells Dev.* **24**, 724–736.
- Sørensen, M., Neilson, E. H. J. & Møller, B. L. (2018). *Mol. Plant.* **11**, 95–117.
- Spackman, P. R., Turner, M. J., McKinnon, J. J., Wolff, S. K., Grimwood, D. J., Jayatilaka, D. & Spackman, M. A. (2021). *J. Appl. Cryst.* **54**, 1006–1011.
- Spek, A. L. (2020). *Acta Cryst. E* **76**, 1–11.
- Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.
- Yuskaitis, C. J. & Jope, R. S. (2009). *Cell. Signal.* **21**, 264–273.

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Computing details

Data collection: *APEX3* (Bruker, 2016); cell refinement: *SAINT* (Bruker, 2016); data reduction: *SAINT* (Bruker, 2016); program(s) used to solve structure: *SHELXT* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2015b); molecular graphics: *DIAMOND* (Brandenburg & Putz, 2012), *PLATON* (Spek, 2020); software used to prepare material for publication: *SHELXL2018/3* (Sheldrick, 2015b), *PLATON* (Spek, 2020) and *publCIF* (Westrip, 2010).

(*E*)-Benzo[*d*][1,3]dioxole-5-carbaldehyde oxime

Crystal data

$C_8H_7NO_3$
 $M_r = 165.15$
Monoclinic, $P2_1/n$
 $a = 6.8724 (14)$ Å
 $b = 33.502 (7)$ Å
 $c = 7.3449 (15)$ Å
 $\beta = 117.238 (3)^\circ$
 $V = 1503.6 (5)$ Å³
 $Z = 8$

$F(000) = 688$
 $D_x = 1.459$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 6661 reflections
 $\theta = 2.4\text{--}28.1^\circ$
 $\mu = 0.11$ mm⁻¹
 $T = 150$ K
Plate, colourless
0.36 × 0.17 × 0.10 mm

Data collection

Bruker SMART APEX CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 8.3333 pixels mm⁻¹
 ϕ and ω scans
Absorption correction: multi-scan
(*SADABS*; Krause *et al.*, 2015)
 $T_{\min} = 0.82$, $T_{\max} = 0.99$

28032 measured reflections
3858 independent reflections
2836 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$
 $\theta_{\max} = 28.7^\circ$, $\theta_{\min} = 2.4^\circ$
 $h = -9 \rightarrow 9$
 $k = -45 \rightarrow 43$
 $l = -9 \rightarrow 9$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.127$
 $S = 1.05$
3858 reflections

217 parameters
0 restraints
Primary atom site location: dual
Secondary atom site location: difference Fourier
map
Hydrogen site location: mixed

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0621P)^2 + 0.2227P]$
 where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.31 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. The diffraction data were obtained from 3 sets of 400 frames, each of width 0.5° in ω , collected at $\varphi = 0.00, 90.00$ and 180.00° and 2 sets of 800 frames, each of width 0.45° in φ , collected at $\omega = -30.00$ and 210.00° . The scan time was 20 sec/frame.

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. H-atoms attached to carbon were placed in calculated positions ($C-H = 0.95 - 0.99 \text{ \AA}$) while those attached to oxygen were placed in locations derived from a difference map and their coordinates adjusted to give $O-H = 0.87 \text{ \AA}$. All were included as riding contributions with isotropic displacement parameters 1.2 - 1.5 times those of the attached atoms.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.21075 (18)	0.58110 (3)	0.12890 (17)	0.0385 (3)
O2	0.50131 (18)	0.54433 (3)	0.13879 (17)	0.0377 (3)
O3	0.40957 (17)	0.36128 (3)	0.45505 (17)	0.0377 (3)
H3A	0.516808	0.349351	0.445154	0.057*
N1	0.43129 (18)	0.40031 (3)	0.39281 (18)	0.0287 (3)
C1	0.2551 (2)	0.46413 (4)	0.30915 (19)	0.0254 (3)
C2	0.0814 (2)	0.48668 (4)	0.3030 (2)	0.0290 (3)
H2	-0.020388	0.474321	0.339202	0.035*
C3	0.0524 (2)	0.52685 (4)	0.2452 (2)	0.0316 (3)
H3	-0.065828	0.542093	0.241589	0.038*
C4	0.2043 (2)	0.54306 (4)	0.1939 (2)	0.0280 (3)
C5	0.3775 (2)	0.52084 (4)	0.1994 (2)	0.0262 (3)
C6	0.4081 (2)	0.48172 (4)	0.25517 (19)	0.0252 (3)
H6	0.527136	0.466906	0.257669	0.030*
C7	0.4031 (3)	0.58311 (4)	0.0990 (3)	0.0379 (4)
H7A	0.363159	0.591287	-0.043339	0.045*
H7B	0.507199	0.602939	0.193416	0.045*
C8	0.2711 (2)	0.42252 (4)	0.3722 (2)	0.0285 (3)
H8	0.159251	0.411565	0.398925	0.034*
O4	0.79029 (19)	0.15551 (3)	0.4126 (2)	0.0489 (3)
O5	0.54405 (19)	0.19665 (3)	0.4559 (2)	0.0488 (3)
O6	0.84324 (18)	0.38394 (3)	0.42029 (17)	0.0392 (3)
H6A	0.725059	0.395443	0.409864	0.059*
N2	0.78297 (19)	0.34360 (4)	0.42083 (18)	0.0299 (3)
C9	0.8894 (2)	0.27623 (4)	0.4103 (2)	0.0266 (3)
C10	1.0360 (2)	0.25112 (4)	0.3838 (2)	0.0319 (3)
H10	1.152810	0.262583	0.366388	0.038*

C11	1.0166 (2)	0.20967 (5)	0.3820 (2)	0.0354 (3)
H11	1.116764	0.192674	0.363450	0.043*
C12	0.8452 (2)	0.19468 (4)	0.4082 (2)	0.0320 (3)
C13	0.6986 (2)	0.21941 (4)	0.4347 (2)	0.0302 (3)
C14	0.7148 (2)	0.25990 (4)	0.4373 (2)	0.0292 (3)
H14	0.613289	0.276437	0.456418	0.035*
C15	0.5989 (3)	0.15609 (4)	0.4420 (2)	0.0369 (3)
H15A	0.476274	0.142790	0.325521	0.044*
H15B	0.626735	0.141616	0.569104	0.044*
C16	0.9197 (2)	0.31931 (4)	0.4083 (2)	0.0299 (3)
H16	1.043047	0.329466	0.397484	0.036*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0392 (6)	0.0285 (6)	0.0509 (7)	0.0074 (4)	0.0234 (5)	0.0055 (5)
O2	0.0400 (6)	0.0297 (6)	0.0560 (7)	0.0021 (4)	0.0328 (5)	0.0076 (5)
O3	0.0355 (6)	0.0243 (5)	0.0599 (7)	-0.0013 (4)	0.0277 (5)	0.0051 (5)
N1	0.0266 (6)	0.0231 (6)	0.0357 (6)	-0.0034 (4)	0.0137 (5)	-0.0002 (5)
C1	0.0204 (6)	0.0296 (7)	0.0245 (6)	-0.0017 (5)	0.0088 (5)	-0.0042 (5)
C2	0.0205 (7)	0.0358 (8)	0.0324 (7)	-0.0030 (5)	0.0137 (6)	-0.0045 (6)
C3	0.0223 (7)	0.0361 (8)	0.0373 (8)	0.0048 (6)	0.0144 (6)	-0.0047 (6)
C4	0.0270 (7)	0.0264 (7)	0.0282 (7)	0.0034 (5)	0.0105 (6)	-0.0011 (5)
C5	0.0240 (7)	0.0308 (7)	0.0257 (6)	-0.0017 (5)	0.0131 (5)	-0.0023 (5)
C6	0.0212 (6)	0.0281 (7)	0.0275 (7)	0.0020 (5)	0.0121 (5)	-0.0020 (5)
C7	0.0463 (9)	0.0327 (8)	0.0414 (9)	0.0046 (7)	0.0259 (7)	0.0068 (6)
C8	0.0229 (7)	0.0298 (7)	0.0343 (7)	-0.0043 (5)	0.0143 (6)	-0.0035 (6)
O4	0.0423 (7)	0.0275 (6)	0.0825 (9)	0.0000 (5)	0.0335 (7)	-0.0056 (5)
O5	0.0397 (6)	0.0279 (6)	0.0940 (10)	-0.0066 (5)	0.0439 (7)	-0.0038 (6)
O6	0.0347 (6)	0.0276 (6)	0.0612 (7)	-0.0021 (4)	0.0269 (5)	0.0071 (5)
N2	0.0272 (6)	0.0265 (6)	0.0360 (6)	-0.0026 (5)	0.0145 (5)	0.0037 (5)
C9	0.0208 (6)	0.0313 (7)	0.0277 (7)	-0.0005 (5)	0.0110 (5)	0.0000 (5)
C10	0.0214 (7)	0.0408 (9)	0.0354 (8)	0.0005 (6)	0.0147 (6)	-0.0002 (6)
C11	0.0240 (7)	0.0399 (9)	0.0435 (8)	0.0058 (6)	0.0165 (6)	-0.0040 (6)
C12	0.0265 (7)	0.0287 (8)	0.0377 (8)	0.0025 (5)	0.0119 (6)	-0.0026 (6)
C13	0.0211 (7)	0.0319 (8)	0.0380 (8)	-0.0029 (5)	0.0139 (6)	-0.0017 (6)
C14	0.0227 (7)	0.0296 (7)	0.0371 (8)	0.0011 (5)	0.0154 (6)	-0.0017 (6)
C15	0.0358 (8)	0.0286 (8)	0.0441 (9)	-0.0017 (6)	0.0164 (7)	0.0016 (6)
C16	0.0237 (7)	0.0357 (8)	0.0331 (7)	-0.0029 (6)	0.0153 (6)	0.0023 (6)

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

O1—C4	1.3685 (17)	O4—C12	1.3699 (18)
O1—C7	1.4377 (19)	O4—C15	1.427 (2)
O2—C5	1.3742 (16)	O5—C13	1.3730 (17)
O2—C7	1.4311 (17)	O5—C15	1.4264 (18)
O3—N1	1.4154 (15)	O6—N2	1.4140 (15)
O3—H3A	0.8702	O6—H6A	0.8701

N1—C8	1.2790 (18)	N2—C16	1.2776 (18)
C1—C2	1.3957 (18)	C9—C10	1.3931 (19)
C1—C6	1.4110 (18)	C9—C14	1.4129 (18)
C1—C8	1.4570 (19)	C9—C16	1.4595 (19)
C2—C3	1.398 (2)	C10—C11	1.394 (2)
C2—H2	0.9500	C10—H10	0.9500
C3—C4	1.373 (2)	C11—C12	1.373 (2)
C3—H3	0.9500	C11—H11	0.9500
C4—C5	1.3890 (19)	C12—C13	1.3849 (19)
C5—C6	1.3603 (19)	C13—C14	1.3606 (19)
C6—H6	0.9500	C14—H14	0.9500
C7—H7A	0.9900	C15—H15A	0.9900
C7—H7B	0.9900	C15—H15B	0.9900
C8—H8	0.9500	C16—H16	0.9500
C4—O1—C7	106.02 (11)	C12—O4—C15	105.87 (11)
C5—O2—C7	106.39 (11)	C13—O5—C15	106.11 (11)
N1—O3—H3A	100.3	N2—O6—H6A	99.2
C8—N1—O3	111.31 (11)	C16—N2—O6	112.48 (11)
C2—C1—C6	120.04 (13)	C10—C9—C14	120.06 (13)
C2—C1—C8	117.85 (12)	C10—C9—C16	118.69 (12)
C6—C1—C8	122.10 (12)	C14—C9—C16	121.24 (12)
C1—C2—C3	122.15 (13)	C9—C10—C11	122.04 (13)
C1—C2—H2	118.9	C9—C10—H10	119.0
C3—C2—H2	118.9	C11—C10—H10	119.0
C4—C3—C2	116.33 (12)	C12—C11—C10	116.57 (13)
C4—C3—H3	121.8	C12—C11—H11	121.7
C2—C3—H3	121.8	C10—C11—H11	121.7
O1—C4—C3	127.89 (13)	O4—C12—C11	128.09 (13)
O1—C4—C5	110.19 (12)	O4—C12—C13	110.11 (13)
C3—C4—C5	121.91 (13)	C11—C12—C13	121.80 (14)
C6—C5—O2	128.03 (12)	C14—C13—O5	127.91 (12)
C6—C5—C4	122.51 (12)	C14—C13—C12	122.58 (13)
O2—C5—C4	109.45 (12)	O5—C13—C12	109.51 (13)
C5—C6—C1	117.05 (12)	C13—C14—C9	116.94 (12)
C5—C6—H6	121.5	C13—C14—H14	121.5
C1—C6—H6	121.5	C9—C14—H14	121.5
O2—C7—O1	107.86 (11)	O5—C15—O4	108.40 (12)
O2—C7—H7A	110.1	O5—C15—H15A	110.0
O1—C7—H7A	110.1	O4—C15—H15A	110.0
O2—C7—H7B	110.1	O5—C15—H15B	110.0
O1—C7—H7B	110.1	O4—C15—H15B	110.0
H7A—C7—H7B	108.4	H15A—C15—H15B	108.4
N1—C8—C1	122.00 (12)	N2—C16—C9	121.07 (12)
N1—C8—H8	119.0	N2—C16—H16	119.5
C1—C8—H8	119.0	C9—C16—H16	119.5
C6—C1—C2—C3	-0.3 (2)	C14—C9—C10—C11	0.2 (2)

C8—C1—C2—C3	179.76 (13)	C16—C9—C10—C11	−179.38 (13)
C1—C2—C3—C4	0.2 (2)	C9—C10—C11—C12	−0.2 (2)
C7—O1—C4—C3	179.08 (14)	C15—O4—C12—C11	−179.48 (15)
C7—O1—C4—C5	−2.05 (15)	C15—O4—C12—C13	0.38 (16)
C2—C3—C4—O1	178.64 (13)	C10—C11—C12—O4	−179.98 (14)
C2—C3—C4—C5	−0.1 (2)	C10—C11—C12—C13	0.2 (2)
C7—O2—C5—C6	−179.40 (14)	C15—O5—C13—C14	179.94 (14)
C7—O2—C5—C4	1.47 (15)	C15—O5—C13—C12	0.07 (17)
O1—C4—C5—C6	−178.80 (12)	O4—C12—C13—C14	179.83 (13)
C3—C4—C5—C6	0.1 (2)	C11—C12—C13—C14	−0.3 (2)
O1—C4—C5—O2	0.38 (16)	O4—C12—C13—O5	−0.29 (17)
C3—C4—C5—O2	179.33 (12)	C11—C12—C13—O5	179.58 (14)
O2—C5—C6—C1	−179.25 (13)	O5—C13—C14—C9	−179.49 (14)
C4—C5—C6—C1	−0.2 (2)	C12—C13—C14—C9	0.4 (2)
C2—C1—C6—C5	0.29 (19)	C10—C9—C14—C13	−0.3 (2)
C8—C1—C6—C5	−179.76 (12)	C16—C9—C14—C13	179.28 (13)
C5—O2—C7—O1	−2.71 (15)	C13—O5—C15—O4	0.16 (17)
C4—O1—C7—O2	2.92 (15)	C12—O4—C15—O5	−0.33 (16)
O3—N1—C8—C1	−179.96 (11)	O6—N2—C16—C9	178.85 (11)
C2—C1—C8—N1	−176.17 (13)	C10—C9—C16—N2	176.16 (13)
C6—C1—C8—N1	3.9 (2)	C14—C9—C16—N2	−3.5 (2)

Hydrogen-bond geometry (\AA , °)

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
O3—H3A···N2	0.87	1.93	2.7549 (16)	158
C7—H7B···O4 ⁱ	0.99	2.58	3.239 (2)	124
C8—H8···O6 ⁱⁱ	0.95	2.43	3.3754 (18)	173
O6—H6A···N1	0.87	1.97	2.7989 (17)	158
C15—H15A···O1 ⁱⁱⁱ	0.99	2.54	3.1775 (19)	122
C16—H16···O3 ^{iv}	0.95	2.59	3.5173 (18)	167

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