



Crystal structure and Hirshfeld surface analysis of 6,6'-(*(1E,1'E)*-{[1,4-phenylenebis(methylene)]bis(azanylylidene)}bis(methaneylylidene))bis(2-methoxyphenol)

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Received 1 November 2021

Accepted 15 December 2021

Edited by V. Jancik, Universidad Nacional Autónoma de México, México

Keywords: crystal structure; tetradentate salen-type; Schiff base; phenol-imine tautomer; Hirshfeld surface analysis.

CCDC reference: 2128953

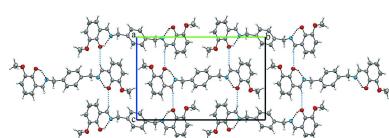
Supporting information: this article has supporting information at journals.iucr.org/e

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The Schiff base compound, $C_{24}H_{24}N_2O_4$, was synthesized by the interaction of 2-hydroxy-3-methoxy benzaldehyde and 1,4-benzene dimethanamine in ethanol, and crystallizes in the monoclinic space group $P2_1/n$ with $Z' = 0.5$. The molecule is not planar, the 1,4-diethylbenzene and the phenol rings are twisted with respect to each other, making a dihedral angle of $74.27(5)^\circ$. The molecular structure is stabilized by an $O-H\cdots N$ hydrogen bond, forming an $S(6)$ ring motif. In the crystal, molecules are linked by $C-H\cdots O$ hydrogen bonds, resulting in the formation of sheets parallel to the bc plane. A Hirshfeld surface analysis was undertaken to investigate the various intermolecular contacts controlling the supramolecular topology, suggesting the $H\cdots O$ (18%) contacts to be the most significant interactions, whereas the $H\cdots H$ (50.5%) and $C\cdots H$ (24.3%) interactions are less significant.

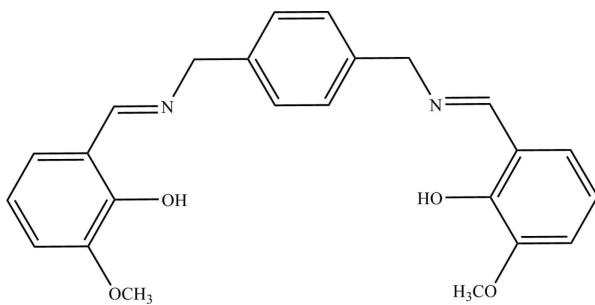
1. Chemical context

Schiff bases are used as pigments and dyes, catalysts, intermediates in organic synthesis, and as polymer stabilizers (Supuran *et al.*, 1996). In azomethine derivatives, the $C\equiv N$ linkage is essential for biological activity and several azomethines have been reported to possess remarkable antibacterial, antifungal, anticancer and diuretic activities (Gaur, 2003). Schiff bases having an azomethine group of general formula $C\equiv N-$ contain various substituted groups (Schiff, 1864). Of particular interest are the two different tautomeric structures for *o*-hydroxy Schiff bases, which are expressed as keto-amine and phenol-imine, with intrinsic $N-H\cdots O$ or $O-H\cdots N$ hydrogen bonds, (Filarowski *et al.*, 2004). There are many studies in the literature on the synthesis of Schiff bases and investigation of tautomeric structures. Phenol-imine and keto-amine tautomeric structures exhibit features of photochromism and thermochromism (Hadjoudis *et al.*, 2004). Tetradentate salen-type ligands have been used in almost all areas of coordination chemistry to prepare complexes that have catalytic and biological activity or which feature interesting structural, electrochemical or magnetic properties (Abd El-Hamid *et al.*, 2019). In this study, a symmetrical tetradentate Schiff base ligand bearing ONNO donor atoms,



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6,6'-((1E,1'E)-{[1,4-phenylenebis(methylene)]bis(azanylidene})bis(methanelylidene)}*)bis(2-methoxyphenol)* was synthesized by the interaction of 2-hydroxy-3-methoxybenzaldehyde and 1,4-benzene dimethanamine in ethanol and its crystal structure determined by single-crystal X-ray diffraction.



2. Structural commentary

The molecular structure of the title Schiff base derivative is illustrated in Fig. 1. The asymmetric unit of the title compound contains one-half of the centrosymmetric molecule ($Z' = 0.5$). There is an intramolecular $O_2-H_2\cdots N_1$ hydrogen bond (Table 1 and Fig. 1); this is a common feature also observed in related phenol-imine Schiff bases. It forms an *S*(6) ring motif and also induces the phenol ring and the Schiff base to be nearly coplanar, as indicated by the $C_6-C_8-N_1-C_9$ torsion angle of $178.54(13)^\circ$. The molecule is non-planar, the 1,4-diethylbenzene ring being inclined to the phenol ring by $74.27(5)^\circ$. The $C_7-C_6-C_8-N_1$ torsion angle [$3.8(2)^\circ$] further supports the co-planarity of the phenol ring and the Schiff base. The C_7-O_2 distance is $1.3438(17)$ Å, which is close to normal values reported for single C–O bonds in phenols and salicylideneamines (Kaştaş & Albayrak Kaştaş, 2019). The N_1-C_8 bond is short at $1.2717(17)$ Å, strongly indicating a $C=N$ double bond, while the long C_6-C_8 bond [$1.451(2)$ Å] implies a single bond. All of these data support the existence of the phenol-imine tautomer for the title compound in the crystalline state.

3. Supramolecular features

In the crystal, molecules are connected into sheets extending in the *bc* plane by $C_8-H_8\cdots O_2^i$ hydrogen bonds (Table 1; Fig. 2).

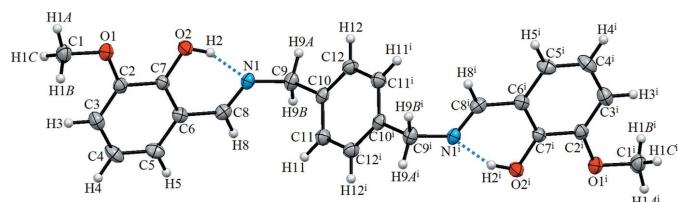


Figure 1

The molecular structure of the title compound with displacement ellipsoids drawn at the 40% probability level. Symmetry code: (i) $-x + 2, -y + 1, -z + 1$.

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O_2-H_2\cdots N_1$	0.86 (1)	1.79 (2)	2.5877 (18)	154 (2)
$C_8-H_8\cdots O_2^i$	0.93	2.51	3.410 (2)	162

Symmetry code: (i) $x + \frac{1}{2}, -y + \frac{1}{2}, z + \frac{1}{2}$.

4. Database survey

A search of the Cambridge Structural Database (CSD Version 5.42, update of May 2021; Groom *et al.*, 2016) for the (1,4-phenylene)bis(*N*-ethylideneimethanamine) moiety revealed some related structures. The most similar structures are 1,4-bis(2-pyridylmethylenaminomethyl)benzene (GOLJUN; Li *et al.*, 2009), 1,4-bis(3-pyridylmethylenaminomethyl)benzene (GOLJOH; He *et al.*, 2009) and 1,4-bis(3,5-di-*t*-butyl-2-hydroxybenzylideneaminomethyl)benzene (OCAPAK; Tooké *et al.*, 2004). In GOLJUN and GOLJOH, the molecules have similar shapes to the title compound. The C–N bond lengths [1.253 (2) Å in GOLJOH and 1.256 (2) Å in GOLJUN] are typical for an azomethine C=N bond and shorter than in the title compound [1.2717 (19) Å]. The torsion angles involving the C=C≡N–C units are $-177.26(11)^\circ$ and $115.21(13)^\circ$ in GOLJUN. These values are similar to those observed in the title compound. In OCAPAK, a *t*-butyl group is present, different from the title compound. In addition, there is an intramolecular O–H···N contact in the title compound. Similarly, in OCAPAK, the hydroxyl H atom is involved in an intramolecular O–H···N hydrogen bond, forming an *S*(6) ring motif as in the title compound. The length of intramolecular O–H···N hydrogen bond in OCAPAK is especially short [1.65 (2) Å] compared to that in the title compound [1.789 (15) Å].

5. Hirshfeld surface analysis

Hirshfeld surface analysis was used to analyse the various intermolecular interactions in the title compound, through mapping of the normalized contact distance (d_{norm}) using *CrystalExplorer17* (Turner *et al.*, 2017; Spackman & Jayatilaka, 2009). Hirshfeld surface analysis is a valuable tool for assessing the strength of intermolecular interactions, predicting the properties of a crystal and its potential applications (Al-Resayes *et al.*, 2020). The Hirshfeld surface was generated using a standard (high) surface resolution with the three-dimensional d_{norm} surface mapped over a fixed color scale of

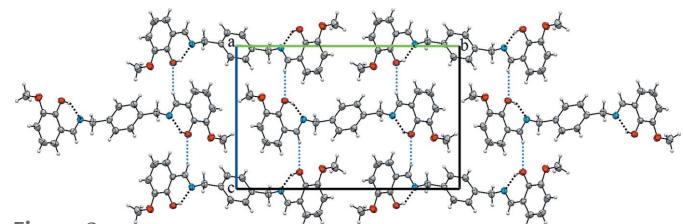


Figure 2

A view of the crystal packing of the title compound in a view parallel to the *bc* plane. C–H···O hydrogen bonds are shown as dashed blue lines.

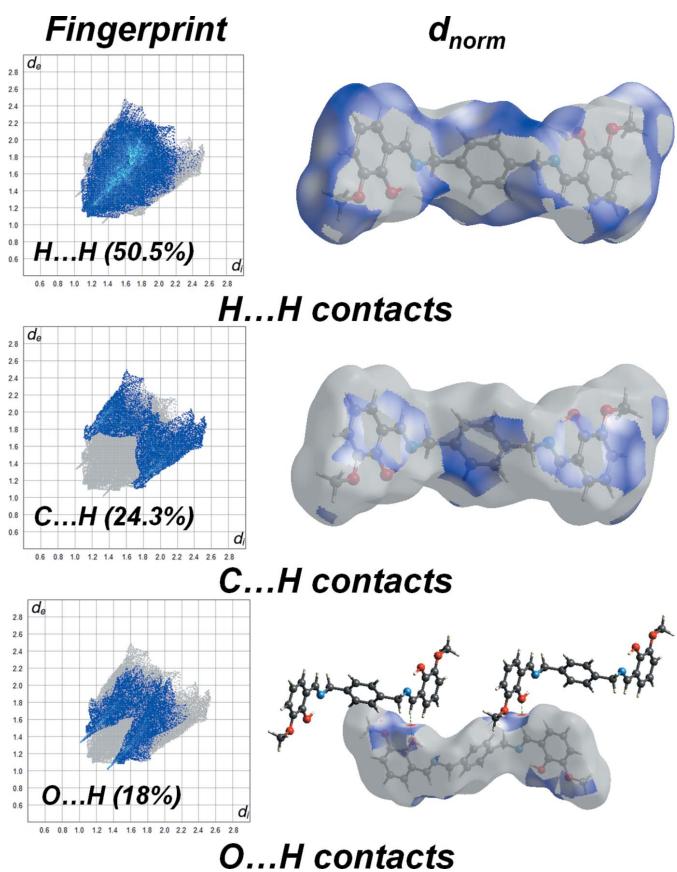


Figure 3

The Hirshfeld surface analysis of the title compound mapped with d_{norm} over -0.175 to 1.404 a.u. showing the $\text{C}-\text{H} \cdots \text{O}$ hydrogen-bonded contacts.

-0.175 (red) to 1.404 a.u. (blue). The packing of molecules is mainly dependent on $\text{H} \cdots \text{H}$ (50.5%) and $\text{C} \cdots \text{H}$ (24.3%) interactions and the significant $\text{C}-\text{H} \cdots \text{O}$ interactions (18%). Blue regions in the d_{norm} map indicate intermolecular interactions with distances longer than van der Waals radius sum of the interacting elements (Fig. 3). The $\text{C}-\text{H} \cdots \text{O}$ interactions, which appear as red spots in the d_{norm} map, have contact distances shorter than the sum of the van der Waals radii of the oxygen and hydrogen atoms

6. Synthesis and crystallization

0.0225 g (0.148 mmol) of 2-hydroxy-3-methoxy benzaldehyde was dissolved in 20 mL of ethanol and mixed with 0.0100 g (0.074 mmol) of 1,4-benzene dimethanamine dissolved in 20 mL of ethanol (Fig. 4). The reaction mixture was refluxed for 6 h and at the end of the reaction, the solution was allowed to cool. The yellow product obtained was washed with ether and crystallized in ethanol at room temperature (m.p. = 431–434 K, yield 85%).

7. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The O-bound H atom was located

Table 2
Experimental details.

Crystal data	$\text{C}_{24}\text{H}_{24}\text{N}_2\text{O}_4$
Chemical formula	404.45
M_r	Monoclinic, $P2_1/n$
Crystal system, space group	296
Temperature (K)	$4.7339 (10), 18.406 (4), 11.880 (2)$
a, b, c (Å)	$98.47 (3)$
β (°)	$1023.8 (4)$
V (Å 3)	Z
Radiation type	2
μ (mm $^{-1}$)	Mo $K\alpha$
Crystal size (mm)	0.09
	$0.61 \times 0.45 \times 0.23$
Data collection	
Diffractometer	Stoe IPDS 2
Absorption correction	Integration (<i>X-RED32</i> ; Stoe & Cie, 2002)
T_{\min}, T_{\max}	$0.963, 0.992$
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	$6766, 1863, 1315$
R_{int}	0.034
$(\sin \theta/\lambda)_{\text{max}}$ (Å $^{-1}$)	0.600
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	$0.037, 0.092, 1.01$
No. of reflections	1863
No. of parameters	140
No. of restraints	1
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å $^{-3}$)	$0.11, -0.09$

Computer programs: *X-AREA* and *X-RED* (Stoe & Cie, 2002), *SHELXT2014/5* (Sheldrick, 2015a), *SHELXL2018/3* (Sheldrick, 2015b) and *PLATON* (Spek, 2020).

in a difference-Fourier map and refined with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$ and a distance restraint. The C-bound H atoms

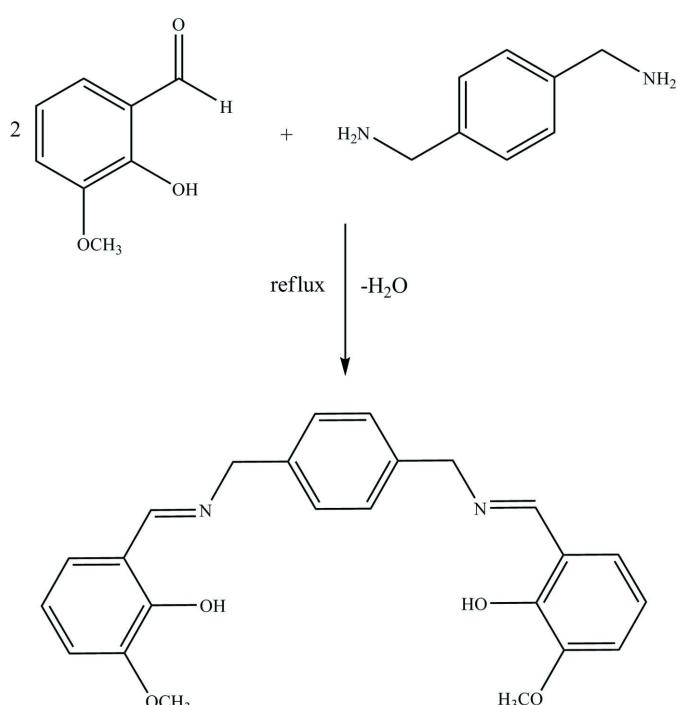


Figure 4
The synthesis of the title compound.

were positioned geometrically ($C-H = 0.93, 0.96$ and 0.97 \AA) and refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms and $1.2U_{\text{eq}}(\text{C})$ for other H atoms.

Acknowledgements

Author contributions are as follows. Conceptualization, SK, EA and ES; synthesis, SY and EA; writing (review and editing of the manuscript) SK and SY; formal analysis, SY, EA, ND and SK; crystal-structure determination, SY, SK and ND; validation, SY, EA and ES; project administration, EA, SY and SK.

Funding information

Funding for this research was provided by Samsun University under Project No. BAP-MÜF.5501.2020.004.

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supporting information

Acta Cryst. (2022). E78, 84-87 [https://doi.org/10.1107/S2056989021013347]

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Computing details

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA* (Stoe & Cie, 2002); data reduction: *X-RED* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXT2014/5* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2015b); molecular graphics: *PLATON* (Spek, 2020).

6,6'-(*(1E,1'E)*-{[1,4-Phenylenebis(methylene)]bis(azanylylidene)}bis(methaneylylidene))bis(2-methoxyphenol)

Crystal data

C ₂₄ H ₂₄ N ₂ O ₄	F(000) = 428
M _r = 404.45	D _x = 1.312 Mg m ⁻³
Monoclinic, P2 ₁ /n	Mo K α radiation, λ = 0.71073 Å
<i>a</i> = 4.7339 (10) Å	Cell parameters from 7423 reflections
<i>b</i> = 18.406 (4) Å	θ = 1.7–31.5°
<i>c</i> = 11.880 (2) Å	μ = 0.09 mm ⁻¹
β = 98.47 (3)°	<i>T</i> = 296 K
<i>V</i> = 1023.8 (4) Å ³	Plate, yellow
Z = 2	0.61 × 0.45 × 0.23 mm

Data collection

Stoe IPDS 2	6766 measured reflections
diffractometer	1863 independent reflections
Radiation source: sealed X-ray tube, 12 x 0.4	1315 reflections with $I > 2\sigma(I)$
mm long-fine focus	R_{int} = 0.034
Detector resolution: 6.67 pixels mm ⁻¹	$\theta_{\text{max}} = 25.3^\circ$, $\theta_{\text{min}} = 2.1^\circ$
rotation method scans	$h = -5 \rightarrow 5$
Absorption correction: integration	$k = -22 \rightarrow 22$
(X-RED32; Stoe & Cie, 2002)	$l = -13 \rightarrow 14$
$T_{\text{min}} = 0.963$, $T_{\text{max}} = 0.992$	

Refinement

Refinement on F^2	Primary atom site location: structure-invariant direct methods
Least-squares matrix: full	Secondary atom site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.037$	Hydrogen site location: mixed
$wR(F^2) = 0.092$	H atoms treated by a mixture of independent and constrained refinement
$S = 1.01$	
1863 reflections	
140 parameters	
1 restraint	

$$w = 1/[\sigma^2(F_o^2) + (0.049P)^2]$$

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

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

$$\Delta\rho_{\max} = 0.11 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.09 \text{ e \AA}^{-3}$$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O2	0.6966 (3)	0.21666 (6)	0.38262 (9)	0.0633 (3)
H2	0.817 (4)	0.2501 (10)	0.4060 (16)	0.095*
O1	0.3093 (3)	0.11352 (6)	0.35977 (10)	0.0722 (4)
N1	1.0411 (3)	0.30049 (6)	0.51303 (11)	0.0550 (3)
C7	0.6181 (3)	0.18967 (7)	0.47846 (12)	0.0499 (4)
C6	0.7362 (3)	0.21470 (8)	0.58611 (12)	0.0528 (4)
C10	1.1273 (3)	0.43140 (8)	0.51623 (13)	0.0533 (4)
C8	0.9571 (3)	0.26998 (8)	0.59825 (13)	0.0561 (4)
H8	1.040345	0.283593	0.671014	0.067*
C2	0.4103 (3)	0.13480 (8)	0.46848 (13)	0.0560 (4)
C12	1.1668 (4)	0.47353 (9)	0.42431 (14)	0.0620 (4)
H12	1.279589	0.456251	0.372334	0.074*
C9	1.2603 (4)	0.35690 (8)	0.53440 (15)	0.0629 (4)
H9A	1.400552	0.350226	0.483419	0.075*
H9B	1.357488	0.352833	0.611939	0.075*
C11	0.9593 (4)	0.45861 (9)	0.59194 (14)	0.0634 (4)
H11	0.930241	0.431002	0.654796	0.076*
C5	0.6416 (4)	0.18568 (10)	0.68255 (14)	0.0697 (5)
H5	0.719087	0.202247	0.754451	0.084*
C3	0.3230 (4)	0.10710 (9)	0.56514 (17)	0.0702 (5)
H3	0.186333	0.070445	0.558980	0.084*
C4	0.4365 (4)	0.13326 (11)	0.67187 (16)	0.0786 (5)
H4	0.372061	0.114863	0.736433	0.094*
C1	0.0961 (4)	0.05885 (10)	0.34579 (18)	0.0816 (6)
H1A	0.044148	0.048311	0.266297	0.122*
H1B	-0.068976	0.075560	0.376457	0.122*
H1C	0.168648	0.015654	0.385164	0.122*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O2	0.0715 (8)	0.0674 (7)	0.0513 (6)	-0.0180 (6)	0.0096 (5)	0.0000 (5)
O1	0.0729 (8)	0.0703 (7)	0.0742 (8)	-0.0230 (6)	0.0139 (6)	-0.0124 (6)
N1	0.0550 (8)	0.0453 (7)	0.0626 (8)	0.0025 (6)	0.0021 (6)	-0.0016 (6)
C7	0.0525 (8)	0.0459 (7)	0.0519 (8)	0.0048 (7)	0.0102 (7)	0.0047 (6)
C6	0.0519 (9)	0.0532 (8)	0.0523 (8)	0.0089 (7)	0.0044 (7)	0.0069 (6)

C10	0.0448 (9)	0.0483 (8)	0.0641 (9)	-0.0039 (6)	-0.0012 (7)	-0.0083 (7)
C8	0.0562 (9)	0.0560 (9)	0.0532 (9)	0.0104 (7)	-0.0019 (7)	-0.0021 (7)
C2	0.0540 (9)	0.0503 (8)	0.0644 (10)	0.0024 (7)	0.0110 (8)	0.0020 (7)
C12	0.0607 (10)	0.0598 (9)	0.0670 (10)	0.0024 (8)	0.0146 (8)	-0.0071 (8)
C9	0.0535 (9)	0.0523 (9)	0.0805 (11)	0.0008 (7)	0.0022 (8)	-0.0051 (8)
C11	0.0675 (11)	0.0570 (9)	0.0665 (10)	0.0020 (8)	0.0128 (9)	0.0029 (7)
C5	0.0690 (11)	0.0868 (12)	0.0527 (9)	0.0083 (10)	0.0066 (8)	0.0126 (8)
C3	0.0629 (11)	0.0631 (10)	0.0865 (13)	-0.0018 (8)	0.0170 (10)	0.0180 (9)
C4	0.0742 (12)	0.0940 (14)	0.0699 (12)	0.0043 (11)	0.0188 (10)	0.0333 (10)
C1	0.0695 (12)	0.0710 (12)	0.1064 (15)	-0.0183 (9)	0.0202 (11)	-0.0169 (10)

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

O2—C7	1.3438 (17)	C2—C3	1.375 (2)
O2—H2	0.858 (13)	C12—C11 ⁱ	1.386 (2)
O1—C2	1.3668 (19)	C12—H12	0.9300
O1—C1	1.418 (2)	C9—H9A	0.9700
N1—C8	1.2717 (19)	C9—H9B	0.9700
N1—C9	1.463 (2)	C11—H11	0.9300
C7—C6	1.396 (2)	C5—C4	1.361 (3)
C7—C2	1.403 (2)	C5—H5	0.9300
C6—C5	1.396 (2)	C3—C4	1.388 (3)
C6—C8	1.451 (2)	C3—H3	0.9300
C10—C12	1.374 (2)	C4—H4	0.9300
C10—C11	1.379 (2)	C1—H1A	0.9600
C10—C9	1.511 (2)	C1—H1B	0.9600
C8—H8	0.9300	C1—H1C	0.9600
C7—O2—H2	104.1 (13)	C10—C9—H9A	109.6
C2—O1—C1	117.26 (13)	N1—C9—H9B	109.6
C8—N1—C9	118.17 (14)	C10—C9—H9B	109.6
O2—C7—C6	122.05 (13)	H9A—C9—H9B	108.1
O2—C7—C2	118.20 (14)	C10—C11—C12 ⁱ	120.95 (15)
C6—C7—C2	119.75 (13)	C10—C11—H11	119.5
C7—C6—C5	119.45 (15)	C12 ⁱ —C11—H11	119.5
C7—C6—C8	120.53 (13)	C4—C5—C6	120.35 (17)
C5—C6—C8	120.02 (15)	C4—C5—H5	119.8
C12—C10—C11	118.18 (14)	C6—C5—H5	119.8
C12—C10—C9	121.52 (14)	C2—C3—C4	120.71 (17)
C11—C10—C9	120.30 (15)	C2—C3—H3	119.6
N1—C8—C6	122.40 (14)	C4—C3—H3	119.6
N1—C8—H8	118.8	C5—C4—C3	120.38 (16)
C6—C8—H8	118.8	C5—C4—H4	119.8
O1—C2—C3	125.21 (15)	C3—C4—H4	119.8
O1—C2—C7	115.45 (13)	O1—C1—H1A	109.5
C3—C2—C7	119.34 (16)	O1—C1—H1B	109.5
C10—C12—C11 ⁱ	120.88 (14)	H1A—C1—H1B	109.5
C10—C12—H12	119.6	O1—C1—H1C	109.5

C11 ⁱ —C12—H12	119.6	H1A—C1—H1C	109.5
N1—C9—C10	110.42 (13)	H1B—C1—H1C	109.5
N1—C9—H9A	109.6		
O2—C7—C6—C5	178.60 (14)	C11—C10—C12—C11 ⁱ	-0.1 (3)
C2—C7—C6—C5	-1.3 (2)	C9—C10—C12—C11 ⁱ	179.25 (15)
O2—C7—C6—C8	-1.9 (2)	C8—N1—C9—C10	-102.38 (16)
C2—C7—C6—C8	178.27 (13)	C12—C10—C9—N1	-108.69 (17)
C9—N1—C8—C6	178.54 (13)	C11—C10—C9—N1	70.69 (19)
C7—C6—C8—N1	3.8 (2)	C12—C10—C11—C12 ⁱ	0.1 (3)
C5—C6—C8—N1	-176.65 (14)	C9—C10—C11—C12 ⁱ	-179.26 (15)
C1—O1—C2—C3	0.4 (2)	C7—C6—C5—C4	0.2 (2)
C1—O1—C2—C7	-179.17 (14)	C8—C6—C5—C4	-179.33 (15)
O2—C7—C2—O1	0.7 (2)	O1—C2—C3—C4	-179.08 (16)
C6—C7—C2—O1	-179.47 (13)	C7—C2—C3—C4	0.5 (2)
O2—C7—C2—C3	-178.96 (14)	C6—C5—C4—C3	1.2 (3)
C6—C7—C2—C3	0.9 (2)	C2—C3—C4—C5	-1.6 (3)

Symmetry code: (i) $-x+2, -y+1, -z+1$.

Hydrogen-bond geometry (\AA , °)

$D\text{—H}\cdots A$	$D\text{—H}$	$\text{H}\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
O2—H2···N1	0.86 (1)	1.79 (2)	2.5877 (18)	154 (2)
C8—H8···O2 ⁱⁱ	0.93	2.51	3.410 (2)	162

Symmetry code: (ii) $x+1/2, -y+1/2, z+1/2$.