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### Polar crystal of vanillylformamide through replacement of the alkene by an isosteric formamide group

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Vanillylformamide [systematic name: *N*-(4-hydroxy-3-methoxybenzyl)formamide],  $C_9H_{11}NO_3$ , (**II**), has been synthesized from vanillylamine hydrochloride and studied by single-crystal X-ray diffraction. Compound (**II**) and the well known biologically active eugenol compound (**I**) can be considered to be 'isosteres' of each other, since they share comparable molecular shape and volume. The product (**II**) crystallizes in the space group *P*1. In the crystal, the vanillylformamide molecules are linked mainly by N-H···O, O-H···O and  $Csp^2-H···O$  hydrogen bonds, forming infinite two-dimensional polar sheets. These two-dimensional layers pack in a parallel fashion, constructing a polar three-dimensional network. Except for van der Waals forces and weak  $Csp^3-$ H···O hydrogen bonds, there are no significant intermolecular interactions between the layers. A Cambridge Structural Database search revealed that vanillylamide-related crystals are scarce.



### Structure description

Eugenol is a natural molecule that exhibits versatile properties useful in various domains. Indeed, there is increasing interest from the scientific and industrial community in eugenol-based polymers (Miao *et al.* 2017; Guzmán *et al.*, 2017; Chen *et al.*, 2017; Modjinou *et al.*, 2016; Wan *et al.*, 2016*a,b*; Deng *et al.*, 2015). Moreover, this bioactive compound has high potential as a therapeutic agent since it has antiparasitic, antiviral, antibacterial, antifungal, anticancer, antioxidant and anti-inflammatory activities (Raja *et al.*, 2015; Khalil *et al.*, 2017). On the other hand, to the best of our knowledge, no study on the bioisosteres of eugenol has been undertaken. The definition of bioisosterism has been



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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdots A$
$N1-H1\cdots O2^{i}$	0.88	2.05	2.910 (4)	167
O3−H3···O1	0.96 (3)	2.15(7)	2.672 (4)	113 (5)
C6-H6···O3 <sup>ii</sup>	0.95	2.54	3.391 (5)	149
O3−H3···O2 <sup>iii</sup>	0.96 (3)	1.90 (4)	2.771 (4)	149 (6)
$C5-H5B\cdots O3^{iv}$	0.99	2.60	3.374 (5)	135
$C5-H5A\cdots O3^{v}$	0.99	2.70	3.676 (5)	167

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

broadened by Burger (1991) as 'Compounds or groups that possess near-equal molecular shapes and volumes, approximately the same distribution of electrons, and which exhibit similar physical properties'. Thus, chemical modification of lead compounds represents a rational approach in drug design (Patani et al., 1996). Alkenes and amides are isosteric since they are both planar and possess two  $sp^2$ -hybridized atoms in the main chain (Choudhary et al., 2011). In this context, effort has been focused in our group to determine the crystal structure of the vanillylformamide (II), which mimics eugenol (I) (Fig. 1) and could have some biological interest. We were also pleased that the title compound crystallized in a polar space group, since several functional properties of advanced materials (piezoelectricity, pyroelectricity, ferroelectricity, second harmonic generation, and electro-optic response) are only allowed or significantly enhanced in polar crystal structures (Centore et al., 2012, 2016; Takahashi et al., 2016).

The molecular structure of the title compound (II) is shown in Fig. 2. All bond lengths and angles are within normal ranges. Although the O3-H3···O1 angle [113 (5)°] is far from linear, we can consider that the phenol and methoxy group are partners in an intramolecular hydrogen bond (Hunt *et al.*, 2005). The *cis/trans* conformational equilibrium of the formamide group is fixed in the solid state in the *trans* conformation, as can be confirmed by the torsion angle C5-N1-C6-O2 [1.3 (6)°].

In the crystal, the vanillylformamide molecules are linked by a series of classical hydrogen bonds (N1-H1···O2, O3-H3···O1, O3-H3···O2; Table 1) and non-conventional  $Csp^2$ -H···O hydrogen bonds (C6-H6···O3; Table 1), forming infinite two-dimensional polar sheets [Fig. 3(*a*)]



Eugenol (I) and the title compound (II).



Figure 2

The molecular structure of compound (II), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are shown as fixed-size spheres of 0.30 Å.

parallel to  $(01\overline{1})$ . Moreover,  $C6\cdots O1$  close contacts stabilize this architecture. The stacking pattern reveals that each layer is oriented in the same direction [Fig. 3(*b*)], generating a complete three-dimensional polar network. The main intermolecular contacts between layers [Fig. 2(*c*)] consist of van der Waals forces (C1 $\cdots$ H7 and H1 $B\cdots$ H7) and weak Cs $p^3$ -H $\cdots$ O hydrogen bonds (C5-H5 $B\cdots$ O3 and C5-H5 $A\cdots$ O3 of neighbouring molecules).





(a) Main hydrogen bonds and dipole-dipole interactions inside the polar sheets of vanillylformamide (II). (b) Stacking of the polar sheets along the *a* axis. (c) Weak interactions between the polar sheets.

Table 2Experimental details.

$ \begin{array}{llllllllllllllllllllllllllllllllllll$	Crystal data	
$\begin{array}{llllllllllllllllllllllllllllllllllll$	Chemical formula	C <sub>9</sub> H <sub>11</sub> NO <sub>3</sub>
$\begin{array}{llllllllllllllllllllllllllllllllllll$	$M_{\rm r}$	181.19
Temperature (K)173 $a, b, c$ (Å)173 $a, b, c$ (Å)4.8011 (2), 6.5522 (3), 7.5052 (3) $\alpha, \beta, \gamma$ (°)93.618 (2), 107.044 (2), 95.658 (2) $V$ (Å <sup>3</sup> )223.58 (2) $Z$ 1Radiation typeCu K $\alpha$ $\mu$ (mm <sup>-1</sup> )0.85Crystal size (mm)0.32 × 0.16 × 0.12Data collectionBruker APEXIIAbsorption correctionMulti-scan (SADABS; Bruker, 2012) $T_{min}, T_{max}$ 0.574, 0.753No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections0.051 $R_{int}$ 0.051 $(\sin \theta/\lambda)_{max}$ (Å <sup>-1</sup> )0.609Refinement1206No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	Crystal system, space group	Triclinic, P1
$a, b, c$ (Å) $4.8011$ (2), $6.5522$ (3), $7.5052$ (3) $\alpha, \beta, \gamma$ (°) $93.618$ (2), $107.044$ (2), $95.658$ (2) $V$ (Å <sup>3</sup> ) $223.58$ (2) $Z$ 1         Radiation type       Cu K $\alpha$ $\mu$ (mm <sup>-1</sup> )       0.85         Crystal size (mm) $0.32 \times 0.16 \times 0.12$ Data collection       Bruker APEXII         Absorption correction       Multi-scan (SADABS; Bruker, 2012) $T_{min}, T_{max}$ $0.574, 0.753$ No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections $0.051$ $R_{int}$ $0.051$ (sin $\theta/\lambda)_{max}$ (Å <sup>-1</sup> ) $0.609$ Refinement $R[F^2 > 2\sigma(F^2)], wR(F^2), S$ No. of restraints       4         H-atom treatment       H atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> ) $0.25, -0.23$	Temperature (K)	173
$\alpha, \beta, \gamma$ (°)93.618 (2), 107.044 (2), 95.658 (2) $V(Å^3)$ 223.58 (2) $Z$ 1Radiation typeCu $K\alpha$ $\mu$ (mm <sup>-1</sup> )0.85Crystal size (mm)0.32 × 0.16 × 0.12Data collectionBruker APEXIIDiffractometerBruker APEXIIAbsorption correctionMulti-scan (SADABS; Bruker, 2012) $T_{min}, T_{max}$ 0.574, 0.753No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections0.051 $R_{int}$ 0.051 $(\sin \theta/\lambda)_{max}$ (Å <sup>-1</sup> )0.609Refinement1206No. of restraints4H-atom treatment4H-atom treatment4 $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	<i>a</i> , <i>b</i> , <i>c</i> (Å)	4.8011 (2), 6.5522 (3), 7.5052 (3)
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	$\alpha, \beta, \gamma$ (°)	93.618 (2), 107.044 (2), 95.658 (2)
$\begin{array}{llllllllllllllllllllllllllllllllllll$	$V(\text{\AA}^3)$	223.58 (2)
Radiation typeCu Kαμ (mm <sup>-1</sup> )0.85Crystal size (mm)0.32 × 0.16 × 0.12Data collection0.32 × 0.16 × 0.12Data collectionBruker APEXIIAbsorption correctionMulti-scan (SADABS; Bruker, 2012) $T_{min}, T_{max}$ 0.574, 0.753No. of measured, independent and observed [ $I > 2\sigma(I)$ ] reflections0.051 $R_{int}$ 0.051 $(\sin \theta   \lambda)_{max}$ (Å <sup>-1</sup> )0.609Refinement1206 $R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , S0.052, 0.154, 1.12No. of reflections1206No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	Ζ	1
μ (mm <sup>-1</sup> )0.85Crystal size (mm)0.32 × 0.16 × 0.12Data collectionBruker APEXIIDiffractometerBruker APEXIIAbsorption correctionMulti-scan (SADABS; Bruker, 2012) $T_{min}, T_{max}$ 0.574, 0.753No. of measured, independent and observed [ $I > 2\sigma(I)$ ] reflections0.051 $R_{int}$ 0.051 $(\sin \theta/\lambda)_{max}$ (Å <sup>-1</sup> )0.609Refinement $R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , S0.052, 0.154, 1.12No. of reflections1206No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	Radiation type	Cu Ka
$\begin{array}{llllllllllllllllllllllllllllllllllll$	$\mu \ (\mathrm{mm}^{-1})$	0.85
Data collectionBruker APEXIIDiffractometerBruker APEXIIAbsorption correctionMulti-scan (SADABS; Bruker, 2012) $T_{min}, T_{max}$ 0.574, 0.753No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections0.574, 0.753 $R_{int}$ 0.051 $(\sin \theta/\lambda)_{max}$ (Å <sup>-1</sup> )0.609Refinement $R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , SNo. of parameters123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	Crystal size (mm)	$0.32 \times 0.16 \times 0.12$
$\begin{array}{llllllllllllllllllllllllllllllllllll$		
$\begin{array}{llllllllllllllllllllllllllllllllllll$	Data collection	
Absorption correctionMulti-scan (SADABS; Bruker, 2012) $T_{\min}, T_{\max}$ 0.574, 0.753No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections0.574, 0.753 $R_{int}$ 0.574, 0.753 $(\sin \theta/\lambda)_{max}$ (Å <sup>-1</sup> )0.051Refinement $R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , S0.052, 0.154, 1.12No. of reflections1206No. of parameters123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{\max}, \Delta \rho_{\min}$ (e Å <sup>-3</sup> )0.25, -0.23	Diffractometer	Bruker APEXII
$\begin{array}{ll} T_{\min}, T_{\max} & 0.574, 0.753 \\ \text{No. of measured, independent and observed } [I > 2\sigma(I)] \text{ reflections} \\ R_{\text{int}} & 0.051 \\ (\sin \theta/\lambda)_{\max} (\text{\AA}^{-1}) & 0.609 \\ \end{array}$ Refinement $\begin{array}{ll} R[F^2 > 2\sigma(F^2)], wR(F^2), S & 0.052, 0.154, 1.12 \\ \text{No. of reflections} & 1206 \\ \text{No. of parameters} & 123 \\ \text{No. of restraints} & 4 \\ \text{H-atom treatment} & H \text{ atoms treated by a mixture of independent and constrained refinement} \\ \Delta\rho_{\max}, \Delta\rho_{\min} (\text{e} \text{\AA}^{-3}) & 0.25, -0.23 \\ \end{array}$	Absorption correction	Multi-scan ( <i>SADABS</i> ; Bruker, 2012)
No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections2726, 1206, 1139 $R_{int}$ 0.051 $(\sin \theta/\lambda)_{max}$ (Å <sup>-1</sup> )0.609Refinement $R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , S $No.$ of reflections1206No. of reflections123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}$ , $\Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	$T_{\min}, T_{\max}$	0.574, 0.753
observed $[I > 2\sigma(I)]$ reflections $R_{int}$ 0.051 $(\sin \theta   \lambda \rangle_{max} (\AA^{-1})$ 0.609Refinement $R[F^2 > 2\sigma(F^2)], wR(F^2), S$ No. of reflections1206No. of parameters123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	No. of measured, independent and	2726, 1206, 1139
$R_{int}$ 0.051 $(\sin \theta / \lambda)_{max}$ $(Å^{-1})$ 0.609Refinement $R[F^2 > 2\sigma(F^2)], wR(F^2), S$ 0.052, 0.154, 1.12No. of reflections1206No. of parameters123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	observed $[I > 2\sigma(I)]$ reflections	
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Refinement $R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , S0.052, 0.154, 1.12No. of reflections1206No. of parameters123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}$ , $\Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	$(\sin \theta / \lambda)_{\text{max}} (\text{\AA}^{-1})$	0.609
Refinement $R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , S0.052, 0.154, 1.12No. of reflections1206No. of parameters123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}$ , $\Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23		
$\begin{split} R[F^2 > 2\sigma(F^2)], & R(F^2), S & 0.052, 0.154, 1.12 \\ \text{No. of reflections} & 1206 \\ \text{No. of parameters} & 123 \\ \text{No. of restraints} & 4 \\ \text{H-atom treatment} & H \text{ atoms treated by a mixture of } \\ & \text{independent and constrained} \\ & \alpha\rho_{\text{max}}, \Delta\rho_{\text{min}} \text{ (e } \text{\AA}^{-3} \text{)} & 0.25, -0.23 \end{split}$	Refinement	
No. of reflections1206No. of parameters123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.052, 0.154, 1.12
No. of parameters123No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	No. of reflections	1206
No. of restraints4H-atom treatmentH atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max}, \Delta \rho_{min}$ (e Å <sup>-3</sup> )0.25, -0.23	No. of parameters	123
H-atom treatment H atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{\text{max}}, \Delta \rho_{\text{min}}$ (e Å <sup>-3</sup> ) 0.25, -0.23	No. of restraints	4
independent and constrained refinement $\Delta \rho_{\text{max}}, \Delta \rho_{\text{min}}$ (e Å <sup>-3</sup> ) 0.25, -0.23	H-atom treatment	H atoms treated by a mixture of
$\Delta \rho_{\text{max}}, \Delta \rho_{\text{min}} \text{ (e Å}^{-3}\text{)}$ 0.25, -0.23		independent and constrained refinement
	$\Delta \rho_{\rm max},  \Delta \rho_{\rm min} \ ({\rm e} \ {\rm \AA}^{-3})$	0.25, -0.23

Computer programs: *APEX2* (Bruker, 2012), *SAINT* (Bruker, 2012), *SORTAV* (Blessing, 1995), *SHELXS* (Sheldrick, 2008), *SHELXL2016/6* (Sheldrick, 2015), *ORTEP* for Windows (Farrugia, 2012) and *Mercury* (Macrae *et al.*, 2008), *WinGX* (Farrugia, 2012) and *publCIF* (Westrip, 2010).

A search of the Cambridge Structural Database (CSD, Version 5.38, update May 2017; Groom *et al.*, 2016) indicated that 11 vanillylamide derivatives have been reported [FABVAF and FABVEJ (Oliver *et al.*, 1985); FABVAF01 (David *et al.*, 1998); FOSXOB (Winkler *et al.*, 2009); FOWTUH (Xia *et al.*, 2009); FOWTUH01 and KUTMAO01 (Wang *et al.*, 2010); KUTMAO (Huang *et al.*, 2010); QUZKOM (Xia *et al.*, 2010); SOFTEN (Zhang *et al.*, 2008); SOFTEN01 (Zhang & Cai, 2008)]. However, there was no structural report on vanillylamide analogues containing small amide units (such as a formamide) that could be considered as true isosteres of eugenol.

### Synthesis and crystallization

Vanillylformamide (II) was already characterized in the literature through NMR spectroscopy (Baldessari *et al.*, 1987). However, this molecule has no known crystal structure.

Compound (II): To a solution of 4-nitrophenylformate (575 mg, 3.44 mmol) in ethyl acetate (10 ml) at room temperature was added potassium carbonate (715 mg, 5.17 mmol) and vanillylamine hydrochloride (717 mg, 3.78 mmol) under an argon atmosphere. After 1 h of stirring at room temperature, a catalytic amount of water (30  $\mu$ l) was added, the resulting mixture was stirred for an additional 3 h.

The reaction was followed by TLC (30/70 acetone/DCM). The reaction mixture was poured in HCl 0.1M (15 ml) and then extracted twice with ethyl acetate (2 × 15 ml). Hexane (50 ml) was added to the combined organic layers and the resulting organic phase was then filtrated directly through a silica gel pad, eluting with 100 ml of acetone/DCM (5:95) and 100 ml of acetone/DCM (30:70), to yield compound (**II**) as a yellowish white crystalline powder (485 mg, 78%). Single crystals suitable for X-ray diffraction were prepared by slow evaporation of an ether/acetone/hexane (70:5:25) solution of (**II**) at room temperature.

 $R_{f} = 0.37$  (acetone/DCM 30:70); <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>, p.p.m.): 8.20 (*s*, 1H), 6.86–6.71 (*m*, 3H), 6.03 (*m*, 1H), 5.86 (*s*, 1H), 4.37 (*d*, J = 5.84 Hz, 2H), 3.85 (*s*, 3H).

### Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The absolute structure could not be determined reliably from the collected diffraction data.

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# full crystallographic data

IUCrData (2018). 3, x181630 [https://doi.org/10.1107/S2414314618016309]

# Polar crystal of vanillylformamide through replacement of the alkene by an isosteric formamide group

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(11)

Crystal data C<sub>9</sub>H<sub>11</sub>NO<sub>3</sub>  $M_r = 181.19$ Triclinic, P1 Hall symbol: P1 a = 4.8011 (2) Å b = 6.5522 (3) Å c = 7.5052 (3) Å a = 93.618 (2)°  $\beta = 107.044$  (2)°  $\gamma = 95.658$  (2)° V = 223.58 (2) Å<sup>3</sup>

### Data collection

Bruker APEXII diffractometer Radiation source: sealed x-ray tube Graphite monochromator  $\varphi$  or  $\omega$  oscillation scans Absorption correction: multi-scan (SADABS; Bruker, 2012)  $T_{\min} = 0.574, T_{\max} = 0.753$ 

### Refinement

Refinement on  $F^2$ Least-squares matrix: full  $R[F^2 > 2\sigma(F^2)] = 0.052$  $wR(F^2) = 0.154$ S = 1.121206 reflections 123 parameters 4 restraints 0 constraints

### Special details

Z = 1 F(000) = 96  $D_x = 1.346 \text{ Mg m}^{-3}$ Cu K $\alpha$  radiation,  $\lambda = 1.54178 \text{ Å}$ Cell parameters from 2759 reflections  $\theta = 6.2-70.7^{\circ}$   $\mu = 0.85 \text{ mm}^{-1}$  T = 173 KPrism, colorless  $0.32 \times 0.16 \times 0.12 \text{ mm}$ 

2726 measured reflections 1206 independent reflections 1139 reflections with  $I > 2\sigma(I)$   $R_{int} = 0.051$   $\theta_{max} = 69.8^{\circ}, \ \theta_{min} = 6.2^{\circ}$   $h = -5 \rightarrow 5$   $k = -7 \rightarrow 7$  $l = -9 \rightarrow 8$ 

Hydrogen site location: mixed H atoms treated by a mixture of independent and constrained refinement  $w = 1/[\sigma^2(F_o^2) + (0.0912P)^2 + 0.0525P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{max} < 0.001$  $\Delta\rho_{max} = 0.25 \text{ e } \text{Å}^{-3}$  $\Delta\rho_{min} = -0.23 \text{ e } \text{Å}^{-3}$ 

**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**. Hydrogen atoms bound to carbon atoms were positioned with idealized geometry and refined isotropically using a riding model, with  $U_{iso}(H) = 1.2 U_{eq}(C)$  and C—H = 0.95 Å for aromatic,  $U_{iso}(H) = 1.2 U_{eq}(C)$  and C—H = 0.99 Å for methylene and  $U_{iso}(H) = 1.5 U_{eq}(C)$  and C—H = 0.98 Å for methyl groups. The N-bound H atom was placed in idealized position with N—H = 0.88 Å and refined in riding mode with  $U_{iso}(H) = 1.2 U_{eq}(N)$ . The hydroxyl H atom was refined independently in isotropic mode, with an O—H distance restraint of 0.96 (2) Å.

	x	у	Ζ	$U_{\rm iso}$ */ $U_{\rm eq}$
O1	0.7872 (8)	0.7526 (5)	0.5754 (4)	0.0436 (9)
O2	1.0943 (7)	-0.0646 (5)	-0.0270 (5)	0.0376 (8)
O3	1.2189 (6)	0.5811 (5)	0.8054 (4)	0.0346 (8)
N1	0.6936 (8)	0.0863 (6)	-0.0091 (5)	0.0313 (8)
H1	0.506646	0.0619	-0.016887	0.038*
C1	0.5812 (13)	0.8622 (8)	0.4558 (8)	0.0457 (12)
H1A	0.673811	0.934512	0.373509	0.069*
H1B	0.513623	0.962513	0.531259	0.069*
H1C	0.413277	0.765528	0.379819	0.069*
C2	0.8948 (9)	0.6005 (6)	0.4953 (6)	0.0300 (10)
C3	0.8016 (9)	0.5279 (6)	0.3069 (6)	0.0313 (9)
H3A	0.648153	0.586589	0.222842	0.038*
C4	0.9284 (9)	0.3719 (6)	0.2397 (6)	0.0296 (9)
C5	0.8312 (9)	0.2985 (7)	0.0327 (6)	0.0330 (10)
H5A	0.690774	0.388824	-0.035372	0.04*
H5B	1.004057	0.312362	-0.014027	0.04*
C6	0.8314 (9)	-0.0728 (6)	-0.0364 (6)	0.0318 (9)
H6	0.718873	-0.204814	-0.065598	0.038*
C7	1.1468 (11)	0.2848 (7)	0.3628 (6)	0.0387 (11)
H7	1.235762	0.177733	0.317846	0.046*
C8	1.2385 (10)	0.3531 (6)	0.5538 (6)	0.0382 (11)
H8	1.385838	0.289743	0.638179	0.046*
C9	1.1167 (9)	0.5113 (6)	0.6200 (5)	0.0295 (10)
Н3	1.144 (16)	0.711 (7)	0.817 (10)	0.061 (18)*

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters  $(Å^2)$ 

Atomic displacement parameters  $(Å^2)$ 

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
01	0.054 (2)	0.0452 (19)	0.0316 (18)	0.0282 (16)	0.0062 (14)	-0.0004 (14)
O2	0.0287 (17)	0.0393 (17)	0.0418 (19)	0.0016 (12)	0.0094 (13)	-0.0082 (14)
03	0.0383 (17)	0.0347 (16)	0.0260 (15)	0.0064 (12)	0.0027 (12)	-0.0021 (12)
N1	0.0234 (16)	0.0360 (18)	0.0309 (19)	0.0017 (13)	0.0047 (14)	-0.0041 (14)
C1	0.054 (3)	0.047 (3)	0.038 (3)	0.024 (2)	0.011 (2)	0.009 (2)
C2	0.033 (2)	0.027 (2)	0.030(2)	0.0017 (17)	0.0105 (19)	0.0021 (17)
C3	0.032 (2)	0.032 (2)	0.026 (2)	0.0069 (17)	0.0025 (17)	0.0053 (16)
C4	0.028 (2)	0.030 (2)	0.029 (2)	0.0017 (16)	0.0052 (17)	0.0032 (17)
C5	0.038 (2)	0.035 (2)	0.024 (2)	0.0041 (17)	0.0066 (18)	0.0056 (16)
C6	0.035 (2)	0.033 (2)	0.025 (2)	-0.0045 (17)	0.0096 (17)	-0.0031 (16)
C7	0.047 (3)	0.037 (2)	0.028 (2)	0.015 (2)	0.0018 (19)	-0.0023 (18)

## data reports

C8	0.040 (3)	0.036 (3)	0.032 (3)	0.014 (2)	-0.002 (2)	0.0008 (19)
C9	0.031 (2)	0.029 (2)	0.025 (2)	-0.0005 (17)	0.0059 (18)	-0.0017 (17)

Geometric parameters (Å, °)

Geometric parameters (11, )			
01-C2	1.352 (6)	С2—С9	1.400 (6)
O1—C1	1.415 (6)	C3—C4	1.382 (6)
O2—C6	1.239 (5)	С3—НЗА	0.95
O3—C9	1.364 (5)	C4—C7	1.376 (6)
O3—H3	0.96 (3)	C4—C5	1.516 (6)
N1—C6	1.325 (6)	С5—Н5А	0.99
N1—C5	1.453 (6)	С5—Н5В	0.99
N1—H1	0.88	С6—Н6	0.95
C1—H1A	0.98	C7—C8	1.400 (6)
C1—H1B	0.98	С7—Н7	0.95
C1—H1C	0.98	C8—C9	1.374 (6)
С2—С3	1.389 (6)	C8—H8	0.95
C2—O1—C1	117.7 (3)	C3—C4—C5	120.6 (4)
С9—О3—Н3	106 (4)	N1—C5—C4	113.5 (3)
C6—N1—C5	124.2 (3)	N1—C5—H5A	108.9
C6—N1—H1	117.9	C4—C5—H5A	108.9
C5—N1—H1	117.9	N1—C5—H5B	108.9
O1—C1—H1A	109.5	C4—C5—H5B	108.9
O1—C1—H1B	109.5	H5A—C5—H5B	107.7
H1A—C1—H1B	109.5	O2—C6—N1	125.7 (4)
01—C1—H1C	109.5	O2—C6—H6	117.1
H1A—C1—H1C	109.5	N1—C6—H6	117.1
H1B—C1—H1C	109.5	C4—C7—C8	120.4 (4)
O1—C2—C3	125.9 (4)	С4—С7—Н7	119.8
O1—C2—C9	114.6 (4)	С8—С7—Н7	119.8
C3—C2—C9	119.5 (4)	C9—C8—C7	120.4 (4)
C4—C3—C2	121.1 (4)	С9—С8—Н8	119.8
С4—С3—Н3А	119.5	С7—С8—Н8	119.8
С2—С3—НЗА	119.5	O3—C9—C8	119.3 (4)
C7—C4—C3	119.2 (4)	O3—C9—C2	121.3 (4)
C7—C4—C5	120.3 (4)	C8—C9—C2	119.4 (4)
C1—O1—C2—C3	-6.5 (6)	C3—C4—C7—C8	0.2 (7)
C1—O1—C2—C9	174.5 (4)	C5—C4—C7—C8	179.6 (4)
O1—C2—C3—C4	179.7 (4)	C4—C7—C8—C9	-1.6 (7)
C9—C2—C3—C4	-1.3 (6)	С7—С8—С9—О3	-177.1 (4)
C2—C3—C4—C7	1.2 (6)	C7—C8—C9—C2	1.5 (7)
C2—C3—C4—C5	-178.1 (3)	O1—C2—C9—O3	-2.3 (5)
C6—N1—C5—C4	-97.8 (5)	C3—C2—C9—O3	178.5 (4)
C7—C4—C5—N1	65.2 (5)	O1—C2—C9—C8	179.0 (4)
C3—C4—C5—N1	-115.5 (4)	C3—C2—C9—C8	-0.1 (6)
C5—N1—C6—O2	1.3 (6)		

### Hydrogen-bond geometry (Å, °)

<i>D</i> —H··· <i>A</i>	D—H	H···A	$D \cdots A$	D—H···A
N1—H1····O2 <sup>i</sup>	0.88	2.05	2.910 (4)	167
O3—H3…O1	0.96 (3)	2.15 (7)	2.672 (4)	113 (5)
С6—Н6…ОЗ <sup>іі</sup>	0.95	2.54	3.391 (5)	149
O3—H3…O2 <sup>iii</sup>	0.96 (3)	1.90 (4)	2.771 (4)	149 (6)
C5—H5 <i>B</i> ····O3 <sup>iv</sup>	0.99	2.60	3.374 (5)	135
С5—Н5А…ОЗ <sup>v</sup>	0.99	2.70	3.676 (5)	167

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