

Crystal structure of 1,3-bis[(*E*)-benzylideneamino]-propan-2-ol

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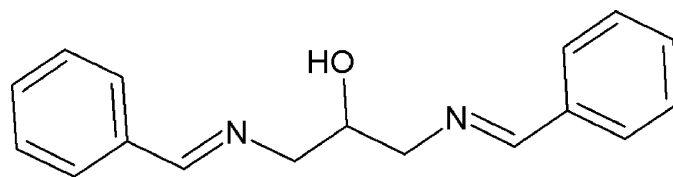
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Keywords: crystal structure; hydrogen bonding; C—H... π interactions; Schiff bases.**CCDC reference:** 1540296**Supporting information:** this article has supporting information at journals.iucr.org/e

In the title compound, C₁₇H₁₈N₂O, the central carbon atom with the OH substituent and one of the (*E*)-benzylideneamino substituents are disordered over two sets of sites with occupancies of 0.851 (4) and 0.149 (4). The relative positions of the two disorder components is equivalent to a rotation of approximately 60° about the C—N single bond. In the crystal, the molecules are held together by O—H...N hydrogen bonds, forming simple C(5) chains along the *b*-axis direction. In addition, pairs of the chains are further aggregated by weak C—H... π interactions.

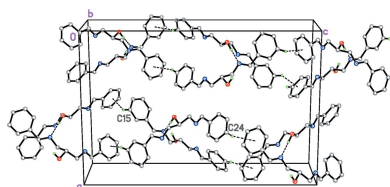
1. Chemical context

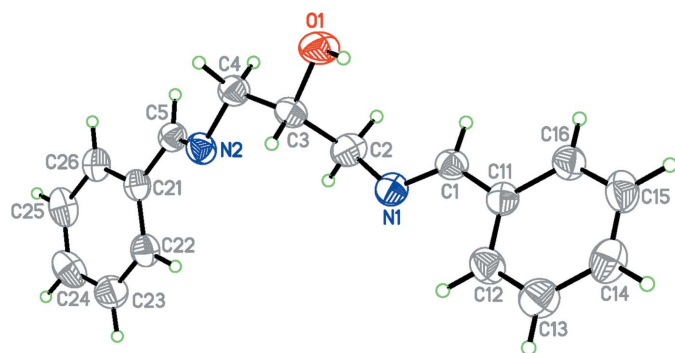
During the last decades, interest in Schiff bases and their complexes has been constant due to their extensive use for industrial purposes and also for their broad range of biological activities (Al Zoubi *et al.* 2016; Sahu *et al.* 2012; Da Silva *et al.*, 2011; Przybylski *et al.* 2009). The common structural feature of these compounds is the presence of a azomethine group ($-R-C=N-$), which can act as a hydrogen-bond acceptor or a ligand. To gain more insight into the structural and spectroscopic properties of this potentially polydentate ligand, we report herein the molecular structure of the title compound.



2. Structural commentary

The molecular structure of the title compound is shown in Fig. 1. The compound exists in an *E,E* conformation with respect to the imine functions. One benzylideneamino segment of the molecule, C3/C4/N2/C5/C21–C26 is disordered over two sets of sites with a refined occupancy ratio of 0.851 (4):0.149 (4). The difference between the two conformers is reflected in the relative arrangement of the central spacer units. In the major disorder component, the torsion angle C3—C4—N2—C5 is $-158.7(2)^\circ$ whereas the corresponding angle C3'—C4'—N2'—C5' in the minor component is $-93.3(14)^\circ$. This translates to a rotation of approximately 60° about the C4—N2 bond. In the second, fully ordered,



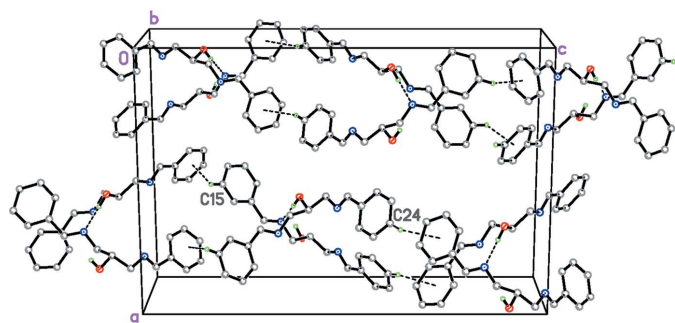

Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level. Only the major occupancy disorder component is shown.

(*E*)-benzylideneamino substituent, the equivalent torsion angles $C1-N1-C2-C3$ and $C1-N1-C2-C3'$ are $-102.03(18)^\circ$ and $-79.8(8)^\circ$, respectively.

Unlike some related structures, which have a well-defined *synclinal* (*-sc*) alignment of the hydroxyl and imine nitrogen atoms around the $N(\text{imine})-C-C-O(\text{hydroxyl})$ bond [$-65.3(3)^\circ$ (Rivera, Miranda-Carvajal, Ríos-Motta & Bolte, 2016) and $-67.6(4)^\circ$ (Moodley & Van Zyl, 2012)], the orientation between these groups in the title compound differs significantly, with the $N1-C2-C3-O1$ and $N1-C2-C3'-O1$ torsion angles being $81.51(19)^\circ$ and $21.2(14)^\circ$, respectively.

The $N1=C1$ and $N2=C5$ distances in the molecule are $1.270(2)$ and $1.259(3)$ Å, respectively, consistent with $C=N$ double bonding. The $C1-N1-C2$ bond angle of $118.61(15)^\circ$ confirms the sp^2 character of $N1$. The bond angles $C5-N2-C4$ and $C5'-N2'-C4'$ [$116.9(2)$ and $114.6(12)^\circ$, respectively] indicate a slight loss of the sp^2 character. The $N1=C1$ azomethine group is essentially co-planar with the attached benzene ring with an $N1-C1-C11-C12$ torsion angle being $2.0(5)^\circ$. In contrast, in the disordered (*E*)-benzylideneamino substituent, the corresponding torsion angles $N2-C5-C21-C22$ and $N2'-C5'-C21'-C22'$ are $-17.6(6)$ and $21(4)^\circ$ for the major and minor disorder components, respectively. All these data suggest that the difference between these (*E*)-


Figure 2

The crystal packing of the title compound showing the extended hydrogen-bonded network.

Table 1

Hydrogen-bond geometry (Å, °).

$Cg1, Cg2$, and $Cg3$, are the centroids of the $C11-C16$, $C21-C26$ and $C21'-C26'$ rings, respectively

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O1-H1\cdots N1^i$	0.93 (3)	1.92 (3)	2.8430 (19)	174 (2)
$C24-H24\cdots Cg1^{ii}$	0.95	2.88	3.802 (5)	164
$C15-H15\cdots Cg2^{iii}$	0.95	2.96	3.796 (3)	148
$C15-H15\cdots Cg3^{iii}$	0.95	2.79	3.640 (12)	150

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

benzylideneamino substituents may result from some loss of conjugation between the phenyl ring and its azomethine substituent in the disordered branch of the molecule.

3. Supramolecular features

As found in related structures (Rivera, Miranda-Carvajal, Ríos-Motta & Bolte, 2016; Moodley & Van Zyl, 2012) in the crystal, molecules are linked by an $O1-H1\cdots N1$ hydrogen bond, Table 1, forming columnar structures built from $C(5)$ chains along the *b*-axis direction. In addition, pairs of the chains are linked by weak $C24-H24\cdots Cg1$ interactions (Table 1 and Fig. 2), involving the $C11-C16$ phenyl ring, together with $C15-H15\cdots Cg2$ and $C15-H15\cdots Cg3$ contacts involving the phenyl rings of the two disorder components; the centroids are defined in Table 1. It is noteworthy that the shortest (and presumably the strongest) of these non-classical contacts is $C15-H15\cdots Cg3$ involving the phenyl ring in the minor disorder component (Table 1).

4. Database survey

A search in the Cambridge Crystallographic Database (CSD Version 5.38, last update 2016; Groom *et al.*, 2016) for the fragment 1,3-bis[(benzylidene)amino]propan-2-ol yielded the following structures: *N,N'*-[(2-hydroxy-1,3-propanediyl)bis(nitrilomethylidene-2,1-phenylene)] bis(4-methylbenzenesulfonamide) (Popov *et al.*, 2009), 2,2'-[(2-hydroxypropane-1,3-diyl)bis(nitrilomethylidene)]diphenol (Azam, Hussain *et al.*, 2012), 1,3-bis(2-hydroxy-5-bromosalicylideneamine)propan-2-ol (Elmali, 2000), 1,3-bis[(*E*)-(2-chlorobenzylidene)amino]propan-2-ol (Azam, Warad *et al.*, 2012) and 1,3-bis[(*E*)-(4-methoxybenzylidene)amino]propan-2-ol (Rivera, Miranda-Carvajal, Ríos-Motta & Bolte, 2016). In each of these structures, the $N=C$ double bonds adopt *E* conformations.

5. Synthesis and crystallization

The title compound was prepared as described by Rivera, Miranda-Carvajal & Ríos-Motta (2016). The crude product was recrystallized from diethyl ether solution with slow evaporation of the solvent, giving colorless crystals suitable for X-ray diffraction, m.p. 396.8–398 K, yield, 40%.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The hydroxyl H atom was refined freely. All remaining H atoms were positioned geometrically and allowed to ride on their parent atoms, with $d(\text{C}-\text{H}) = 0.95 \text{ \AA}$ for aromatic and azomethine atoms, $d(\text{C}-\text{H}) = 0.99 \text{ \AA}$ for methylene and $d(\text{C}-\text{H}) = 1.00 \text{ \AA}$ for C3–H3. The $U_{\text{iso}}(\text{H})$ values were constrained to $1.2U_{\text{eq}}(\text{C})$. The C3/C4/N2/C5/C21–C26 segment of the molecule is disordered over two sets of sites with a refined occupancy ratio of 0.851 (4):0.149 (4).

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Table 2

Experimental details.

Crystal data	
Chemical formula	C ₁₇ H ₁₈ N ₂ O
M_r	266.33
Crystal system, space group	Orthorhombic, <i>Pbca</i>
Temperature (K)	173
a, b, c (Å)	16.4313 (7), 7.1909 (3), 24.7345 (11)
V (Å ³)	2922.5 (2)
Z	8
Radiation type	Mo $K\alpha$
μ (mm ⁻¹)	0.08
Crystal size (mm)	0.24 × 0.22 × 0.18
Data collection	
Diffractometer	STOE IPDS II two-circle
Absorption correction	Multi-scan (<i>X-AREA</i> ; Stoe & Cie, 2001)
$T_{\text{min}}, T_{\text{max}}$	0.742, 1.000
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	25768, 2574, 2200
R_{int}	0.054
$(\sin \theta/\lambda)_{\text{max}}$ (Å ⁻¹)	0.595
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.046, 0.111, 1.10
No. of reflections	2574
No. of parameters	276
No. of restraints	84
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)	0.24, -0.21

Computer programs: *X-AREA* (Stoe & Cie, 2001), *SHELXS2014/7* and *XP* in *SHELXTL-Plus* (Sheldrick, 2008) and *SHELXL2014/7* (Sheldrick, 2015).

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supporting information

Acta Cryst. (2017). E73, 627-629 [https://doi.org/10.1107/S2056989017004741]

Crystal structure of 1,3-bis[(*E*)-benzylideneamino]propan-2-ol

Augusto Rivera, Ingrid Miranda-Carvajal, Jaime Ríos-Motta and Michael Bolte

Computing details

Data collection: *X-AREA* (Stoe & Cie, 2001); cell refinement: *X-AREA* (Stoe & Cie, 2001); data reduction: *X-AREA* (Stoe & Cie, 2001); program(s) used to solve structure: *SHELXS2014/7* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2014/7* (Sheldrick, 2015); molecular graphics: *XP* in *SHELXTL-Plus* (Sheldrick, 2008).

1,3-Bis[(*E*)-benzylideneamino]propan-2-ol

Crystal data

$C_{17}H_{18}N_2O$

$M_r = 266.33$

Orthorhombic, *Pbca*

$a = 16.4313$ (7) Å

$b = 7.1909$ (3) Å

$c = 24.7345$ (11) Å

$V = 2922.5$ (2) Å³

$Z = 8$

$F(000) = 1136$

$D_x = 1.211$ Mg m⁻³

Mo *K*α radiation, $\lambda = 0.71073$ Å

Cell parameters from 24714 reflections

$\theta = 2.1$ – 25.5°

$\mu = 0.08$ mm⁻¹

$T = 173$ K

Block, colourless

$0.24 \times 0.22 \times 0.18$ mm

Data collection

STOE IPDS II two-circle

diffractometer

Radiation source: Genix 3D IμS microfocus X-

ray source

ω scans

Absorption correction: multi-scan

(*X-Area*; Stoe & Cie, 2001)

$T_{\min} = 0.742$, $T_{\max} = 1.000$

25768 measured reflections

2574 independent reflections

2200 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.054$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.1^\circ$

$h = -18 \rightarrow 19$

$k = -8 \rightarrow 8$

$l = -29 \rightarrow 29$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.046$

$wR(F^2) = 0.111$

$S = 1.10$

2574 reflections

276 parameters

84 restraints

Hydrogen site location: mixed

H atoms treated by a mixture of independent

and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0542P)^2 + 0.639P]$

where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.24$ e Å⁻³

$\Delta\rho_{\min} = -0.21$ e Å⁻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>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1	0.63353 (8)	0.70433 (16)	0.38320 (5)	0.0509 (4)	
H1	0.6763 (14)	0.763 (3)	0.3660 (9)	0.075 (7)*	
N1	0.74117 (8)	0.41062 (18)	0.33137 (5)	0.0392 (3)	
C1	0.73647 (10)	0.4889 (2)	0.28545 (6)	0.0364 (4)	
H1A	0.6840	0.5185	0.2718	0.044*	
C2	0.66704 (11)	0.3836 (2)	0.36223 (7)	0.0470 (4)	
H2A	0.6668	0.2584	0.3789	0.056*	0.851 (4)
H2B	0.6190	0.3942	0.3383	0.056*	0.851 (4)
H2C	0.6278	0.3353	0.3353	0.056*	0.149 (4)
H2D	0.6797	0.2768	0.3861	0.056*	0.149 (4)
C3	0.66388 (15)	0.5357 (3)	0.40687 (9)	0.0350 (5)	0.851 (4)
H3	0.7194	0.5566	0.4222	0.042*	0.851 (4)
C4	0.60471 (12)	0.4859 (3)	0.45148 (8)	0.0394 (6)	0.851 (4)
H4A	0.6007	0.5912	0.4771	0.047*	0.851 (4)
H4B	0.5501	0.4654	0.4356	0.047*	0.851 (4)
N2	0.62980 (16)	0.3185 (3)	0.48085 (10)	0.0391 (5)	0.851 (4)
C5	0.57527 (15)	0.2327 (3)	0.50643 (8)	0.0361 (5)	0.851 (4)
H5	0.5210	0.2773	0.5038	0.043*	0.851 (4)
C21	0.5901 (2)	0.0675 (5)	0.54025 (17)	0.0352 (7)	0.851 (4)
C22	0.6613 (3)	-0.0353 (9)	0.5350 (3)	0.0425 (15)	0.851 (4)
H22	0.7005	-0.0016	0.5085	0.051*	0.851 (4)
C23	0.6751 (4)	-0.1864 (10)	0.5681 (3)	0.0544 (14)	0.851 (4)
H23	0.7232	-0.2582	0.5639	0.065*	0.851 (4)
C24	0.6186 (3)	-0.2338 (6)	0.6078 (2)	0.0561 (10)	0.851 (4)
H24	0.6295	-0.3338	0.6318	0.067*	0.851 (4)
C25	0.5473 (3)	-0.1359 (6)	0.61203 (15)	0.0503 (10)	0.851 (4)
H25	0.5080	-0.1711	0.6383	0.060*	0.851 (4)
C26	0.53203 (18)	0.0143 (5)	0.57820 (13)	0.0409 (7)	0.851 (4)
H26	0.4822	0.0805	0.5809	0.049*	0.851 (4)
C11	0.80792 (10)	0.5363 (2)	0.25222 (6)	0.0356 (4)	
C12	0.88687 (10)	0.4938 (2)	0.26791 (7)	0.0440 (4)	
H12	0.8961	0.4264	0.3004	0.053*	
C13	0.95204 (11)	0.5492 (3)	0.23646 (7)	0.0499 (5)	
H13	1.0058	0.5180	0.2472	0.060*	
C14	0.93971 (12)	0.6499 (2)	0.18938 (7)	0.0477 (4)	
H14	0.9848	0.6897	0.1683	0.057*	
C15	0.86123 (12)	0.6919 (2)	0.17337 (7)	0.0474 (4)	
H15	0.8522	0.7605	0.1411	0.057*	
C16	0.79573 (11)	0.6343 (2)	0.20430 (6)	0.0412 (4)	

H16	0.7419	0.6619	0.1927	0.049*	
C3'	0.6215 (9)	0.4977 (16)	0.3939 (5)	0.042 (3)	0.149 (4)
H3'	0.5637	0.4580	0.3990	0.050*	0.149 (4)
C4'	0.6731 (7)	0.4957 (14)	0.4447 (5)	0.040 (3)	0.149 (4)
H4'1	0.7295	0.5323	0.4356	0.049*	0.149 (4)
H4'2	0.6512	0.5870	0.4709	0.049*	0.149 (4)
N2'	0.6732 (9)	0.3067 (15)	0.4696 (4)	0.044 (3)	0.149 (4)
C5'	0.6203 (12)	0.283 (2)	0.5052 (6)	0.041 (3)	0.149 (4)
H5'	0.5840	0.3821	0.5129	0.049*	0.149 (4)
C21'	0.6110 (12)	0.104 (2)	0.5364 (9)	0.032 (4)	0.149 (4)
C22'	0.6731 (19)	-0.025 (5)	0.5415 (17)	0.035 (5)	0.149 (4)
H22'	0.7195	-0.0144	0.5188	0.042*	0.149 (4)
C23'	0.6693 (15)	-0.169 (5)	0.5788 (14)	0.039 (5)	0.149 (4)
H23'	0.7164	-0.2393	0.5873	0.047*	0.149 (4)
C24'	0.5946 (13)	-0.208 (4)	0.6038 (14)	0.048 (6)	0.149 (4)
H24'	0.5856	-0.3222	0.6219	0.057*	0.149 (4)
C25'	0.5353 (11)	-0.075 (3)	0.6010 (8)	0.034 (4)	0.149 (4)
H25'	0.4873	-0.0892	0.6221	0.040*	0.149 (4)
C26'	0.5437 (12)	0.079 (3)	0.5682 (9)	0.040 (4)	0.149 (4)
H26'	0.5019	0.1705	0.5676	0.048*	0.149 (4)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0560 (8)	0.0352 (6)	0.0615 (8)	0.0078 (6)	0.0122 (6)	0.0145 (6)
N1	0.0464 (8)	0.0304 (6)	0.0409 (7)	-0.0005 (6)	0.0045 (6)	0.0015 (6)
C1	0.0401 (9)	0.0291 (7)	0.0402 (9)	0.0019 (6)	-0.0012 (7)	-0.0004 (6)
C2	0.0511 (10)	0.0366 (9)	0.0534 (10)	0.0021 (8)	0.0146 (8)	0.0083 (8)
C3	0.0352 (12)	0.0296 (10)	0.0401 (12)	-0.0021 (9)	-0.0030 (10)	0.0047 (9)
C4	0.0412 (13)	0.0369 (10)	0.0400 (10)	0.0021 (8)	0.0023 (8)	0.0030 (8)
N2	0.0373 (13)	0.0433 (11)	0.0367 (13)	-0.0005 (10)	0.0011 (11)	0.0064 (9)
C5	0.0352 (11)	0.0372 (11)	0.0360 (10)	-0.0012 (10)	-0.0007 (9)	-0.0028 (8)
C21	0.0364 (18)	0.0357 (14)	0.0334 (15)	-0.0080 (12)	-0.0046 (13)	-0.0033 (11)
C22	0.044 (2)	0.0458 (18)	0.038 (2)	-0.0041 (18)	-0.003 (2)	-0.0014 (17)
C23	0.066 (2)	0.044 (2)	0.053 (3)	0.0077 (14)	-0.0057 (16)	-0.003 (2)
C24	0.068 (3)	0.042 (2)	0.0583 (17)	-0.0111 (18)	-0.018 (2)	0.0112 (16)
C25	0.061 (2)	0.048 (2)	0.0417 (15)	-0.021 (2)	-0.0047 (15)	0.0085 (15)
C26	0.0428 (13)	0.042 (2)	0.0376 (19)	-0.0106 (14)	-0.0008 (11)	-0.0017 (15)
C11	0.0431 (9)	0.0272 (7)	0.0366 (8)	-0.0011 (6)	0.0013 (7)	-0.0019 (6)
C12	0.0458 (10)	0.0428 (9)	0.0435 (9)	-0.0006 (7)	-0.0010 (8)	0.0081 (7)
C13	0.0408 (10)	0.0540 (11)	0.0549 (11)	-0.0038 (8)	0.0022 (8)	0.0029 (9)
C14	0.0540 (11)	0.0424 (9)	0.0468 (10)	-0.0112 (8)	0.0105 (8)	-0.0020 (8)
C15	0.0643 (12)	0.0406 (9)	0.0373 (9)	-0.0027 (8)	0.0040 (8)	0.0043 (7)
C16	0.0486 (10)	0.0374 (8)	0.0377 (8)	0.0034 (7)	-0.0001 (7)	0.0005 (7)
C3'	0.046 (7)	0.030 (6)	0.050 (6)	-0.017 (5)	0.000 (5)	0.001 (4)
C4'	0.045 (7)	0.041 (5)	0.036 (6)	-0.007 (4)	0.007 (5)	-0.003 (4)
N2'	0.048 (7)	0.047 (5)	0.038 (5)	-0.004 (5)	0.001 (5)	0.002 (4)
C5'	0.046 (7)	0.040 (6)	0.035 (6)	-0.006 (5)	-0.003 (6)	0.000 (5)

C21'	0.039 (8)	0.033 (6)	0.024 (6)	-0.007 (5)	-0.004 (5)	-0.005 (5)
C22'	0.029 (7)	0.033 (6)	0.043 (9)	-0.007 (5)	-0.022 (6)	-0.012 (5)
C23'	0.036 (7)	0.035 (8)	0.046 (11)	0.006 (5)	-0.014 (5)	-0.006 (8)
C24'	0.042 (8)	0.029 (7)	0.073 (13)	0.010 (7)	-0.002 (8)	0.011 (6)
C25'	0.030 (6)	0.029 (8)	0.042 (10)	0.006 (5)	-0.006 (5)	0.005 (5)
C26'	0.047 (7)	0.030 (8)	0.042 (7)	0.003 (5)	-0.001 (5)	0.005 (6)

Geometric parameters (Å, °)

O1—C3	1.436 (2)	C26—H26	0.9500
O1—C3'	1.522 (11)	C11—C12	1.388 (2)
O1—H1	0.93 (3)	C11—C16	1.394 (2)
N1—C1	1.270 (2)	C12—C13	1.382 (2)
N1—C2	1.451 (2)	C12—H12	0.9500
C1—C11	1.473 (2)	C13—C14	1.386 (3)
C1—H1A	0.9500	C13—H13	0.9500
C2—C3'	1.359 (13)	C14—C15	1.382 (3)
C2—C3	1.555 (3)	C14—H14	0.9500
C2—H2A	0.9900	C15—C16	1.384 (2)
C2—H2B	0.9900	C15—H15	0.9500
C2—H2C	0.9900	C16—H16	0.9500
C2—H2D	0.9900	C3'—C4'	1.515 (14)
C3—C4	1.513 (3)	C3'—H3'	1.0000
C3—H3	1.0000	C4'—N2'	1.492 (13)
C4—N2	1.465 (3)	C4'—H4'1	0.9900
C4—H4A	0.9900	C4'—H4'2	0.9900
C4—H4B	0.9900	N2'—C5'	1.249 (15)
N2—C5	1.259 (3)	C5'—C21'	1.508 (15)
C5—C21	1.473 (3)	C5'—H5'	0.9500
C5—H5	0.9500	C21'—C26'	1.367 (15)
C21—C22	1.390 (4)	C21'—C22'	1.382 (16)
C21—C26	1.392 (4)	C22'—C23'	1.389 (17)
C22—C23	1.380 (4)	C22'—H22'	0.9500
C22—H22	0.9500	C23'—C24'	1.402 (17)
C23—C24	1.392 (5)	C23'—H23'	0.9500
C23—H23	0.9500	C24'—C25'	1.364 (16)
C24—C25	1.372 (5)	C24'—H24'	0.9500
C24—H24	0.9500	C25'—C26'	1.384 (15)
C25—C26	1.389 (4)	C25'—H25'	0.9500
C25—H25	0.9500	C26'—H26'	0.9500
C3—O1—H1	108.2 (15)	C12—C11—C16	118.91 (15)
C3'—O1—H1	128.8 (16)	C12—C11—C1	122.55 (14)
C1—N1—C2	118.61 (15)	C16—C11—C1	118.49 (15)
N1—C1—C11	123.57 (15)	C13—C12—C11	120.21 (16)
N1—C1—H1A	118.2	C13—C12—H12	119.9
C11—C1—H1A	118.2	C11—C12—H12	119.9
C3'—C2—N1	133.2 (6)	C12—C13—C14	120.68 (17)

N1—C2—C3	107.89 (15)	C12—C13—H13	119.7
N1—C2—H2A	110.1	C14—C13—H13	119.7
C3—C2—H2A	110.1	C15—C14—C13	119.42 (16)
N1—C2—H2B	110.1	C15—C14—H14	120.3
C3—C2—H2B	110.1	C13—C14—H14	120.3
H2A—C2—H2B	108.4	C14—C15—C16	120.11 (16)
C3'—C2—H2C	103.9	C14—C15—H15	119.9
N1—C2—H2C	103.9	C16—C15—H15	119.9
C3'—C2—H2D	103.9	C15—C16—C11	120.65 (16)
N1—C2—H2D	103.9	C15—C16—H16	119.7
H2C—C2—H2D	105.4	C11—C16—H16	119.7
O1—C3—C4	105.89 (16)	C2—C3'—C4'	99.4 (10)
O1—C3—C2	108.42 (17)	C2—C3'—O1	114.7 (8)
C4—C3—C2	111.88 (17)	C4'—C3'—O1	94.6 (9)
O1—C3—H3	110.2	C2—C3'—H3'	115.1
C4—C3—H3	110.2	C4'—C3'—H3'	115.1
C2—C3—H3	110.2	O1—C3'—H3'	115.1
N2—C4—C3	112.03 (18)	N2'—C4'—C3'	110.6 (9)
N2—C4—H4A	109.2	N2'—C4'—H4'1	109.5
C3—C4—H4A	109.2	C3'—C4'—H4'1	109.5
N2—C4—H4B	109.2	N2'—C4'—H4'2	109.5
C3—C4—H4B	109.2	C3'—C4'—H4'2	109.5
H4A—C4—H4B	107.9	H4'1—C4'—H4'2	108.1
C5—N2—C4	116.9 (2)	C5'—N2'—C4'	114.6 (12)
N2—C5—C21	124.3 (2)	N2'—C5'—C21'	123.3 (14)
N2—C5—H5	117.9	N2'—C5'—H5'	118.3
C21—C5—H5	117.9	C21'—C5'—H5'	118.3
C22—C21—C26	119.6 (3)	C26'—C21'—C22'	117.3 (15)
C22—C21—C5	121.0 (3)	C26'—C21'—C5'	119.1 (15)
C26—C21—C5	119.4 (3)	C22'—C21'—C5'	122.8 (16)
C23—C22—C21	120.1 (4)	C21'—C22'—C23'	122 (2)
C23—C22—H22	120.0	C21'—C22'—H22'	119.2
C21—C22—H22	120.0	C23'—C22'—H22'	119.2
C22—C23—C24	120.1 (4)	C22'—C23'—C24'	118.8 (19)
C22—C23—H23	119.9	C22'—C23'—H23'	120.6
C24—C23—H23	119.9	C24'—C23'—H23'	120.6
C25—C24—C23	119.9 (3)	C25'—C24'—C23'	117.6 (17)
C25—C24—H24	120.1	C25'—C24'—H24'	121.2
C23—C24—H24	120.1	C23'—C24'—H24'	121.2
C24—C25—C26	120.5 (3)	C24'—C25'—C26'	121.2 (16)
C24—C25—H25	119.8	C24'—C25'—H25'	119.4
C26—C25—H25	119.8	C26'—C25'—H25'	119.4
C25—C26—C21	119.7 (3)	C21'—C26'—C25'	121.4 (15)
C25—C26—H26	120.1	C21'—C26'—H26'	119.3
C21—C26—H26	120.1	C25'—C26'—H26'	119.3
C2—N1—C1—C11	175.36 (14)	C1—C11—C12—C13	-176.91 (15)
C1—N1—C2—C3'	-79.8 (8)	C11—C12—C13—C14	0.9 (3)

C1—N1—C2—C3	-102.03 (18)	C12—C13—C14—C15	-1.2 (3)
N1—C2—C3—O1	81.51 (19)	C13—C14—C15—C16	0.2 (3)
N1—C2—C3'—O1	21.2 (14)	C14—C15—C16—C11	1.1 (3)
N1—C2—C3—C4	-162.08 (16)	C12—C11—C16—C15	-1.3 (2)
O1—C3—C4—N2	-177.91 (17)	C1—C11—C16—C15	176.04 (14)
C2—C3—C4—N2	64.2 (2)	N1—C2—C3'—C4'	-78.4 (8)
C3—C4—N2—C5	-158.7 (2)	N1—C2—C3'—O1	21.2 (14)
C4—N2—C5—C21	-176.9 (3)	C2—C3'—C4'—N2'	-66.3 (11)
N2—C5—C21—C22	-17.6 (6)	O1—C3'—C4'—N2'	177.7 (9)
N2—C5—C21—C26	162.1 (3)	C3'—C4'—N2'—C5'	-93.3 (14)
C26—C21—C22—C23	-1.6 (10)	C4'—N2'—C5'—C21'	-178.5 (15)
C5—C21—C22—C23	178.1 (6)	N2'—C5'—C21'—C26'	-169.5 (18)
C21—C22—C23—C24	-1.5 (12)	N2'—C5'—C21'—C22'	21 (4)
C22—C23—C24—C25	3.3 (11)	C26'—C21'—C22'—C23'	-3 (6)
C23—C24—C25—C26	-2.1 (7)	C5'—C21'—C22'—C23'	167 (4)
C24—C25—C26—C21	-0.9 (5)	C21'—C22'—C23'—C24'	14 (7)
C22—C21—C26—C25	2.8 (6)	C22'—C23'—C24'—C25'	-17 (6)
C5—C21—C26—C25	-176.9 (3)	C23'—C24'—C25'—C26'	10 (5)
N1—C1—C11—C12	2.0 (2)	C22'—C21'—C26'—C25'	-5 (4)
N1—C1—C11—C16	-175.34 (15)	C5'—C21'—C26'—C25'	-174.7 (19)
C16—C11—C12—C13	0.4 (2)	C24'—C25'—C26'—C21'	1 (4)

Hydrogen-bond geometry (Å, °)

*Cg*1, *Cg*2, and *Cg*3, are the centroids of the C11—C16, C21—C26 and C21'—C26' rings, respectively

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O1—H1...N1 ⁱ	0.93 (3)	1.92 (3)	2.8430 (19)	174 (2)
C24—H24... <i>Cg</i> 1 ⁱⁱ	0.95	2.88	3.802 (5)	164
C15—H15... <i>Cg</i> 2 ⁱⁱⁱ	0.95	2.96	3.796 (3)	148
C15—H15... <i>Cg</i> 3 ⁱⁱⁱ	0.95	2.79	3.640 (12)	150

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