

Crystal structure of (*E*)-5-diethylamino-2-([4-(dimethylamino)phenyl]imino)methylphenol

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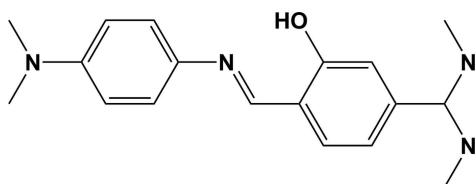
The title Schiff base compound, $C_{19}H_{25}N_3O$, is approximately planar, with a dihedral angle of $9.03(13)^\circ$ between the planes of the aromatic rings, and has an *E* conformation about the $N\equiv C$ bond. The molecular structure is stabilized by an intramolecular $O-H\cdots N$ hydrogen bond, with an *S*(6) ring motif. In the crystal, molecules are linked by $C-H\cdots \pi$ interactions, forming sheets parallel to the *bc* plane.

Keywords: crystal structure; Schiff base; intramolecular $O-H\cdots N$ hydrogen bond; $C-H\cdots \pi$ interactions.

CCDC reference: 1407678

1. Related literature

For biological activities of Schiff base derivatives, see: Savaliya *et al.* (2010); Xu *et al.* (2012). For the structures of similar compounds, see: Manvizhi *et al.* (2011); Thirugnanasundar *et al.* (2011).



2. Experimental

2.1. Crystal data

$C_{19}H_{25}N_3O$
 $M_r = 311.42$
Monoclinic, $P2_1$
 $a = 8.8201(7)\text{ \AA}$

$b = 7.8850(7)\text{ \AA}$
 $c = 13.0639(10)\text{ \AA}$
 $\beta = 108.407(3)^\circ$
 $V = 862.06(12)\text{ \AA}^3$

$Z = 2$
Mo $K\alpha$ radiation
 $\mu = 0.08\text{ mm}^{-1}$

$T = 295\text{ K}$
 $0.26 \times 0.22 \times 0.20\text{ mm}$

2.2. Data collection

Bruker Kappa APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.981$, $T_{\max} = 0.985$

13009 measured reflections
3825 independent reflections
2438 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.159$
 $S = 1.03$
3825 reflections
214 parameters

6 restraints
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.29\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.16\text{ e \AA}^{-3}$

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

Cg1 and *Cg2* are the centroids of rings C3–C8 and C10–C15, respectively.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1—H1 \cdots N2	0.82	1.85	2.585 (3)	148
C11—H11 \cdots Cg1 ⁱ	0.93	2.71	3.517 (3)	145
C17—H17B \cdots Cg2 ⁱⁱ	0.96	2.90	3.743 (5)	147

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

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *SHELXL97* and *PLATON*.

Acknowledgements

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Supporting information for this paper is available from the IUCr electronic archives (Reference: SU5157).

References

- Bruker (2004). *APEX2* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Macrae, C. F., Bruno, I. J., Chisholm, J. A., Edgington, P. R., McCabe, P., Pidcock, E., Rodriguez-Monge, L., Taylor, R., van de Streek, J. & Wood, P. A. (2008). *J. Appl. Cryst.* **41**, 466–470.
- Manvizhi, K., Chakkavarthi, G., Anbalagan, G. & Rajagopal, G. (2011). *Acta Cryst. E* **67**, o2500.
- Savaliya, M. D., Dobaria, J. G. & Purohit, D. M. (2010). *An Indian J.* **6**, 267–271.
- Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Spek, A. L. (2009). *Acta Cryst. D* **65**, 148–155.
- Thirugnanasundar, A., Suresh, J., Ramu, A. & RajaGopal, G. (2011). *Acta Cryst. E* **67**, o2303.
- Xu, R.-B., Zhang, N., Zhou, H.-Y., Yang, S.-P., Li, Y.-Y., Shi, D.-H., Ma, W.-X. & Xu, X.-Y. (2012). *J. Chem. Crystallogr.* **42**, 928–932.

supporting information

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Crystal structure of (*E*)-5-diethylamino-2-({[4-(dimethylamino)phenyl]imino}-methyl)phenol

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S1. Structural commentary

Schiff base derivatives are known to exhibit antimicrobial (Savaliya *et al.*, 2010) and antibacterial (Xu *et al.*, 2012) activities. Herein we report on the synthesis and the crystal structure of a new Schiff base compound.

The molecular structure of the title compound is illustrated in Fig. 1. The geometric parameters are comparable to those reported for similar structures (Manvizhi *et al.*, 2011; Thirugnanasundar *et al.*, 2011). The dihedral angle between the benzene rings (C3—C8) and (C10—C15) is 9.03 (13)°. The molecular structure is stabilized by an intramolecular O—H···N hydrogen bond (Table 1 and Fig. 1).

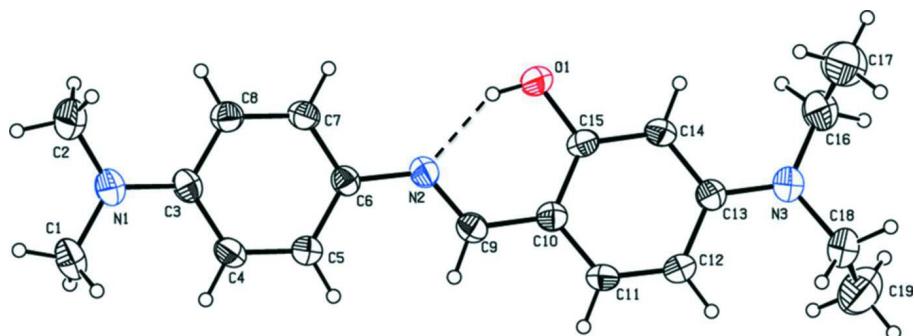
In the crystal, molecules are linked by C—H···π interactions forming sheets parallel to the bc plane (Table 1 and Fig. 2).

S2. Synthesis and crystallization

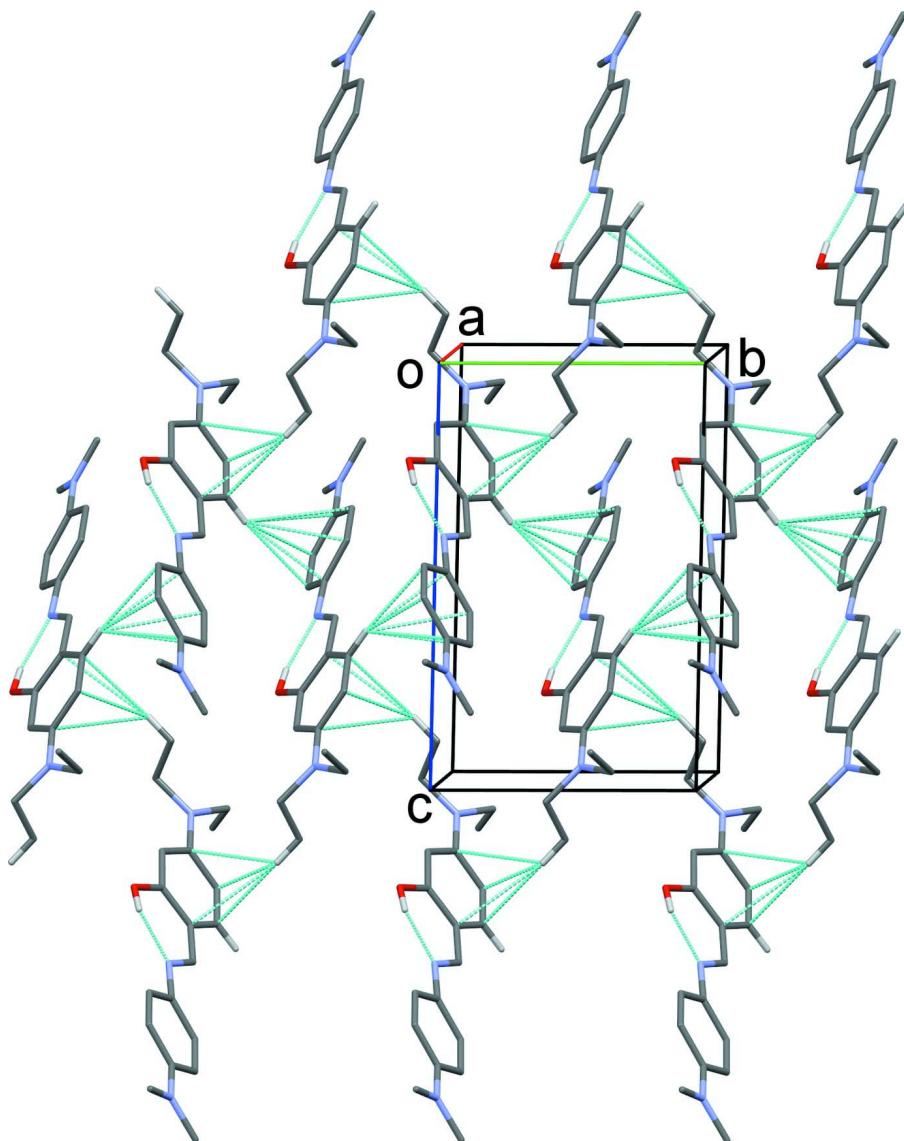
To an ethanol solution (10 ml) of 5-(diethylamino)-2-hydroxybenzaldehyde (96.5 mg, 0.5 mol) was added *N,N'*-dimethylbenzene-1,4-diamine (68 mg, 0.5 mol). The mixture was stirred and 2 to 3 drops of glacial acetic acid were added. Stirring was continued for 30 mins and then the reaction mixture was refluxed for 2 h. On completion of the reaction, monitored by TLC, the mixture was allowed to cool to room temperature and the solid yellow precipitate that formed was filtered, dried, and recrystallized from DMF, giving colourless block-like crystals.

S3. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. H atoms were positioned geometrically and refined using riding model: O—H = 0.82 Å, C—H = 0.93 - 0.97 Å with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O,C})$ for the hydroxyl and methyl H atoms and $1.2U_{\text{eq}}(\text{C})$ for other H atoms. The components of the anisotropic displacement parameters of the atoms in bonds N3—C16, N3—C18 and N1—C2 were restrained to be equal within an effective standard deviation of 0.001 using the DELU command, and the C16—C17 bond distance was restrained to 1.54 (1) Å.

**Figure 1**

The molecular structure of the title compound, with atom labelling. Displacement ellipsoids are drawn at the 30% probability level. The intramolecular O—H..N hydrogen bonds is shown as a dashd lines (see Table 1 for details).

**Figure 2**

A view along the *a* axis of the crystal apcking of the title compound. The O—H..N and C-H \cdots π interactions are illustrated by dashed lines (see Table 1 for details).

(E)-5-Diethylamino-2-((4-(dimethylamino)phenyl)imino)methylphenol

Crystal data

C₁₉H₂₅N₃O
 $M_r = 311.42$
 Monoclinic, P2₁
 Hall symbol: P 2yb
 $a = 8.8201 (7)$ Å
 $b = 7.8850 (7)$ Å
 $c = 13.0639 (10)$ Å
 $\beta = 108.407 (3)$ °
 $V = 862.06 (12)$ Å³
 $Z = 2$

$F(000) = 336$
 $D_x = 1.200 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 4428 reflections
 $\theta = 2.4\text{--}27.2$ °
 $\mu = 0.08 \text{ mm}^{-1}$
 $T = 295 \text{ K}$
 Block, colourless
 $0.26 \times 0.22 \times 0.20 \text{ mm}$

Data collection

Bruker Kappa APEXII CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω and φ scan
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.981$, $T_{\max} = 0.985$

13009 measured reflections
3825 independent reflections
2438 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$
 $\theta_{\max} = 27.3^\circ$, $\theta_{\min} = 2.4^\circ$
 $h = -11 \rightarrow 10$
 $k = -10 \rightarrow 10$
 $l = -16 \rightarrow 16$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.159$
 $S = 1.03$
3825 reflections
214 parameters
6 restraints
Primary atom site location: structure-invariant
direct methods
Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0751P)^2 + 0.1538P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.29 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.16 \text{ e } \text{\AA}^{-3}$
Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.014 (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.

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.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.4431 (4)	0.6814 (5)	0.2015 (2)	0.0780 (10)
H1A	0.3484	0.6253	0.1573	0.117*
H1B	0.5213	0.6840	0.1644	0.117*
H1C	0.4171	0.7953	0.2159	0.117*
C2	0.6641 (4)	0.5264 (6)	0.3287 (3)	0.0902 (12)
H2A	0.7316	0.5867	0.3901	0.135*
H2B	0.7039	0.5405	0.2688	0.135*
H2C	0.6635	0.4081	0.3459	0.135*
C3	0.4220 (3)	0.5850 (3)	0.3736 (2)	0.0502 (6)
C4	0.2713 (3)	0.6598 (4)	0.3525 (2)	0.0547 (7)
H4	0.2275	0.7180	0.2880	0.066*
C5	0.1861 (3)	0.6502 (4)	0.42371 (19)	0.0512 (6)
H5	0.0860	0.7008	0.4067	0.061*
C6	0.2487 (3)	0.5648 (3)	0.52175 (18)	0.0444 (6)
C7	0.3965 (3)	0.4922 (3)	0.5425 (2)	0.0508 (6)

H7	0.4405	0.4352	0.6075	0.061*
C8	0.4820 (3)	0.5003 (4)	0.4715 (2)	0.0537 (7)
H8	0.5816	0.4483	0.4889	0.064*
C9	0.0435 (3)	0.6249 (4)	0.59847 (19)	0.0481 (6)
H9	-0.0034	0.6975	0.5411	0.058*
C10	-0.0294 (3)	0.6066 (3)	0.68154 (19)	0.0462 (6)
C11	-0.1669 (3)	0.6947 (3)	0.6784 (2)	0.0537 (7)
H11	-0.2150	0.7626	0.6188	0.064*
C12	-0.2347 (3)	0.6864 (4)	0.7586 (2)	0.0596 (8)
H12	-0.3278	0.7467	0.7524	0.072*
C13	-0.1648 (3)	0.5871 (4)	0.8507 (2)	0.0528 (6)
C14	-0.0279 (3)	0.4940 (3)	0.8547 (2)	0.0523 (7)
H14	0.0198	0.4255	0.9141	0.063*
C15	0.0370 (3)	0.5025 (3)	0.7722 (2)	0.0472 (6)
C16	-0.1784 (4)	0.4511 (5)	1.0194 (3)	0.0834 (10)
H16A	-0.2702	0.4176	1.0403	0.100*
H16B	-0.1393	0.3520	0.9915	0.100*
C17	-0.0541 (5)	0.5166 (6)	1.1125 (3)	0.1089 (14)
H17A	0.0389	0.5432	1.0925	0.163*
H17B	-0.0274	0.4327	1.1686	0.163*
H17C	-0.0917	0.6173	1.1380	0.163*
C18	-0.3507 (4)	0.7080 (5)	0.9409 (3)	0.0835 (10)
H18A	-0.3375	0.7326	1.0160	0.100*
H18B	-0.3372	0.8131	0.9062	0.100*
C19	-0.5106 (4)	0.6420 (8)	0.8890 (4)	0.1178 (15)
H19A	-0.5223	0.6129	0.8156	0.177*
H19B	-0.5881	0.7267	0.8905	0.177*
H19C	-0.5270	0.5428	0.9268	0.177*
N1	0.5063 (3)	0.5915 (4)	0.30117 (18)	0.0693 (7)
N2	0.1707 (2)	0.5451 (3)	0.60028 (16)	0.0495 (5)
N3	-0.2261 (3)	0.5836 (4)	0.9344 (2)	0.0826 (9)
O1	0.1678 (2)	0.4075 (3)	0.77919 (17)	0.0695 (6)
H1	0.2010	0.4308	0.7289	0.104*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.097 (2)	0.090 (3)	0.0568 (17)	-0.0042 (19)	0.0384 (16)	0.0063 (17)
C2	0.0743 (17)	0.127 (3)	0.082 (2)	0.0100 (19)	0.0428 (18)	0.012 (2)
C3	0.0561 (14)	0.0511 (15)	0.0454 (13)	-0.0092 (13)	0.0189 (11)	0.0004 (12)
C4	0.0608 (16)	0.0587 (17)	0.0431 (13)	0.0034 (14)	0.0143 (12)	0.0113 (13)
C5	0.0487 (13)	0.0557 (16)	0.0487 (14)	0.0064 (12)	0.0148 (11)	0.0060 (13)
C6	0.0510 (14)	0.0410 (14)	0.0418 (13)	-0.0028 (12)	0.0154 (11)	0.0013 (11)
C7	0.0506 (14)	0.0535 (16)	0.0453 (14)	0.0033 (13)	0.0110 (11)	0.0073 (12)
C8	0.0494 (14)	0.0554 (16)	0.0544 (16)	0.0018 (12)	0.0137 (12)	0.0038 (13)
C9	0.0505 (14)	0.0473 (15)	0.0436 (13)	-0.0018 (13)	0.0107 (11)	0.0001 (11)
C10	0.0455 (13)	0.0444 (14)	0.0467 (13)	-0.0009 (11)	0.0116 (10)	-0.0012 (11)
C11	0.0547 (14)	0.0533 (16)	0.0496 (14)	0.0138 (12)	0.0113 (12)	0.0110 (12)

C12	0.0529 (14)	0.066 (2)	0.0604 (16)	0.0188 (13)	0.0188 (13)	0.0095 (14)
C13	0.0533 (14)	0.0548 (16)	0.0530 (14)	0.0039 (13)	0.0207 (11)	0.0075 (13)
C14	0.0543 (15)	0.0523 (16)	0.0506 (15)	0.0101 (13)	0.0167 (12)	0.0147 (13)
C15	0.0403 (12)	0.0443 (14)	0.0554 (15)	0.0053 (11)	0.0129 (11)	0.0050 (12)
C16	0.086 (2)	0.093 (3)	0.081 (2)	0.0042 (18)	0.0414 (19)	0.0168 (16)
C17	0.126 (3)	0.098 (3)	0.097 (3)	0.010 (3)	0.028 (3)	-0.008 (3)
C18	0.080 (2)	0.107 (3)	0.075 (2)	0.0190 (17)	0.0400 (18)	0.0039 (19)
C19	0.094 (3)	0.134 (4)	0.131 (4)	0.014 (3)	0.043 (3)	-0.015 (3)
N1	0.0737 (14)	0.083 (2)	0.0611 (15)	0.0044 (13)	0.0357 (13)	0.0146 (14)
N2	0.0530 (12)	0.0515 (14)	0.0459 (11)	0.0007 (10)	0.0184 (9)	0.0030 (10)
N3	0.0769 (16)	0.112 (2)	0.0696 (16)	0.0288 (15)	0.0381 (13)	0.0210 (15)
O1	0.0666 (12)	0.0777 (14)	0.0737 (14)	0.0302 (11)	0.0356 (10)	0.0283 (11)

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

C1—N1	1.432 (4)	C11—C12	1.363 (4)
C1—H1A	0.9600	C11—H11	0.9300
C1—H1B	0.9600	C12—C13	1.404 (4)
C1—H1C	0.9600	C12—H12	0.9300
C2—N1	1.419 (4)	C13—N3	1.364 (3)
C2—H2A	0.9600	C13—C14	1.400 (3)
C2—H2B	0.9600	C14—C15	1.373 (4)
C2—H2C	0.9600	C14—H14	0.9300
C3—N1	1.377 (3)	C15—O1	1.354 (3)
C3—C8	1.390 (4)	C16—C17	1.451 (5)
C3—C4	1.400 (4)	C16—N3	1.485 (4)
C4—C5	1.370 (4)	C16—H16A	0.9700
C4—H4	0.9300	C16—H16B	0.9700
C5—C6	1.397 (3)	C17—H17A	0.9600
C5—H5	0.9300	C17—H17B	0.9600
C6—C7	1.370 (3)	C17—H17C	0.9600
C6—N2	1.413 (3)	C18—C19	1.454 (5)
C7—C8	1.370 (4)	C18—N3	1.495 (4)
C7—H7	0.9300	C18—H18A	0.9700
C8—H8	0.9300	C18—H18B	0.9700
C9—N2	1.280 (3)	C19—H19A	0.9600
C9—C10	1.433 (3)	C19—H19B	0.9600
C9—H9	0.9300	C19—H19C	0.9600
C10—C11	1.386 (3)	O1—H1	0.8200
C10—C15	1.407 (3)		
N1—C1—H1A	109.5	N3—C13—C14	121.0 (2)
N1—C1—H1B	109.5	N3—C13—C12	121.3 (2)
H1A—C1—H1B	109.5	C14—C13—C12	117.7 (2)
N1—C1—H1C	109.5	C15—C14—C13	120.9 (2)
H1A—C1—H1C	109.5	C15—C14—H14	119.5
H1B—C1—H1C	109.5	C13—C14—H14	119.5
N1—C2—H2A	109.5	O1—C15—C14	118.5 (2)

N1—C2—H2B	109.5	O1—C15—C10	119.9 (2)
H2A—C2—H2B	109.5	C14—C15—C10	121.6 (2)
N1—C2—H2C	109.5	C17—C16—N3	109.7 (3)
H2A—C2—H2C	109.5	C17—C16—H16A	109.7
H2B—C2—H2C	109.5	N3—C16—H16A	109.7
N1—C3—C8	121.2 (2)	C17—C16—H16B	109.7
N1—C3—C4	122.3 (2)	N3—C16—H16B	109.7
C8—C3—C4	116.4 (2)	H16A—C16—H16B	108.2
C5—C4—C3	122.2 (2)	C16—C17—H17A	109.5
C5—C4—H4	118.9	C16—C17—H17B	109.5
C3—C4—H4	118.9	H17A—C17—H17B	109.5
C4—C5—C6	120.4 (2)	C16—C17—H17C	109.5
C4—C5—H5	119.8	H17A—C17—H17C	109.5
C6—C5—H5	119.8	H17B—C17—H17C	109.5
C7—C6—C5	117.3 (2)	C19—C18—N3	111.1 (3)
C7—C6—N2	117.5 (2)	C19—C18—H18A	109.4
C5—C6—N2	125.2 (2)	N3—C18—H18A	109.4
C8—C7—C6	122.6 (2)	C19—C18—H18B	109.4
C8—C7—H7	118.7	N3—C18—H18B	109.4
C6—C7—H7	118.7	H18A—C18—H18B	108.0
C7—C8—C3	121.0 (2)	C18—C19—H19A	109.5
C7—C8—H8	119.5	C18—C19—H19B	109.5
C3—C8—H8	119.5	H19A—C19—H19B	109.5
N2—C9—C10	122.5 (2)	C18—C19—H19C	109.5
N2—C9—H9	118.8	H19A—C19—H19C	109.5
C10—C9—H9	118.8	H19B—C19—H19C	109.5
C11—C10—C15	116.4 (2)	C3—N1—C2	120.8 (2)
C11—C10—C9	121.6 (2)	C3—N1—C1	120.2 (2)
C15—C10—C9	121.9 (2)	C2—N1—C1	118.6 (2)
C12—C11—C10	123.1 (2)	C9—N2—C6	123.9 (2)
C12—C11—H11	118.5	C13—N3—C16	121.5 (3)
C10—C11—H11	118.5	C13—N3—C18	121.1 (3)
C11—C12—C13	120.3 (2)	C16—N3—C18	117.4 (2)
C11—C12—H12	119.8	C15—O1—H1	109.5
C13—C12—H12	119.8		
N1—C3—C4—C5	178.6 (3)	C13—C14—C15—C10	1.1 (4)
C8—C3—C4—C5	-0.3 (4)	C11—C10—C15—O1	177.5 (3)
C3—C4—C5—C6	0.5 (4)	C9—C10—C15—O1	-4.4 (4)
C4—C5—C6—C7	-0.2 (4)	C11—C10—C15—C14	-2.3 (4)
C4—C5—C6—N2	-179.1 (3)	C9—C10—C15—C14	175.8 (3)
C5—C6—C7—C8	-0.3 (4)	C8—C3—N1—C2	-6.4 (5)
N2—C6—C7—C8	178.7 (2)	C4—C3—N1—C2	174.8 (3)
C6—C7—C8—C3	0.5 (4)	C8—C3—N1—C1	-179.2 (3)
N1—C3—C8—C7	-179.0 (3)	C4—C3—N1—C1	2.1 (4)
C4—C3—C8—C7	-0.2 (4)	C10—C9—N2—C6	-177.7 (2)
N2—C9—C10—C11	179.7 (2)	C7—C6—N2—C9	170.0 (2)
N2—C9—C10—C15	1.7 (4)	C5—C6—N2—C9	-11.0 (4)

C15—C10—C11—C12	1.4 (4)	C14—C13—N3—C16	−15.8 (5)
C9—C10—C11—C12	−176.7 (3)	C12—C13—N3—C16	165.5 (3)
C10—C11—C12—C13	0.8 (4)	C14—C13—N3—C18	167.5 (3)
C11—C12—C13—N3	176.7 (3)	C12—C13—N3—C18	−11.3 (5)
C11—C12—C13—C14	−2.1 (4)	C17—C16—N3—C13	97.4 (4)
N3—C13—C14—C15	−177.6 (3)	C17—C16—N3—C18	−85.7 (4)
C12—C13—C14—C15	1.2 (4)	C19—C18—N3—C13	91.4 (4)
C13—C14—C15—O1	−178.8 (3)	C19—C18—N3—C16	−85.5 (4)

Hydrogen-bond geometry (Å, °)

Cg1 and Cg2 are the centroids of rings C3-C8 and C10-C15, respectively.

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···N2	0.82	1.85	2.585 (3)	148
C11—H11···Cg1 ⁱ	0.93	2.71	3.517 (3)	145
C17—H17B···Cg2 ⁱⁱ	0.96	2.90	3.743 (5)	147

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