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Crystal structure of *N*-(1*H*-indol-2-ylmethylidene)-4-methoxyaniline

Masatsugu Taneda,^{a*} Masato Nishi,^a Koji Kubono,^b Yukiyasu Kashiwagi^c and Taisuke Matsumoto^d

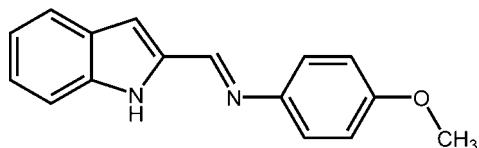
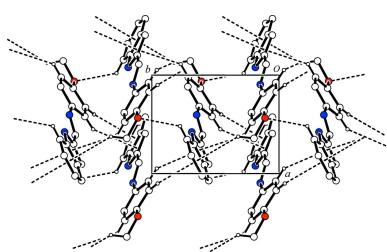
^aDepartment of Science Education, Faculty of Education, Osaka Kyoiku University, Kashiwara, Osaka 582-8582, Japan,

^bDivision of Natural Sciences, Osaka Kyoiku University, Kashiwara, Osaka 582-8582, Japan, ^cOsaka Research Institute of Industrial Science and Technology, 1-6-50 Morinomiya, Joto-ku, Osaka 536-8553, Japan, and ^dInstitute for Materials Chemistry and Engineering, Kyushu University, Kasuga, Fukuoka 816-8580, Japan. *Correspondence e-mail: tane@cc.osaka-kyoiku.ac.jp

The molecule of the title compound, $C_{16}H_{14}N_2O$, contains an essentially planar indole ring system and a phenyl ring. In the crystal, the molecules are linked by a weak intermolecular C—H···O hydrogen bond and C—H···π interactions, forming a one-dimensional column structure along the *b*-axis direction. These columns are linked by other C—H···π interactions, forming a two-dimensional network structure.

1. Chemical context

Indole and its derivatives are useful starting compounds to derive pharmaceutical (Nalli *et al.*, 2020) and biological materials (Arumugam *et al.*, 2021). Indole can function as a hydrogen-bond donor because of the high acidity of the hydrogen atom at position 1. The introduction of a hydrogen-bond acceptor to position 2 of the indole ring forms a five-to-seven-membered intramolecular hydrogen-bonded ring (Nosenko, *et al.*, 2008). In this work, a Schiff base including an indole ring, *N*-(indol-2-ylmethylidene)-4-methoxyaniline, was newly synthesized. Similar Schiff bases such as salicylideneamines often function as bidentate ligands (Wang *et al.*, 2018). Whereas salicylideneamines form intramolecular hydrogen bonds between coordination site atoms, such intramolecular interactions are absent from the crystal structure of the title compound. We report herein on its molecular and crystal structure.



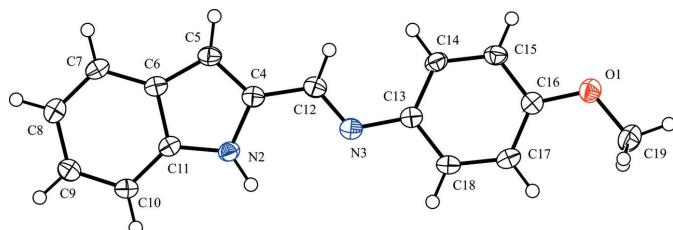
2. Structural commentary

The molecular structure of the title compound is shown in Fig. 1. The C=N double bond adopts an *E* configuration. The indole moiety is almost planar with an r.m.s. deviation of 0.009 (1) Å. The bond lengths and angles in the title molecule are normal and agree with those in other indole imine compounds (IWIGUS; Suresh *et al.*, 2016; KEVLON; Ho *et al.*,



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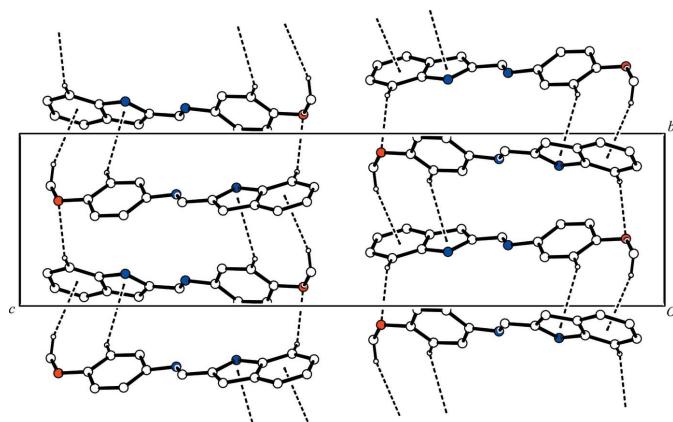
**Figure 1**

The molecular structure of the title compound with atom labelling. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by spheres of arbitrary radius.

2006). The dihedral angle between the indole system and the benzene ring is $9.89(5)^\circ$. In the related compound IWIGUS, the dihedral angles between the indole system and the benzene ring disordered over two sets of sites are widened to $81.8(3)$ and $85.2(3)^\circ$ due to two isopropyl substituents in the benzene ring. There is no intramolecular hydrogen bond in the title compound, because the $\text{N}2-\text{H}2\cdots\text{N}3$ angle is as small as $94.4(10)^\circ$; however, the $\text{N}2\cdots\text{N}3$ distance is $2.8633(16)$ Å, and the $\text{N}2-\text{C}4-\text{C}12-\text{N}3$ torsion angle is $3.94(19)^\circ$. Although no intramolecular hydrogen bond is observed, a broad peak assigned for the $\text{N}-\text{H}$ proton is seen in the ^1H NMR spectrum of the title compound in a CDCl_3 medium and this suggests that the compound forms an intramolecular hydrogen bond in solution (see *Synthesis and crystallization*).

3. Supramolecular features

The title compound contains an $\text{N}-\text{H}$ group, which is a hydrogen-bond donor, and an imino group, which is a hydrogen-bond acceptor, but neither of them forms an intermolecular hydrogen bond in the crystal. Compounds containing a similar indol-2-ylmethyldene-aniline fragment with a *cis*-conformation of the C–C single bond between the N atoms often form dimers by intermolecular $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds (see *Database survey*). However, in the

**Figure 2**

One-dimensional column structure in the crystal of the title compound viewed along the a axis. The $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds and the $\text{C}-\text{H}\cdots\pi$ interactions are shown as dashed lines. H atoms not involved in these interactions are omitted for clarity.

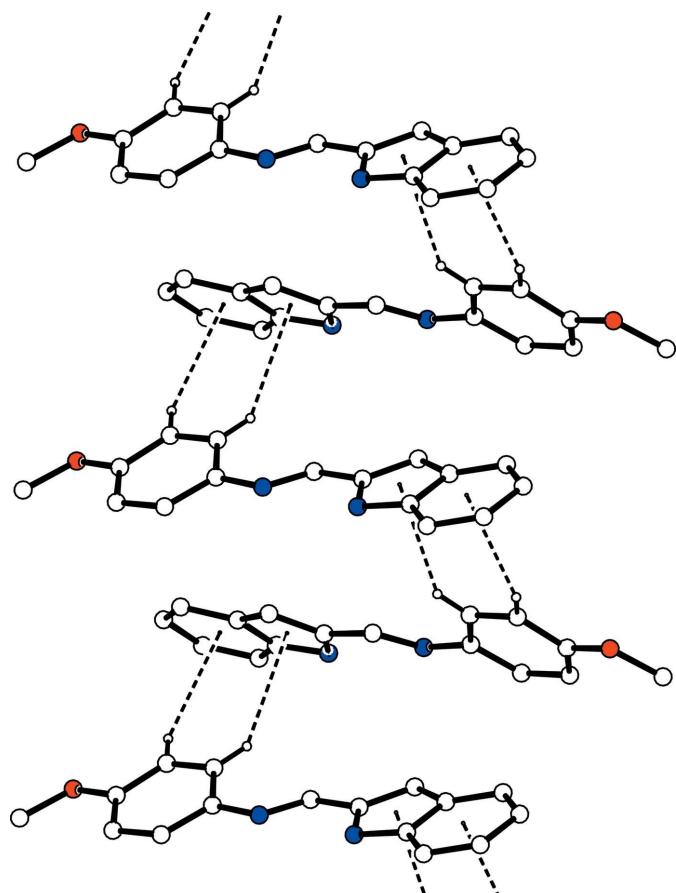
Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$Cg1$ and $Cg2$ are the centroids of the $\text{N}2/\text{C}4-\text{C}6/\text{C}11$ and $\text{C}6-\text{C}11$ rings, respectively.

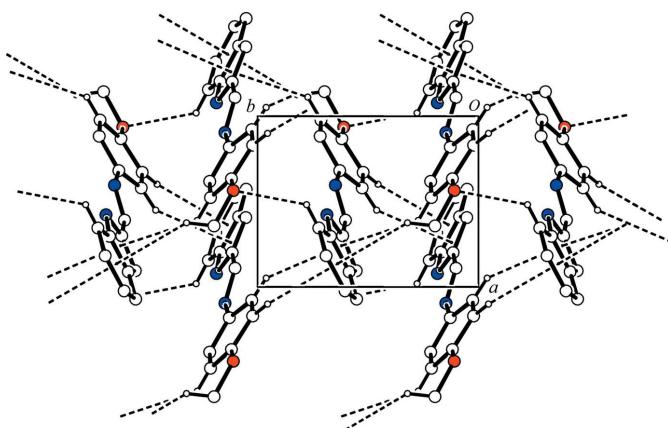
$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C}10-\text{H}10\cdots\text{O}1^i$	0.93	2.58	3.2479 (16)	129
$\text{C}14-\text{H}14\cdots\text{C}g1^{ii}$	0.93	2.81	3.6006 (14)	143
$\text{C}15-\text{H}15\cdots\text{C}g2^{ii}$	0.93	2.79	3.5153 (14)	136
$\text{C}17-\text{H}17\cdots\text{C}g1^i$	0.93	2.89	3.5718 (14)	131
$\text{C}19-\text{H}19\text{C}\cdots\text{C}g2^i$	0.96	2.97	3.7716 (16)	142

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

crystal the molecules of the title compound are linked by a weak intermolecular $\text{C}10-\text{H}10\cdots\text{O}1^i$ hydrogen bond and $\text{C}-\text{H}\cdots\pi$ interactions [$\text{C}17-\text{H}17\cdots\text{C}g1^i$ and $\text{C}19-\text{H}19\text{C}\cdots\text{C}g2^i$; $\text{C}g1$ is the centroid of the $\text{N}2/\text{C}4-\text{C}6/\text{C}11$ ring and $\text{C}g2$ is the centroid of the $\text{C}6-\text{C}11$ ring; symmetry code: (i) $\frac{1}{2} - x, \frac{1}{2} + y, \frac{3}{2} - z$], forming columns along the b -axis direction (Fig. 2, Table 1). Besides this, the molecules belonging to different columns are joined by other $\text{C}-\text{H}\cdots\pi$ interactions [$\text{C}14-\text{H}14\cdots\text{C}g1^{ii}$ and $\text{C}15-\text{H}15\cdots\text{C}g2^{ii}$; symmetry code: (ii) $\frac{3}{2} - x, -\frac{1}{2} + y, \frac{3}{2} - z$] (Fig. 3, Table 1). As a result, the intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds and $\text{C}-\text{H}\cdots\pi$ interactions form a two-dimensional network structure (Fig. 4).

**Figure 3**

Part of the crystal structure of the title compound showing the formation of ribbons along the b -axis direction. The $\text{C}-\text{H}\cdots\pi$ interactions are shown as dashed lines. H atoms not involved in these interactions are omitted for clarity.

**Figure 4**

A packing diagram of the title compound viewed along the c axis, showing the two-dimensional network. The $\text{C}—\text{H}\cdots\text{O}$ hydrogen bonds and $\text{C}—\text{H}\cdots\pi$ interactions are shown as dashed lines. H atoms not involved in these interactions are omitted for clarity.

4. Database survey

A search of the Cambridge Structural Database (CSD, Version 5.42, update of May 2021; Groom *et al.*, 2016) using *ConQuest* (Bruno *et al.*, 2002) for indole derivatives gave 5272 hits, and for the (*1H*-indol-2-yl)methanimine skeleton gave 86 hits. Among these, the imino N atom bonded to an H atom gave one hit, to an N atom gave 24 hits, and to a C atom gave 61 hits. A search for the indol-2-ylmethylidene-aniline fragment gave 30 hits, and those containing a (*1H*-indol-2-yl)methylidene-aniline fragment with a *cis*-conformation of the C—C single bond gave seven hits. These seven compounds include five examples of dimers linked by complementary N—H···N hydrogen bonds (FORJAA; Li *et al.*, 2019; IWIGUS; Suresh *et al.*, 2016; KEZCUQ; Ariyasu *et al.*, 2016; VACKES; Gadekar *et al.*, 2016; WAGCEP; Tian *et al.*, 2016), one example of a one-dimensional-chain structure (UWUSA; Kalalbandi & Seetharamappa, 2016), and one example of a monomer protected from hydrogen bonding by steric hindrance (KEVLON; Ho *et al.*, 2006). These structures contain intermolecular or intramolecular hydrogen bonds involving the N—H or the imino groups. Of these structures, the compounds most closely related to the title compound are *N*-(2,6-diisopropylphenyl)-1-(*1H*-indol-2-yl)methanimine (IWIGUS; Suresh *et al.*, 2016), 4,6-dimethoxy-3-methyl-2,7-bis[(phenyl-imino)methyl]indole (KEVLON; Ho *et al.*, 2006) and 2-(phenyl-*N*-oxidoiminomethyl)-3-phenylaminoindole (CIP-WED; Greci & Sgarabotto, 1984). In the crystal of IWIGUS, which features a large dihedral angle between the indole and benzene rings, two neighbouring molecules are associated through pairs of N—H···N intermolecular hydrogen bonds, forming a centrosymmetric dimer. The crystal structure of an indol-2-ylmethylidene-aniline compound without a hydrogen bond between the N—H and imino groups has not yet been reported. In an almost planar molecule without a bulky substituent such as the tile compound, the formation of a dimer by intermolecular N—H···N hydrogen bonding is probably not appropriate for the crystal packing.

Table 2
Experimental details.

Crystal data	$\text{C}_{16}\text{H}_{14}\text{N}_2\text{O}$
Chemical formula	$\text{C}_{16}\text{H}_{14}\text{N}_2\text{O}$
M_r	250.29
Crystal system, space group	Monoclinic, $P2_1/n$
Temperature (K)	123
a, b, c (Å)	5.87685 (19), 7.5999 (3), 28.4578 (11)
β (°)	90.604 (3)
V (Å ³)	1270.95 (8)
Z	4
Radiation type	Mo $K\alpha$
μ (mm ⁻¹)	0.08
Crystal size (mm)	0.30 × 0.20 × 0.10
Data collection	
Diffractometer	Rigaku AFC10 Saturn70 area detector
Absorption correction	Multi-scan <i>CrysAlis PRO</i> ; Rigaku OD, 2018)
T_{\min}, T_{\max}	0.608, 0.992
No. of measured, independent and observed [$F^2 > 2.0\sigma(F^2)$] reflections	11128, 2907, 2525
R_{int}	0.056
(sin θ/λ) _{max} (Å ⁻¹)	0.649
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.049, 0.127, 1.05
No. of reflections	2907
No. of parameters	178
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)	0.39, -0.29

Computer programs: *CrysAlis PRO* (Rigaku OD, 2018), *SIR92* (Altomare *et al.*, 1993), *SHELXL2018/3* (Sheldrick, 2015), *PLATON* (Spek, 2020) and *CrystalStructure* (Rigaku, 2019).

5. Synthesis and crystallization

Indole-2-carbaldehyde (145 mg, 1.00 mmol) and *p*-anisidine (148 mg, 1.20 mmol) were dissolved in toluene (20 mL), and the solution was refluxed under inert gas for 6 h, followed by evaporation. The residue was purified by recrystallization from a solvent mixture of acetone and *n*-hexane (1:1), and the title compound was then obtained (212 mg, 0.848 mmol, 84.8%) as a pale-red powder. The recrystallization of the title compound from a mixture of acetone and methanol afforded single crystals suitable for X-ray structure analysis. ¹H NMR (CDCl_3 , 400 MHz) δ = 3.84 (*s*, 3H, OCH₃), 6.93–6.97 (*m*, 3H, ArH), 7.13 (*td*, 1H, $J_{\text{ortho}} = 7.5$ Hz, $J_{\text{meta}} = 1.0$ Hz, ArH), 7.25–7.31 (*m*, 3H, ArH), 7.40 (*dd*, 1H, $J_{\text{ortho}} = 8.3$ Hz, $J_{\text{meta}} = 0.9$ Hz, ArH), 7.66 (*d*, 1H, $J_{\text{ortho}} = 8.0$ Hz, ArH), 8.48 (*s*, 1H, N=CH), 9.25 (*br*, 1H, NH). HR-MS (*m/z*): calculated for [C₁₆H₁₅N₂O]⁺, *m/z* = 251.1179; found, 251.1192.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The H atom attached to N2 was located in a difference-Fourier map and freely refined. The C-bound H atoms were positioned geometrically and refined using a riding model: C—H = 0.93–0.96 Å with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

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Crystal structure of *N*-(1*H*-indol-2-ylmethylidene)-4-methoxyaniline

Masatsugu Taneda, Masato Nishi, Koji Kubono, Yukiyasu Kashiwagi and Taisuke Matsumoto

Computing details

Data collection: *CrysAlis PRO* (Rigaku OD, 2018); cell refinement: *CrysAlis PRO* (Rigaku OD, 2018); data reduction: *CrysAlis PRO* (Rigaku OD, 2018); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2015); molecular graphics: *PLATON* (Spek, 2020); software used to prepare material for publication: *CrystalStructure* (Rigaku, 2019).

N-(1*H*-Indol-2-ylmethylidene)-4-methoxyaniline

Crystal data

$C_{16}H_{14}N_2O$
 $M_r = 250.29$
Monoclinic, $P2_1/n$
 $a = 5.87685 (19)$ Å
 $b = 7.5999 (3)$ Å
 $c = 28.4578 (11)$ Å
 $\beta = 90.604 (3)^\circ$
 $V = 1270.95 (8)$ Å³
 $Z = 4$

$F(000) = 528.00$
 $D_x = 1.308 \text{ Mg m}^{-3}$
 $Mo K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 6329 reflections
 $\theta = 2.8\text{--}30.8^\circ$
 $\mu = 0.08 \text{ mm}^{-1}$
 $T = 123$ K
Prism, colourless
 $0.30 \times 0.20 \times 0.10$ mm

Data collection

Rigaku AFC10 Saturn70 area detector
dифрактометр
Radiation source: rotating anode X-ray
generator, micromax007
Multi-layer mirror optics monochromator
Detector resolution: 28.5714 pixels mm⁻¹
 ω scans
Absorption correction: multi-scan
CrysAlisPro; Rigaku OD, 2018)

$T_{\min} = 0.608$, $T_{\max} = 0.992$
11128 measured reflections
2907 independent reflections
2525 reflections with $F^2 > 2.0\sigma(F^2)$
 $R_{\text{int}} = 0.056$
 $\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 2.8^\circ$
 $h = -7 \rightarrow 7$
 $k = -8 \rightarrow 9$
 $l = -36 \rightarrow 35$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.127$
 $S = 1.05$
2907 reflections
178 parameters
0 restraints
Primary atom site location: structure-invariant
direct methods
Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites
H atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0686P)^2 + 0.3253P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.39 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.29 \text{ e } \text{\AA}^{-3}$
Extinction correction: SHELXL
Extinction coefficient: 0.259 (14)

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 was performed using all reflections. The weighted R-factor (wR) and goodness of fit (S) are based on F^2 . R-factor (gt) are based on F . The threshold expression of $F^2 > 2.0 \text{ sigma}(F^2)$ is used only for calculating R-factor (gt).

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	0.06455 (16)	0.60985 (13)	0.93970 (3)	0.0292 (3)
N2	0.57779 (18)	0.68608 (14)	0.66418 (4)	0.0237 (3)
N3	0.40410 (18)	0.64748 (14)	0.75724 (4)	0.0248 (3)
C4	0.7005 (2)	0.61479 (16)	0.70151 (4)	0.0240 (3)
C5	0.9058 (2)	0.55340 (17)	0.68542 (5)	0.0249 (3)
H5	1.0199	0.5007	0.7034	0.030*
C6	0.9115 (2)	0.58536 (16)	0.63609 (4)	0.0221 (3)
C7	1.0717 (2)	0.55168 (17)	0.60070 (5)	0.0253 (3)
H7	1.2079	0.4950	0.6079	0.030*
C8	1.0237 (2)	0.60388 (17)	0.55530 (5)	0.0272 (3)
H8	1.1287	0.5822	0.5318	0.033*
C9	0.8174 (2)	0.68975 (17)	0.54398 (5)	0.0266 (3)
H9	0.7899	0.7250	0.5131	0.032*
C10	0.6548 (2)	0.72278 (17)	0.57776 (4)	0.0253 (3)
H10	0.5182	0.7780	0.5701	0.030*
C11	0.7037 (2)	0.67011 (16)	0.62387 (4)	0.0219 (3)
C12	0.6095 (2)	0.60207 (17)	0.74822 (4)	0.0245 (3)
H12	0.7019	0.5599	0.7724	0.029*
C13	0.3193 (2)	0.62794 (16)	0.80344 (4)	0.0231 (3)
C14	0.4210 (2)	0.52631 (17)	0.83908 (5)	0.0264 (3)
H14	0.5511	0.4612	0.8327	0.032*
C15	0.3296 (2)	0.52215 (17)	0.88355 (5)	0.0266 (3)
H15	0.3993	0.4548	0.9069	0.032*
C16	0.1332 (2)	0.61828 (16)	0.89388 (4)	0.0233 (3)
C17	0.0233 (2)	0.71292 (17)	0.85845 (5)	0.0253 (3)
H17	-0.1111	0.7729	0.8645	0.030*
C18	0.1179 (2)	0.71653 (17)	0.81370 (5)	0.0247 (3)
H18	0.0446	0.7798	0.7900	0.030*
C19	-0.1427 (3)	0.6969 (2)	0.95149 (5)	0.0367 (4)
H19A	-0.1793	0.6722	0.9836	0.044*
H19B	-0.2632	0.6554	0.9313	0.044*
H19C	-0.1247	0.8215	0.9474	0.044*
H2	0.438 (3)	0.737 (2)	0.6678 (5)	0.032 (4)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0294 (5)	0.0304 (5)	0.0279 (5)	0.0010 (4)	0.0024 (4)	-0.0028 (4)
N2	0.0196 (5)	0.0258 (6)	0.0258 (5)	0.0019 (4)	-0.0007 (4)	-0.0004 (4)
N3	0.0257 (5)	0.0218 (6)	0.0270 (6)	-0.0006 (4)	-0.0012 (4)	0.0001 (4)
C4	0.0223 (6)	0.0226 (6)	0.0270 (7)	-0.0032 (5)	-0.0028 (5)	-0.0007 (5)
C5	0.0216 (6)	0.0242 (6)	0.0288 (7)	-0.0008 (5)	-0.0046 (5)	0.0014 (5)
C6	0.0194 (6)	0.0180 (6)	0.0289 (7)	-0.0027 (4)	-0.0030 (5)	-0.0015 (5)
C7	0.0192 (6)	0.0228 (6)	0.0339 (7)	0.0008 (5)	-0.0006 (5)	-0.0029 (5)
C8	0.0247 (6)	0.0259 (7)	0.0310 (7)	-0.0034 (5)	0.0029 (5)	-0.0056 (5)
C9	0.0289 (6)	0.0249 (7)	0.0258 (6)	-0.0022 (5)	-0.0032 (5)	-0.0005 (5)
C10	0.0235 (6)	0.0238 (6)	0.0283 (7)	0.0021 (5)	-0.0040 (5)	0.0000 (5)
C11	0.0194 (6)	0.0189 (6)	0.0273 (6)	-0.0015 (5)	-0.0009 (4)	-0.0023 (5)
C12	0.0237 (6)	0.0239 (7)	0.0259 (6)	-0.0032 (5)	-0.0042 (5)	-0.0007 (5)
C13	0.0221 (6)	0.0200 (6)	0.0270 (6)	-0.0023 (5)	-0.0017 (5)	-0.0005 (5)
C14	0.0216 (6)	0.0242 (7)	0.0335 (7)	0.0047 (5)	0.0029 (5)	0.0042 (5)
C15	0.0250 (6)	0.0243 (7)	0.0305 (7)	0.0019 (5)	-0.0022 (5)	0.0053 (5)
C16	0.0239 (6)	0.0192 (6)	0.0266 (6)	-0.0043 (5)	-0.0009 (5)	-0.0025 (5)
C17	0.0206 (6)	0.0212 (6)	0.0340 (7)	0.0012 (5)	-0.0017 (5)	-0.0023 (5)
C18	0.0227 (6)	0.0222 (6)	0.0289 (7)	0.0009 (5)	-0.0054 (5)	0.0016 (5)
C19	0.0341 (8)	0.0398 (9)	0.0364 (8)	0.0040 (6)	0.0069 (6)	-0.0064 (7)

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

O1—C16	1.3705 (15)	C9—H9	0.9300
O1—C19	1.4287 (17)	C10—C11	1.3989 (17)
N2—C11	1.3769 (16)	C10—H10	0.9300
N2—C4	1.3876 (16)	C12—H12	0.9300
N2—H2	0.912 (17)	C13—C18	1.3950 (17)
N3—C12	1.2840 (17)	C13—C14	1.4035 (18)
N3—C13	1.4189 (17)	C14—C15	1.3803 (18)
C4—C5	1.3768 (18)	C14—H14	0.9300
C4—C12	1.4413 (18)	C15—C16	1.4000 (18)
C5—C6	1.4252 (18)	C15—H15	0.9300
C5—H5	0.9300	C16—C17	1.3918 (18)
C6—C7	1.4094 (18)	C17—C18	1.3950 (18)
C6—C11	1.4205 (17)	C17—H17	0.9300
C7—C8	1.3781 (19)	C18—H18	0.9300
C7—H7	0.9300	C19—H19A	0.9600
C8—C9	1.4111 (18)	C19—H19B	0.9600
C8—H8	0.9300	C19—H19C	0.9600
C9—C10	1.3852 (19)		
C16—O1—C19	117.50 (11)	C10—C11—C6	121.79 (12)
C11—N2—C4	108.89 (10)	N3—C12—C4	121.67 (12)
C11—N2—H2	128.4 (10)	N3—C12—H12	119.2
C4—N2—H2	122.7 (10)	C4—C12—H12	119.2

C12—N3—C13	119.83 (11)	C18—C13—C14	118.05 (12)
C5—C4—N2	109.15 (11)	C18—C13—N3	116.78 (11)
C5—C4—C12	128.24 (12)	C14—C13—N3	125.17 (12)
N2—C4—C12	122.51 (11)	C15—C14—C13	120.54 (12)
C4—C5—C6	107.42 (11)	C15—C14—H14	119.7
C4—C5—H5	126.3	C13—C14—H14	119.7
C6—C5—H5	126.3	C14—C15—C16	120.68 (12)
C7—C6—C11	119.10 (12)	C14—C15—H15	119.7
C7—C6—C5	134.07 (12)	C16—C15—H15	119.7
C11—C6—C5	106.83 (11)	O1—C16—C17	125.06 (12)
C8—C7—C6	119.08 (12)	O1—C16—C15	115.28 (11)
C8—C7—H7	120.5	C17—C16—C15	119.65 (12)
C6—C7—H7	120.5	C16—C17—C18	119.03 (12)
C7—C8—C9	120.94 (12)	C16—C17—H17	120.5
C7—C8—H8	119.5	C18—C17—H17	120.5
C9—C8—H8	119.5	C13—C18—C17	121.91 (11)
C10—C9—C8	121.49 (12)	C13—C18—H18	119.0
C10—C9—H9	119.3	C17—C18—H18	119.0
C8—C9—H9	119.3	O1—C19—H19A	109.5
C9—C10—C11	117.58 (12)	O1—C19—H19B	109.5
C9—C10—H10	121.2	H19A—C19—H19B	109.5
C11—C10—H10	121.2	O1—C19—H19C	109.5
N2—C11—C10	130.50 (11)	H19A—C19—H19C	109.5
N2—C11—C6	107.71 (11)	H19B—C19—H19C	109.5
C11—N2—C4—C5	0.24 (14)	C5—C6—C11—C10	179.05 (11)
C11—N2—C4—C12	-176.34 (11)	C13—N3—C12—C4	178.13 (11)
N2—C4—C5—C6	-0.69 (14)	C5—C4—C12—N3	-171.94 (13)
C12—C4—C5—C6	175.63 (12)	N2—C4—C12—N3	3.94 (19)
C4—C5—C6—C7	-179.42 (14)	C12—N3—C13—C18	165.19 (12)
C4—C5—C6—C11	0.87 (14)	C12—N3—C13—C14	-15.23 (19)
C11—C6—C7—C8	0.81 (18)	C18—C13—C14—C15	-3.23 (19)
C5—C6—C7—C8	-178.88 (13)	N3—C13—C14—C15	177.21 (12)
C6—C7—C8—C9	-0.08 (19)	C13—C14—C15—C16	0.4 (2)
C7—C8—C9—C10	-0.8 (2)	C19—O1—C16—C17	3.66 (18)
C8—C9—C10—C11	0.89 (19)	C19—O1—C16—C15	-176.29 (12)
C4—N2—C11—C10	-179.44 (13)	C14—C15—C16—O1	-177.25 (11)
C4—N2—C11—C6	0.32 (14)	C14—C15—C16—C17	2.80 (19)
C9—C10—C11—N2	179.59 (12)	O1—C16—C17—C18	177.07 (11)
C9—C10—C11—C6	-0.14 (19)	C15—C16—C17—C18	-2.98 (18)
C7—C6—C11—N2	179.51 (11)	C14—C13—C18—C17	3.04 (19)
C5—C6—C11—N2	-0.73 (13)	N3—C13—C18—C17	-177.36 (11)
C7—C6—C11—C10	-0.71 (19)	C16—C17—C18—C13	0.05 (19)

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

Cg1 and Cg2 are the centroids of the N2/C4–C6/C11 and C6–C11 rings, respectively.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C10—H10···O1 ⁱ	0.93	2.58	3.2479 (16)	129
C14—H14···Cg1 ⁱⁱ	0.93	2.81	3.6006 (14)	143
C15—H15···Cg2 ⁱⁱ	0.93	2.79	3.5153 (14)	136
C17—H17···Cg1 ⁱ	0.93	2.89	3.5718 (14)	131
C19—H19C···Cg2 ⁱ	0.96	2.97	3.7716 (16)	142

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