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Crystal structure of 2-{5-[2-(2-hydroxyphenyl)diazen-1-yl]-1-methylpyrrol-2-yl}phenol methanol monosolvate

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In the title azopyrrole compound, $C_{17}H_{15}N_3O_2 \cdot CH_3OH$, the azo N=N bond adopts a *trans* configuration and the pyrrole N and azo group are in an *anti* orientation. The dihedral angles between the pyrrole ring and the two phenyl rings are 6.7 (3) and 54.7 (3)°. In the crystal, a supramolecular ring structure is formed between two azopyrrole and two methanol solvent molecules through four $O-H \cdots O$ hydrogen bonds.

1. Chemical context

2. Structural commentary

Recently, azopyrrole dyes have received much attention for their promising use in the design of advanced materials and devices. For example, some thienylpyrrole azo dyes bearing heterocyclic groups have good non-linear optical properties (Raposo *et al.*, 2011). Mikroyannidis and coworkers found that many azopyrrole dyes are efficient bulk heterojunction solar cell materials (Sharma *et al.*, 2012). In a previous work, we reported the crystal engineering of some 5,5'-bis(phenyldiazo)dipyrromethane compounds and demonstrated their interlocked type self-assemblies in the solid state *via* quadruple N-H···N hydrogen bonds (Yin *et al.*, 2008, 2009). In a continuation of this research, we report herein the crystal structure of 2-{5-[2-(2-hydroxyphenyl)diazen-1-yl]-1-methylpyrrol-2-yl}phenol methanol monosolvate.



The structure of the title compound is shown in Fig. 1. The asymmetric unit contains one azopyrrole molecule and one methanol solvent molecule. The azoylazopyrrole group is almost planar, reflected by the dihedral angle between the pyrrole ring (N3/C7–C10) and the benzene ring (C1–C6) of

only 6.7 $(3)^{\circ}$, which may be due to the existence of the intra-





research communications



Figure 1

ORTEP diagram for the title compound, with displacement ellipsoids drawn at the 30% probability level. The methanol solvent molecule was omitted for clarity.

molecular O1-H1···N2 hydrogen bond (Table 1) between the hydroxy group and the azo N atom. The dihedral angle between pyrrole ring and the other benzene ring (C12-C17) is 54.7 (3)°, which may be caused by the steric repulsion between hydroxy group and methyl group. The azo N=N bond adopts a *trans* configuration and its length is 1.286 (2) Å, which is shorter than that in the crystal of 2,5-bis(2-hydroxyphenylazo)-1*H*-pyrrole (1.293 Å; Li *et al.* 2009). It is worth mentioning that the N1 atom of the azo group and the N3 atom of the pyrrole ring are arranged on opposite sides with respect to the C7-N2 bond, which is the same as in the crystal of 2-phenylazo-1-vinyl pyrrole (Trofimov *et al.*, 2006) but different to many other observations (Li *et al.*, 2009; Yin *et al.*, 2008). The bond lengths in the pyrrole ring are more equal compared to those normally observed.

3. Supramolecular features

In the crystal, two azopyrrole molecules are bridged by two methanol solvent molecules through four $O-H\cdots O$



Figure 2 O-H···O hydrogen-bonded (Table 1) supramolecular ring structure.

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

Cg1 and Cg2 are the centroids of the C1–C6 and C12-C17 rings, respectively

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O1−H1···N2	0.84	1.81	2.530 (2)	143
$O2-H2\cdots O3^{i}$	0.84	1.81	2.641 (2)	171
$O3-H3A\cdots O1^{ii}$	0.84	1.97	2.763 (2)	157
$C11 - H11B \cdots Cg1^{iii}$	0.98	2.73	3.587 (2)	147
$C18 - H18C \cdots Cg2^{iv}$	0.98	2.75	3.483 (3)	132

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

hydrogen bonds forming a large supramolecular ring structure, in which the methanol acts as both a hydrogen-bond acceptor and a donor (Fig. 2, Table 1). This type of coordination environment is most populated (occupying 70%) for methanol molecules as revealed by a search of the Cambridge Structural Database (CSD) (Brychczynska *et al.* 2008). The methyl groups point to the inside of the ring. The rings are further held together through $C-H\cdots\pi$ contacts involving the benzene rings (Table 1). There are no $\pi-\pi$ interactions between the aromatic rings. The packing is shown in Fig. 3.

4. Database survey

A search in the Cambridge Structural Database (Version 5.38; Groom *et al.*, 2016) returned 45 entries for azopyrrole derivatives, including three entries for *N*-vinylphenylazopyrrole (Trofimov *et al.*, 2006; Rusakov *et al.*, 2007), four entries for mono- or bisazopyrroles (Li *et al.*, 2009), two entries for azo calix[4]pyrroles (Nishiyabu *et al.*, 2006), five entries for pyrrole-azocrown ethers (Wagner-Wysiecka *et al.*, 2011; Szczygelska-Tao *et al.*, 2008), two entries for azopyrrole boron difluoride complexes (Li *et al.*, 2009; Lee *et al.*, 2012), ten entries for metal complexes (Li *et al.*, 2008; Li, & Dolphin, 2011; Yin *et al.*, 2012; Zhang *et al.*, 2015; Ghorui *et al.*, 2016);



Figure 3 A view of the crystal packing along the *b* axis.

5. Synthesis and crystallization

A 273 K solution of 2-aminophenol 0.272 g (2.5 mmol) and aqueous HCl (2 mL) in water (2 mL) was treated with another 273 K solution of NaNO₂ (0.18 g, 2.5 mmol) in 3 mL water, and the mixture was stirred at 273 K for 30 min. The diazonium salt solution was added dropwise to a solution of *N*methylpyrrole (81 mg, 1 mmol) in acetonitrile (25 mL) and three drops of acetic acid. The combined solution was maintained at 273 K for 2 h with stirring. After that, EtOAc (25 mL) and water (25 mL) were added. The organic layer was separated and washed with water (20 mL) and dried with anhydrous MgSO₄. The solution was evaporated and the residue was purified by column chromatography on silica (ethyl acetate/petroleum ether = 1:2), which gave the title compound as an orange powder (200 mg, 68%, m.p. 404 K).

¹H NMR (400MHz, DMSO-*d*₆): δ 3.73 (*s*, 3H, –CH₃), 6.32 (*d*, *J* = 4 Hz, 1H, pyrrole C–H), 6.86 (*d*, *J* = 4 Hz, 1H, pyrrole C–H), 6.90–6.95 (*m*, 2H, Ar C-H), 6.99 (*t*, 2H, *J* = 8 Hz, Ar C–H), 7.23–7.31 (*m*, 3H, Ar C-H), 7.65 (*d*, *J* = 8Hz, 1H, Ar C–H), 9.95 (*s*, 1H, –OH), 10.43 (*s*, 1H, –OH). Crystals suitable X-ray diffraction analysis were obtained by the slow evaporation of a CHCl₃/CH₃OH solution of the title compound.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. O–H atoms were located in a difference-Fourier map and refined freely. Other H atoms were positioned geometrically (C–H = 0.95 or 0.98 Å) and included in the final cycles of refinement using a riding model, with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(C)$.

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Table	2	
Experi	mental	details.

Crystal data	
Chemical formula	$C_{17}H_{15}N_3O_2 \cdot CH_4O$
M _r	325.36
Crystal system, space group	Triclinic, $P\overline{1}$
Temperature (K)	173
a, b, c (Å)	7.1597 (11), 9.8762 (16), 12.243 (2)
α, β, γ (°)	110.426 (3), 94.051 (3), 93.378 (3)
$V(\dot{A}^3)$	806.0 (2)
Z	2
Radiation type	Μο Κα
$\mu (\text{mm}^{-1})$	0.09
Crystal size (mm)	$0.14 \times 0.13 \times 0.12$
Data collection	
Diffractometer	Bruker APEXII CCD
Absorption correction	Multi-scan (<i>SADABS</i> ; Bruker, 2005)
T_{\min}, T_{\max}	0.987, 0.989
No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections	4168, 2830, 2071
R _{int}	0.025
$(\sin \theta / \lambda)_{\rm max} ({\rm \AA}^{-1})$	0.595
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.045, 0.109, 1.03
No. of reflections	2830
No. of parameters	222
H-atom treatment	H-atom parameters constrained
$\Delta \rho_{\rm max}, \Delta \rho_{\rm min} ({\rm e} {\rm \AA}^{-3})$	0.21, -0.23

Computer programs: *APEX2* and *SAINT* (Bruker, 2005), *SHELXS97*, *SHELXL97* and *SHELXTL* (Sheldrick, 2008).

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Crystal structure of 2-{5-[2-(2-hydroxyphenyl)diazen-1-yl]-1-methylpyrrol-2yl}phenol methanol monosolvate

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Computing details

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT* (Bruker, 2005); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

2-{5-[2-(2-Hydroxyphenyl)diazen-1-yl]-1-methylpyrrol-2-yl}phenol methanol monosolvate

Crystal data

 $C_{17}H_{15}N_{3}O_{2} \cdot CH_{4}O$ $M_{r} = 325.36$ Triclinic, *P*1 *a* = 7.1597 (11) Å *b* = 9.8762 (16) Å *c* = 12.243 (2) Å *a* = 110.426 (3)° *β* = 94.051 (3)° *γ* = 93.378 (3)° *V* = 806.0 (2) Å³

Data collection

Bruker APEXII CCD diffractometer Radiation source: fine-focus sealed tube Graphite monochromator φ and ω scans Absorption correction: multi-scan (SADABS; Bruker, 2005) $T_{\min} = 0.987, T_{\max} = 0.989$

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.109$ S = 1.032830 reflections 222 parameters 0 restraints Z = 2 F(000) = 344 $D_x = 1.341 \text{ Mg m}^{-3}$ Mo K α radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 1104 reflections $\theta = 2.3-28.1^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 173 KBLOCK, yellow $0.14 \times 0.13 \times 0.12 \text{ mm}$

4168 measured reflections 2830 independent reflections 2071 reflections with $I > 2\sigma(I)$ $R_{int} = 0.025$ $\theta_{max} = 25.0^{\circ}, \theta_{min} = 1.8^{\circ}$ $h = -8 \rightarrow 8$ $k = -11 \rightarrow 9$ $l = -14 \rightarrow 14$

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 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0437P)^{2} + 0.2373P] \qquad \Delta \rho_{max} = 0.21 \text{ e } \text{\AA}^{-3}$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3 \qquad \Delta \rho_{min} = -0.23 \text{ e } \text{\AA}^{-3}$ $(\Delta/\sigma)_{max} < 0.001$

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 $F^2 > 2sigma(F^2)$ 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 an	d isotropic or e	quivalent isotrop	oic displacement	parameters	$(Å^2)$)
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	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
01	0.2218 (2)	0.53355 (15)	-0.07978 (12)	0.0254 (4)	
H1	0.2260	0.5941	-0.0113	0.038*	
O2	0.4830(2)	1.13606 (14)	0.27561 (13)	0.0221 (4)	
H2	0.5527	1.1998	0.2636	0.033*	
O3	0.2725 (2)	0.68674 (17)	0.77244 (15)	0.0304 (4)	
H3A	0.2900	0.6375	0.8154	0.046*	
N1	0.2920 (2)	0.48800 (18)	0.14369 (15)	0.0191 (4)	
N2	0.2614 (2)	0.61395 (18)	0.14132 (14)	0.0181 (4)	
N3	0.2453 (2)	0.86125 (17)	0.24967 (14)	0.0166 (4)	
C1	0.2840 (3)	0.3796 (2)	0.03169 (18)	0.0180 (5)	
C2	0.2539 (3)	0.4013 (2)	-0.07572 (18)	0.0196 (5)	
C3	0.2592 (3)	0.2857 (2)	-0.18007 (19)	0.0247 (5)	
H3	0.2390	0.3001	-0.2526	0.030*	
C4	0.2933 (3)	0.1506 (2)	-0.17897 (19)	0.0243 (5)	
H4	0.2988	0.0728	-0.2508	0.029*	
C5	0.3199 (3)	0.1269 (2)	-0.07380 (19)	0.0235 (5)	
Н5	0.3407	0.0329	-0.0735	0.028*	
C6	0.3159 (3)	0.2406 (2)	0.02965 (19)	0.0206 (5)	
H6	0.3353	0.2244	0.1015	0.025*	
C7	0.2752 (3)	0.7239 (2)	0.24820 (17)	0.0170 (5)	
C8	0.3197 (3)	0.7316 (2)	0.36227 (17)	0.0190 (5)	
H8	0.3478	0.6533	0.3870	0.023*	
C9	0.3152 (3)	0.8753 (2)	0.43346 (18)	0.0190 (5)	
H9	0.3386	0.9128	0.5164	0.023*	
C10	0.2708 (3)	0.9552 (2)	0.36319 (17)	0.0172 (5)	
C11	0.1888 (3)	0.8954 (2)	0.14600 (18)	0.0210 (5)	
H11A	0.1466	0.9933	0.1702	0.032*	
H11B	0.0858	0.8251	0.0986	0.032*	
H11C	0.2959	0.8908	0.0997	0.032*	
C12	0.2552 (3)	1.1127 (2)	0.39959 (17)	0.0167 (5)	
C13	0.3645 (3)	1.2011 (2)	0.35662 (17)	0.0166 (5)	
C14	0.3540 (3)	1.3501 (2)	0.39739 (17)	0.0184 (5)	

H14	0.4266	1.4092	0.3664	0.022*	
C15	0.2372 (3)	1.4124 (2)	0.48346 (18)	0.0203 (5)	
H15	0.2299	1.5144	0.5113	0.024*	
C16	0.1314 (3)	1.3269 (2)	0.52895 (18)	0.0211 (5)	
H16	0.0526	1.3699	0.5886	0.025*	
C17	0.1411 (3)	1.1790 (2)	0.48707 (18)	0.0201 (5)	
H17	0.0682	1.1207	0.5187	0.024*	
C18	0.0860 (3)	0.7274 (3)	0.7761 (2)	0.0310 (6)	
H18A	0.0712	0.7970	0.8541	0.046*	
H18B	-0.0020	0.6411	0.7594	0.046*	
H18C	0.0596	0.7720	0.7173	0.046*	

Atomic displacement parameters $(Å^2)$

	U^{11}	U ²²	U ³³	U ¹²	U ¹³	U^{23}
01	0.0358 (9)	0.0192 (8)	0.0215 (8)	0.0050 (7)	0.0042 (7)	0.0068 (7)
O2	0.0216 (8)	0.0161 (8)	0.0286 (8)	0.0020 (6)	0.0093 (7)	0.0066 (7)
O3	0.0220 (8)	0.0362 (10)	0.0448 (10)	0.0055 (7)	0.0064 (7)	0.0279 (8)
N1	0.0188 (9)	0.0144 (9)	0.0233 (10)	0.0014 (7)	0.0030 (8)	0.0053 (8)
N2	0.0164 (9)	0.0155 (10)	0.0216 (9)	0.0004 (7)	0.0024 (7)	0.0057 (8)
N3	0.0157 (9)	0.0155 (9)	0.0188 (9)	0.0018 (7)	0.0022 (7)	0.0062 (7)
C1	0.0120 (10)	0.0167 (11)	0.0226 (11)	0.0005 (8)	0.0008 (9)	0.0040 (9)
C2	0.0165 (11)	0.0176 (12)	0.0253 (12)	0.0020 (9)	0.0015 (9)	0.0083 (9)
C3	0.0235 (12)	0.0296 (13)	0.0199 (11)	0.0021 (10)	0.0022 (10)	0.0072 (10)
C4	0.0209 (11)	0.0210 (12)	0.0236 (12)	0.0036 (9)	0.0026 (9)	-0.0015 (10)
C5	0.0212 (12)	0.0160 (12)	0.0304 (13)	0.0036 (9)	0.0001 (10)	0.0049 (10)
C6	0.0172 (11)	0.0211 (12)	0.0230 (11)	0.0008 (9)	-0.0021 (9)	0.0080 (9)
C7	0.0152 (10)	0.0136 (11)	0.0215 (11)	0.0004 (8)	0.0032 (9)	0.0052 (9)
C8	0.0222 (11)	0.0143 (11)	0.0216 (11)	0.0022 (9)	0.0025 (9)	0.0077 (9)
С9	0.0198 (11)	0.0184 (11)	0.0179 (11)	0.0012 (9)	0.0031 (9)	0.0051 (9)
C10	0.0143 (10)	0.0160 (11)	0.0195 (11)	0.0004 (8)	0.0028 (9)	0.0042 (9)
C11	0.0239 (12)	0.0176 (11)	0.0198 (11)	0.0019 (9)	-0.0034 (9)	0.0054 (9)
C12	0.0151 (10)	0.0142 (11)	0.0197 (11)	0.0000 (8)	-0.0022 (9)	0.0055 (9)
C13	0.0148 (10)	0.0174 (11)	0.0160 (10)	0.0031 (8)	-0.0001 (8)	0.0040 (9)
C14	0.0181 (11)	0.0154 (11)	0.0211 (11)	-0.0006 (8)	-0.0010 (9)	0.0066 (9)
C15	0.0210 (11)	0.0141 (11)	0.0227 (11)	0.0027 (9)	-0.0009 (9)	0.0031 (9)
C16	0.0206 (11)	0.0200 (12)	0.0216 (11)	0.0047 (9)	0.0048 (9)	0.0050 (9)
C17	0.0171 (11)	0.0202 (12)	0.0239 (11)	-0.0005 (9)	0.0028 (9)	0.0090 (9)
C18	0.0216 (12)	0.0363 (14)	0.0367 (14)	0.0030 (10)	0.0015 (11)	0.0153 (12)

Geometric parameters (Å, °)

01—C2	1.357 (2)	С7—С8	1.385 (3)	
01—H1	0.8400	C8—C9	1.386 (3)	
O2—C13	1.362 (2)	C8—H8	0.9500	
O2—H2	0.8400	C9—C10	1.389 (3)	
O3—C18	1.416 (3)	С9—Н9	0.9500	
O3—H3A	0.8400	C10—C12	1.475 (3)	

supporting information

N1—N2	1.286 (2)	C11—H11A	0.9800
N1—C1	1.410 (2)	C11—H11B	0.9800
N2—C7	1.371 (2)	C11—H11C	0.9800
N3—C10	1.367 (2)	C12—C17	1.393 (3)
N3—C7	1.380 (3)	C12—C13	1.396 (3)
N3—C11	1.460 (3)	C13—C14	1.388 (3)
C1—C6	1.397 (3)	C14—C15	1.386 (3)
C1-C2	1 409 (3)	C14—H14	0.9500
$C^2 - C^3$	1 390 (3)	C15-C16	1 382 (3)
$C_2 C_3$	1.375 (3)	C15 H15	0.9500
C_{2} H_{2}	0.0500	C16 C17	0.9500
	0.9300		1.578 (5)
C4—C5	1.389 (3)		0.9500
C4—H4	0.9500	С17—Н17	0.9500
C5—C6	1.372 (3)	C18—H18A	0.9800
С5—Н5	0.9500	C18—H18B	0.9800
С6—Н6	0.9500	C18—H18C	0.9800
C2_01_H1	109.5	C10_C9_H9	125 7
$C_1 = C_1 = C_1 = C_1 = C_1 = C_2 $	109.5	$N_3 C_{10} C_9$	123.7 107.62 (17)
$C_{12} = 02 = 112$	100.5	$N_{3} = C_{10} = C_{12}$	107.02(17) 124.25(10)
10 - 03 - 113	109.3	N_{3} $-C_{10}$ $-C_{12}$ C_{10} $-C_{12}$	124.33(19) 128.02(19)
N2—N1—C1	115.04 (18)	$C_{2} = C_{10} = C_{12}$	128.05 (18)
NI—N2—C7	115.61 (18)	N3—CII—HIIA	109.5
C10—N3—C7	108.49 (17)	N3—C11—H11B	109.5
C10—N3—C11	127.06 (17)	H11A—C11—H11B	109.5
C7—N3—C11	124.41 (17)	N3—C11—H11C	109.5
C6—C1—C2	118.50 (19)	H11A—C11—H11C	109.5
C6-C1-N1	115.80 (19)	H11B—C11—H11C	109.5
C2-C1-N1	125.66 (19)	C17—C12—C13	118.08 (19)
O1—C2—C3	119.0 (2)	C17—C12—C10	119.67 (19)
O1—C2—C1	121.41 (18)	C13—C12—C10	122.00 (19)
C3—C2—C1	119.6 (2)	O2—C13—C14	121.74 (19)
C4-C3-C2	120.4 (2)	O2—C13—C12	117.65 (18)
C4—C3—H3	119.8	C_{14} C_{13} C_{12}	120 60 (19)
C2_C3_H3	119.8	C_{15} C_{14} C_{13}	1198(2)
$C_{2} = C_{3} = C_{4} = C_{5}$	120.6(2)	C_{15} C_{14} H_{14}	120.1
$C_3 = C_4 = C_3$	110.7	C_{13} C_{14} H_{14}	120.1
$C_5 = C_4 = H_4$	119.7	C16 C15 C14	120.1
C3-C4-H4	119.7	C16 - C15 - C14	120.3 (2)
C_{0}	119.3 (2)	C16—C15—H15	119.9
С6—С5—Н5	120.3	C14—C15—H15	119.9
C4—C5—H5	120.3	C17—C16—C15	119.5 (2)
C5—C6—C1	121.5 (2)	С17—С16—Н16	120.3
С5—С6—Н6	119.3	C15—C16—H16	120.3
С1—С6—Н6	119.3	C16—C17—C12	121.7 (2)
N2—C7—N3	117.55 (18)	С16—С17—Н17	119.2
N2—C7—C8	133.93 (19)	С12—С17—Н17	119.2
N3—C7—C8	108.49 (17)	O3—C18—H18A	109.5
С7—С8—С9	106.85 (19)	O3—C18—H18B	109.5
С7—С8—Н8	126.6	H18A—C18—H18B	109.5

С9—С8—Н8	126.6	O3—C18—H18C	109.5
C8—C9—C10	108.55 (18)	H18A—C18—H18C	109.5
С8—С9—Н9	125.7	H18B—C18—H18C	109.5
C1—N1—N2—C7	-177.49 (17)	C7—C8—C9—C10	-0.8 (2)
N2—N1—C1—C6	-179.85 (17)	C7—N3—C10—C9	-0.6 (2)
N2—N1—C1—C2	2.6 (3)	C11—N3—C10—C9	177.13 (18)
C6-C1-C2-O1	179.88 (19)	C7—N3—C10—C12	178.51 (18)
N1-C1-C2-O1	-2.6 (3)	C11—N3—C10—C12	-3.7 (3)
C6—C1—C2—C3	-0.9 (3)	C8—C9—C10—N3	0.9 (2)
N1—C1—C2—C3	176.63 (19)	C8—C9—C10—C12	-178.2 (2)
O1—C2—C3—C4	179.32 (19)	N3—C10—C12—C17	128.7 (2)
C1—C2—C3—C4	0.1 (3)	C9—C10—C12—C17	-52.3 (3)
C2—C3—C4—C5	1.1 (3)	N3—C10—C12—C13	-57.2 (3)
C3—C4—C5—C6	-1.4 (3)	C9—C10—C12—C13	121.8 (2)
C4—C5—C6—C1	0.6 (3)	C17—C12—C13—O2	176.58 (17)
C2-C1-C6-C5	0.6 (3)	C10-C12-C13-O2	2.4 (3)
N1-C1-C6-C5	-177.19 (18)	C17—C12—C13—C14	-2.3 (3)
N1—N2—C7—N3	179.37 (17)	C10-C12-C13-C14	-176.47 (18)
N1—N2—C7—C8	1.5 (3)	O2—C13—C14—C15	-177.29 (17)
C10—N3—C7—N2	-178.23 (17)	C12—C13—C14—C15	1.5 (3)
C11—N3—C7—N2	3.9 (3)	C13—C14—C15—C16	0.0 (3)
C10—N3—C7—C8	0.2 (2)	C14—C15—C16—C17	-0.8 (3)
C11—N3—C7—C8	-177.69 (18)	C15—C16—C17—C12	0.0 (3)
N2-C7-C8-C9	178.4 (2)	C13—C12—C17—C16	1.5 (3)
N3—C7—C8—C9	0.4 (2)	C10—C12—C17—C16	175.89 (18)

Hydrogen-bond geometry (Å, °)

Cg1 and Cg2 are the centroids of the C1–C6 and C12-C17 rings, respectively

D—H···A	D—H	H···A	D····A	D—H…A
O1—H1…N2	0.84	1.81	2.530 (2)	143
O2—H2···O3 ⁱ	0.84	1.81	2.641 (2)	171
O3—H3A···O1 ⁱⁱ	0.84	1.97	2.763 (2)	157
C11—H11 <i>B</i> ··· <i>Cg</i> 1 ⁱⁱⁱ	0.98	2.73	3.587 (2)	147
C18—H18 C ··· $Cg2^{iv}$	0.98	2.75	3.483 (3)	132

Symmetry codes: (i) -*x*+1, -*y*+2, -*z*+1; (ii) *x*, *y*, *z*+1; (iii) -*x*, -*y*+1, -*z*; (iv) -*x*, -*y*+2, -*z*+1.