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N'-(2-Chlorobenzylidene)-3,4,5-trimethoxybenzohydrazide methanol solvate

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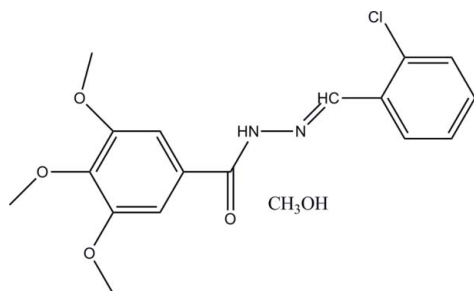
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Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.032; wR factor = 0.081; data-to-parameter ratio = 16.3.

In the title compound, $\text{C}_{17}\text{H}_{17}\text{ClN}_2\text{O}_4 \cdot \text{CH}_4\text{O}$, the dihedral angle between the benzene ring planes is $5.29(6)^\circ$. Inter-molecular $\text{N}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds link the molecules into a chain along the a axis.

Related literature

For related literature, see: Allen *et al.* (1987), Bernardino *et al.* (2006); Ganjali *et al.* (2006); Gardner *et al.* (1991); Patole *et al.* (2003)



Experimental

Crystal data

$\text{C}_{17}\text{H}_{17}\text{ClN}_2\text{O}_4 \cdot \text{CH}_4\text{O}$
 $M_r = 380.82$
Orthorhombic, $Pna2_1$

$a = 12.9356(7)$ Å
 $b = 4.8718(3)$ Å
 $c = 29.4119(16)$ Å

$V = 1853.53(18)$ Å³
 $Z = 4$
Mo $K\alpha$ radiation

$\mu = 0.24$ mm⁻¹
 $T = 173(2)$ K
 $0.48 \times 0.40 \times 0.39$ mm

Data collection

Bruker SMART 1000 CCD diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)
 $T_{\min} = 0.895$, $T_{\max} = 0.913$

9008 measured reflections
3916 independent reflections
3561 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.021$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.031$
 $wR(F^2) = 0.080$
 $S = 1.09$
3916 reflections
240 parameters
1 restraint

H-atom parameters constrained
 $\Delta\rho_{\max} = 0.20$ e Å⁻³
 $\Delta\rho_{\min} = -0.17$ e Å⁻³
Absolute structure: Flack (1983),
1846 Friedel pairs
Flack parameter: 0.04 (5)

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{N1}-\text{H1} \cdots \text{O5}^i$	0.88	2.01	2.8673 (19)	165
$\text{O5}-\text{H5} \cdots \text{O4}$	0.84	1.97	2.7904 (18)	166

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

Data collection: SMART (Bruker, 2001); cell refinement: SAINT-Plus (Bruker, 2003); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2761).

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supporting information

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N'*-(2-Chlorobenzylidene)-3,4,5-trimethoxybenzohydrazide methanol solvate*Dao-Hang He, Yong-Chuang Zhu, Zhuo-Ru Yang, Shao-Yun Song and Qi-Jin Chen****S1. Comment**

Molecules involving Schiff bases have attracted much attention due to their diverse range of bioactivities in pharmaceutical and agrochemical field (e.g. Bernardino *et al.*, 2006; Ganjali *et al.*, 2006). We now report the synthesis and structure of the title compound, (I), obtained by the condensation of 3,4,5-trimethoxybenzohydrazide with 2-chlorobenzaldehyde as a methanol solvate (Fig. 1).

The bond lengths and bond angles for (I) are within normal ranges (Allen *et al.*, 1987). The two benzene rings are approximately planar, with a dihedral angle of 5.29 (6)°. The methanol molecules in the crystal are lined to the Schiff base moieties through intermolecular N—H···O and O—H···O hydrogen bonds to form a chain along the *a* axis, which helps to consolidate the packing (Fig 2).

S2. Experimental

A mixture of 3,4,5-trimethoxybenzohydrazide (1 mmol) and 2-chlorobenzaldehyde in anhydrous ethanol (10 ml) was refluxed for 2 h. When the solution was cooled to room temperature, some white needles separated out. After filtration, colorless blocks of (I) were obtained by slow evaporation of a methanol solution.

S3. Refinement

All H atoms were placed in geometrically idealized positions and allowed to ride on their parent atoms, with N—H = 0.88 Å, O—H = 0.84 Å, C—H = 0.95 (aromatic and N=CH), 0.98 (methyl) Å) and $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C, N, O})$, where $x = 1.5$ for the methyl and hydroxyl groups, $x = 1.2$ for all other H atoms.

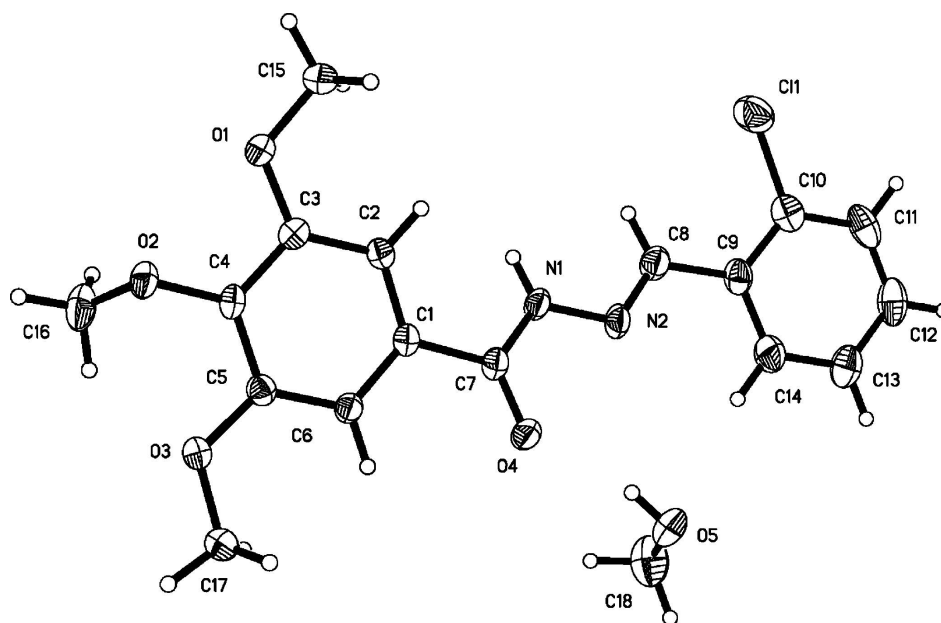
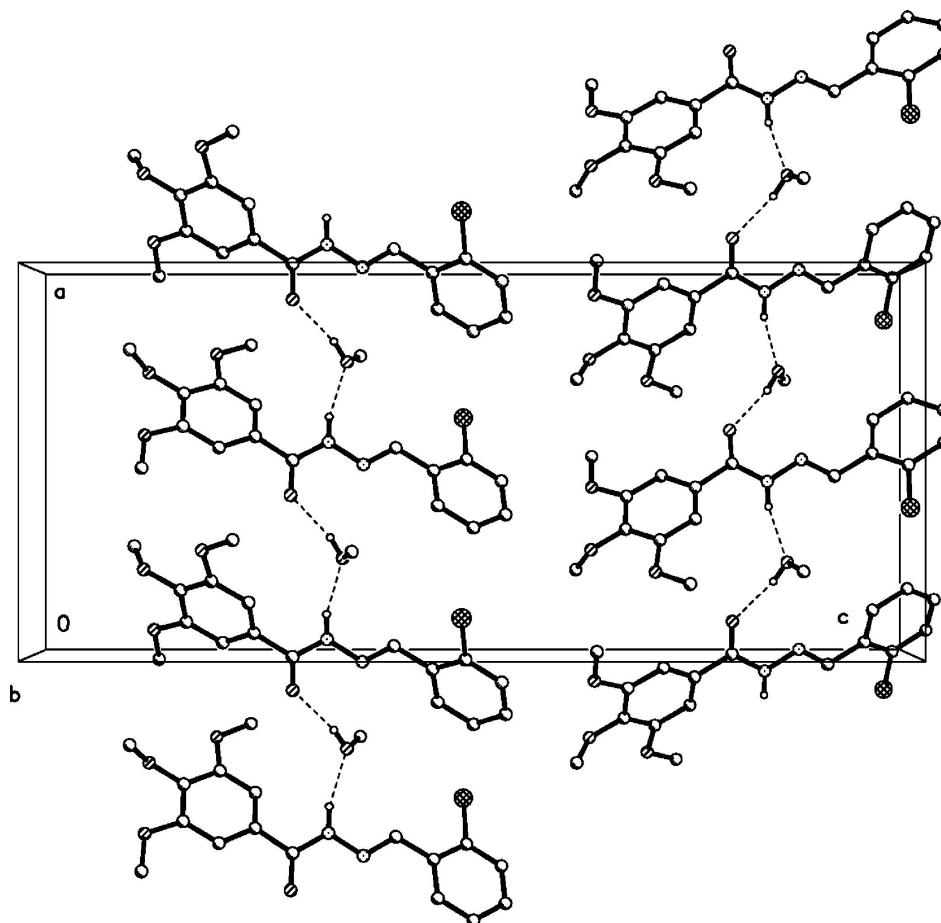


Figure 1

The molecular structure of (I) with displacement ellipsoids for the non-hydrogen atoms drawn at the 50% probability level.

**Figure 2**

The packing of (I), viewed down the *b* axis. The dashed lines represent the hydrogen bonding interactions.

(I)*Crystal data*

$C_{17}H_{17}ClN_2O_4 \cdot CH_4O$

$M_r = 380.82$

Orthorhombic, *Pna*2₁

Hall symbol: P 2c -2n

$a = 12.9356 (7) \text{ \AA}$

$b = 4.8718 (3) \text{ \AA}$

$c = 29.4119 (16) \text{ \AA}$

$V = 1853.53 (18) \text{ \AA}^3$

$Z = 4$

$F(000) = 800$

$D_x = 1.365 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 5318 reflections

$\theta = 2.6\text{--}27.0^\circ$

$\mu = 0.24 \text{ mm}^{-1}$

$T = 173 \text{ K}$

Block, colorless

$0.48 \times 0.40 \times 0.39 \text{ mm}$

Data collection

Bruker SMART 1000 CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 2003)

$T_{\min} = 0.895$, $T_{\max} = 0.913$

9008 measured reflections

3916 independent reflections

3561 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.021$
 $\theta_{\text{max}} = 27.0^\circ$, $\theta_{\text{min}} = 2.8^\circ$
 $h = -16 \rightarrow 8$

$k = -5 \rightarrow 6$
 $l = -37 \rightarrow 36$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.031$
 $wR(F^2) = 0.080$
 $S = 1.09$
 3916 reflections
 240 parameters
 1 restraint
 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
 $w = 1/[\sigma^2(F_o^2) + (0.039P)^2 + 0.3064P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.012$
 $\Delta\rho_{\text{max}} = 0.20 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.17 \text{ e } \text{\AA}^{-3}$
 Absolute structure: Flack (1983), 1846 Friedel
 pairs
 Absolute structure parameter: 0.04 (5)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s 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 $F^2 > \sigma(F^2)$ 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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.88399 (4)	-0.29291 (12)	0.98890 (2)	0.04522 (15)
C1	0.93423 (13)	0.2105 (4)	0.75620 (6)	0.0215 (3)
C2	0.85309 (13)	0.0229 (4)	0.75340 (6)	0.0211 (3)
H2	0.8352	-0.0866	0.7789	0.025*
C3	0.79858 (12)	-0.0025 (3)	0.71285 (6)	0.0213 (3)
C4	0.82468 (13)	0.1595 (3)	0.67522 (6)	0.0209 (3)
C5	0.90935 (13)	0.3386 (4)	0.67796 (6)	0.0221 (3)
C6	0.96347 (13)	0.3656 (4)	0.71860 (6)	0.0216 (3)
H6	1.0201	0.4892	0.7207	0.026*
C7	0.99585 (13)	0.2484 (4)	0.79886 (6)	0.0231 (4)
C8	0.96354 (14)	0.0864 (4)	0.91319 (7)	0.0308 (4)
H8	0.8949	0.0174	0.9121	0.037*
C9	1.02227 (15)	0.0827 (4)	0.95597 (6)	0.0286 (4)
C10	0.99287 (15)	-0.0824 (4)	0.99280 (7)	0.0319 (4)
C11	1.04956 (19)	-0.0869 (5)	1.03275 (7)	0.0403 (5)
H11	1.0289	-0.2029	1.0571	0.048*
C12	1.13525 (18)	0.0753 (5)	1.03723 (7)	0.0435 (5)
H12	1.1731	0.0753	1.0649	0.052*
C13	1.16701 (18)	0.2405 (5)	1.00118 (7)	0.0416 (5)
H13	1.2270	0.3517	1.0041	0.050*

C14	1.11093 (16)	0.2419 (5)	0.96120 (7)	0.0362 (5)
H14	1.1334	0.3541	0.9367	0.043*
C15	0.69115 (15)	-0.3585 (4)	0.74240 (6)	0.0260 (4)
H15A	0.7517	-0.4661	0.7515	0.039*
H15B	0.6361	-0.4827	0.7324	0.039*
H15C	0.6668	-0.2497	0.7683	0.039*
C16	0.71561 (18)	0.3559 (4)	0.61922 (7)	0.0386 (5)
H16A	0.6581	0.3954	0.6399	0.058*
H16B	0.6884	0.3175	0.5888	0.058*
H16C	0.7619	0.5150	0.6179	0.058*
C17	1.01472 (14)	0.6703 (4)	0.63995 (7)	0.0302 (4)
H17A	0.9989	0.8111	0.6627	0.045*
H17B	1.0217	0.7565	0.6100	0.045*
H17C	1.0796	0.5785	0.6480	0.045*
C18	1.2215 (2)	0.7735 (5)	0.86998 (9)	0.0501 (6)
H18A	1.1903	0.7780	0.9003	0.075*
H18B	1.1739	0.8569	0.8480	0.075*
H18C	1.2866	0.8763	0.8703	0.075*
N1	0.94896 (12)	0.1728 (3)	0.83814 (5)	0.0268 (3)
H1	0.8841	0.1185	0.8385	0.032*
N2	1.00645 (11)	0.1840 (3)	0.87756 (5)	0.0269 (3)
O1	0.71872 (9)	-0.1791 (3)	0.70587 (4)	0.0259 (3)
O2	0.77131 (9)	0.1235 (3)	0.63530 (4)	0.0259 (3)
O3	0.93290 (10)	0.4729 (3)	0.63872 (4)	0.0308 (3)
O4	1.08391 (10)	0.3399 (3)	0.79778 (4)	0.0298 (3)
O5	1.24118 (9)	0.4964 (3)	0.85732 (5)	0.0328 (3)
H5	1.1900	0.4338	0.8431	0.049*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0469 (3)	0.0485 (3)	0.0403 (3)	-0.0043 (2)	0.0087 (2)	0.0094 (3)
C1	0.0212 (8)	0.0249 (9)	0.0183 (8)	0.0027 (7)	-0.0023 (7)	-0.0007 (7)
C2	0.0216 (8)	0.0220 (8)	0.0196 (8)	0.0027 (6)	0.0005 (7)	0.0018 (7)
C3	0.0197 (8)	0.0211 (8)	0.0231 (8)	0.0018 (6)	0.0002 (7)	-0.0038 (7)
C4	0.0212 (8)	0.0242 (9)	0.0171 (8)	0.0044 (7)	-0.0033 (6)	-0.0027 (6)
C5	0.0219 (8)	0.0239 (9)	0.0205 (8)	0.0015 (7)	0.0007 (6)	0.0009 (7)
C6	0.0205 (8)	0.0243 (8)	0.0201 (8)	-0.0005 (7)	-0.0017 (6)	-0.0003 (7)
C7	0.0219 (8)	0.0280 (9)	0.0195 (8)	0.0017 (7)	-0.0028 (7)	-0.0007 (7)
C8	0.0255 (9)	0.0425 (11)	0.0244 (9)	-0.0010 (8)	-0.0022 (7)	0.0010 (8)
C9	0.0308 (10)	0.0370 (10)	0.0179 (8)	0.0040 (8)	-0.0003 (7)	-0.0002 (8)
C10	0.0353 (10)	0.0339 (10)	0.0266 (9)	0.0074 (8)	0.0055 (8)	-0.0001 (8)
C11	0.0580 (14)	0.0406 (12)	0.0221 (9)	0.0135 (11)	0.0047 (9)	0.0069 (9)
C12	0.0540 (14)	0.0538 (14)	0.0227 (9)	0.0139 (11)	-0.0115 (9)	-0.0045 (10)
C13	0.0419 (12)	0.0519 (14)	0.0309 (11)	0.0028 (10)	-0.0109 (9)	-0.0042 (10)
C14	0.0397 (11)	0.0458 (12)	0.0233 (9)	-0.0009 (10)	-0.0031 (8)	0.0043 (9)
C15	0.0269 (9)	0.0232 (9)	0.0278 (9)	-0.0019 (7)	0.0025 (7)	-0.0004 (7)
C16	0.0433 (12)	0.0374 (11)	0.0351 (11)	0.0081 (10)	-0.0179 (9)	-0.0010 (9)

C17	0.0269 (9)	0.0345 (10)	0.0293 (9)	-0.0029 (8)	-0.0006 (8)	0.0078 (8)
C18	0.0638 (15)	0.0453 (13)	0.0412 (13)	0.0120 (12)	-0.0099 (12)	-0.0078 (11)
N1	0.0206 (7)	0.0415 (9)	0.0184 (7)	-0.0044 (6)	-0.0041 (6)	0.0019 (6)
N2	0.0257 (7)	0.0370 (9)	0.0180 (7)	0.0010 (6)	-0.0053 (6)	0.0009 (6)
O1	0.0263 (6)	0.0282 (7)	0.0232 (6)	-0.0049 (5)	-0.0047 (5)	0.0008 (5)
O2	0.0292 (6)	0.0276 (6)	0.0208 (6)	0.0011 (5)	-0.0064 (5)	-0.0028 (5)
O3	0.0309 (6)	0.0399 (7)	0.0217 (6)	-0.0096 (6)	-0.0052 (6)	0.0076 (6)
O4	0.0243 (6)	0.0426 (8)	0.0225 (7)	-0.0089 (6)	-0.0040 (5)	0.0029 (6)
O5	0.0231 (6)	0.0405 (7)	0.0347 (7)	0.0025 (6)	-0.0042 (5)	-0.0078 (6)

Geometric parameters (Å, °)

C11—C10	1.746 (2)	C12—H12	0.9500
C1—C6	1.392 (2)	C13—C14	1.382 (3)
C1—C2	1.394 (2)	C13—H13	0.9500
C1—C7	1.498 (2)	C14—H14	0.9500
C2—C3	1.391 (2)	C15—O1	1.430 (2)
C2—H2	0.9500	C15—H15A	0.9800
C3—O1	1.360 (2)	C15—H15B	0.9800
C3—C4	1.401 (2)	C15—H15C	0.9800
C4—O2	1.373 (2)	C16—O2	1.423 (2)
C4—C5	1.402 (2)	C16—H16A	0.9800
C5—O3	1.361 (2)	C16—H16B	0.9800
C5—C6	1.391 (2)	C16—H16C	0.9800
C6—H6	0.9500	C17—O3	1.430 (2)
C7—O4	1.224 (2)	C17—H17A	0.9800
C7—N1	1.356 (2)	C17—H17B	0.9800
C8—N2	1.278 (2)	C17—H17C	0.9800
C8—C9	1.470 (2)	C18—O5	1.423 (3)
C8—H8	0.9500	C18—H18A	0.9800
C9—C14	1.393 (3)	C18—H18B	0.9800
C9—C10	1.402 (3)	C18—H18C	0.9800
C10—C11	1.385 (3)	N1—N2	1.378 (2)
C11—C12	1.368 (3)	N1—H1	0.8800
C11—H11	0.9500	O5—H5	0.8400
C12—C13	1.393 (3)		
C6—C1—C2	120.90 (15)	C14—C13—H13	120.1
C6—C1—C7	117.00 (15)	C12—C13—H13	120.1
C2—C1—C7	122.07 (14)	C13—C14—C9	121.6 (2)
C3—C2—C1	119.38 (15)	C13—C14—H14	119.2
C3—C2—H2	120.3	C9—C14—H14	119.2
C1—C2—H2	120.3	O1—C15—H15A	109.5
O1—C3—C2	124.80 (15)	O1—C15—H15B	109.5
O1—C3—C4	114.85 (15)	H15A—C15—H15B	109.5
C2—C3—C4	120.34 (15)	O1—C15—H15C	109.5
O2—C4—C3	118.83 (15)	H15A—C15—H15C	109.5
O2—C4—C5	121.43 (15)	H15B—C15—H15C	109.5

C3—C4—C5	119.55 (14)	O2—C16—H16A	109.5
O3—C5—C6	124.76 (15)	O2—C16—H16B	109.5
O3—C5—C4	115.15 (14)	H16A—C16—H16B	109.5
C6—C5—C4	120.09 (15)	O2—C16—H16C	109.5
C5—C6—C1	119.63 (16)	H16A—C16—H16C	109.5
C5—C6—H6	120.2	H16B—C16—H16C	109.5
C1—C6—H6	120.2	O3—C17—H17A	109.5
O4—C7—N1	122.52 (16)	O3—C17—H17B	109.5
O4—C7—C1	121.23 (15)	H17A—C17—H17B	109.5
N1—C7—C1	116.24 (15)	O3—C17—H17C	109.5
N2—C8—C9	118.82 (17)	H17A—C17—H17C	109.5
N2—C8—H8	120.6	H17B—C17—H17C	109.5
C9—C8—H8	120.6	O5—C18—H18A	109.5
C14—C9—C10	117.21 (17)	O5—C18—H18B	109.5
C14—C9—C8	120.89 (18)	H18A—C18—H18B	109.5
C10—C9—C8	121.89 (18)	O5—C18—H18C	109.5
C11—C10—C9	121.40 (19)	H18A—C18—H18C	109.5
C11—C10—C11	118.27 (16)	H18B—C18—H18C	109.5
C9—C10—C11	120.33 (15)	C7—N1—N2	117.68 (14)
C12—C11—C10	120.1 (2)	C7—N1—H1	121.2
C12—C11—H11	119.9	N2—N1—H1	121.2
C10—C11—H11	119.9	C8—N2—N1	116.16 (15)
C11—C12—C13	119.97 (19)	C3—O1—C15	117.56 (13)
C11—C12—H12	120.0	C4—O2—C16	115.90 (14)
C13—C12—H12	120.0	C5—O3—C17	117.86 (14)
C14—C13—C12	119.7 (2)	C18—O5—H5	109.5
C6—C1—C2—C3	-2.2 (2)	C14—C9—C10—C11	0.1 (3)
C7—C1—C2—C3	179.86 (15)	C8—C9—C10—C11	179.16 (19)
C1—C2—C3—O1	179.51 (15)	C14—C9—C10—C11	-178.94 (15)
C1—C2—C3—C4	-0.2 (2)	C8—C9—C10—C11	0.2 (3)
O1—C3—C4—O2	-1.7 (2)	C9—C10—C11—C12	1.1 (3)
C2—C3—C4—O2	178.07 (15)	C11—C10—C11—C12	-179.86 (17)
O1—C3—C4—C5	-176.74 (15)	C10—C11—C12—C13	-1.5 (3)
C2—C3—C4—C5	3.0 (2)	C11—C12—C13—C14	0.7 (3)
O2—C4—C5—O3	0.7 (2)	C12—C13—C14—C9	0.5 (3)
C3—C4—C5—O3	175.62 (15)	C10—C9—C14—C13	-0.8 (3)
O2—C4—C5—C6	-178.40 (16)	C8—C9—C14—C13	-180.0 (2)
C3—C4—C5—C6	-3.5 (2)	O4—C7—N1—N2	-4.4 (3)
O3—C5—C6—C1	-177.87 (16)	C1—C7—N1—N2	174.82 (16)
C4—C5—C6—C1	1.1 (3)	C9—C8—N2—N1	177.74 (17)
C2—C1—C6—C5	1.7 (3)	C7—N1—N2—C8	-173.39 (17)
C7—C1—C6—C5	179.79 (15)	C2—C3—O1—C15	-2.3 (2)
C6—C1—C7—O4	-21.9 (3)	C4—C3—O1—C15	177.43 (14)
C2—C1—C7—O4	156.15 (17)	C3—C4—O2—C16	118.71 (19)
C6—C1—C7—N1	158.87 (16)	C5—C4—O2—C16	-66.3 (2)
C2—C1—C7—N1	-23.1 (2)	C6—C5—O3—C17	-4.5 (3)
N2—C8—C9—C14	16.3 (3)	C4—C5—O3—C17	176.48 (15)

N2—C8—C9—C10 -162.78 (19)

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N1—H1...O5 ⁱ	0.88	2.01	2.8673 (19)	165
O5—H5...O4	0.84	1.97	2.7904 (18)	166

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