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(E)-N'-(5-Bromo-2-hydroxybenzylidene)-3,4,5-trimethoxybenzohydrazide

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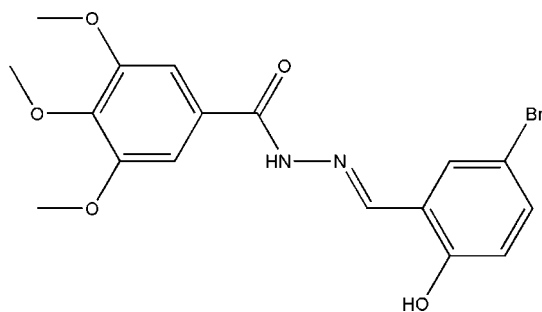
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Key indicators: single-crystal X-ray study; $T = 273$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.037; wR factor = 0.097; data-to-parameter ratio = 13.3.

The molecule of the title compound, $\text{C}_{17}\text{H}_{17}\text{BrN}_2\text{O}_5$, assumes an *E* configuration, with the 5-bromo-2-hydroxyphenyl and benzohydrazide units located on opposite sites of the $\text{C}=\text{N}$ double bond. The dihedral angle between the planes of the two benzene rings is $32.48(15)^\circ$. The crystal structure is stabilized by intramolecular $\text{O}-\text{H}\cdots\text{N}$ and intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For related literature, see: Yang *et al.* (1996); Nawar & Hosny (2000); Pelagatti *et al.* (1999); Ainscough *et al.* (1998). For related structures, see: Diao & Yu (2006); Jing *et al.* (2005); Wang *et al.* (2008).



Experimental

Crystal data

$\text{C}_{17}\text{H}_{17}\text{BrN}_2\text{O}_5$
 $M_r = 409.24$

Monoclinic, $P2_1/c$
 $a = 11.4157(19)$ Å

$b = 16.279(3)$ Å
 $c = 9.3738(16)$ Å
 $\beta = 100.210(3)^\circ$
 $V = 1714.4(5)$ Å³
 $Z = 4$

Mo $K\alpha$ radiation
 $\mu = 2.43$ mm⁻¹
 $T = 273(2)$ K [Unusual. Please check]
 $0.15 \times 0.10 \times 0.06$ mm

Data collection

Bruker APEX CCD area-detector diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.712$, $T_{\max} = 0.868$

8954 measured reflections
3037 independent reflections
2065 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.033$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.097$
 $S = 1.01$
3037 reflections

228 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.43$ e Å⁻³
 $\Delta\rho_{\min} = -0.54$ e Å⁻³

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O1}-\text{H1}\cdots\text{N2}$	0.82	1.92	2.642 (3)	145
$\text{N1}-\text{H1A}\cdots\text{O2}^i$	0.86	2.14	2.888 (3)	146

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

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; 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.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2120).

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supporting information

Acta Cryst. (2008). E64, o2408 [doi:10.1107/S1600536808038464]

(E)-N'-(5-Bromo-2-hydroxybenzylidene)-3,4,5-trimethoxybenzohydrazide**Zhen-dong Zhao, Yu-min Wang, Yu-xiang Chen and Liang-wu Bi****S1. Comment**

Salicylaldehyde hydrazones have received considerable attention because of their possible applications in catalysis and pharmacology (Yang *et al.*, 1996; Nawar & Hosny, 2000; Pelagatti *et al.*, 1999; Ainscough *et al.*, 1998). As part of our ongoing investigation on acylhydrazone compounds (Wang *et al.*, 2008) and their metal complexes, we have obtained the title compound, (I), from a condensation of 3,4,5-trimethoxybenzohydrazide and 5-bromo-2-hydroxybenzaldehyde, and report its crystal structure in this paper.

The molecular structure of (I) (Fig. 1) contains 3,4,5-trimethoxybenzohydrazide and 5-bromo-2-hydroxybenzaldehyde moieties that are located on opposite sides of the C=N double bond exhibiting an E configuration about C=N. The dihedral angle between the mean-planes of the two benzene rings is 32.48 (15)°. The crystal structure involves an intramolecular hydrogen bond O1–H1···N2, and is further stabilized by an intermolecular hydrogen bonds N1–H1A···O2, resulting in chains of molecules extended along the *c*-axis; details are given in Table 1 and Fig. 2. The bond distances and bond angles in (I) are in agreement with the corresponding dimensions reported for some structures closely related to (I) (Diao & Yu, 2006; Jing *et al.*, 2005; Wang *et al.*, 2008). (

S2. Experimental

An ethanol solution (50 ml) of 3,4,5-trimethoxybenzohydrazide (0.01 mol) and 5-bromo-2-hydroxybenzaldehyde (0.01 mol) was refluxed and stirred for 2h ; the mixture was cooled and the resulting solid product, (I), was collected by filtration. Crystals suitable for single-crystal X-ray diffraction were grown by slow evaporation of a solution in THF.

S3. Refinement

All H atoms were placed geometrically at the distances: C—H(methyl) = 0.96, C—H(aryl) = 0.93, N—H = 0.86 and O—H = 0.82 Å and included in the refinement in riding motion approximation with $U_{\text{iso}}(\text{H}) = 1.2$ or $1.5U_{\text{eq}}$ of the carrier atom.

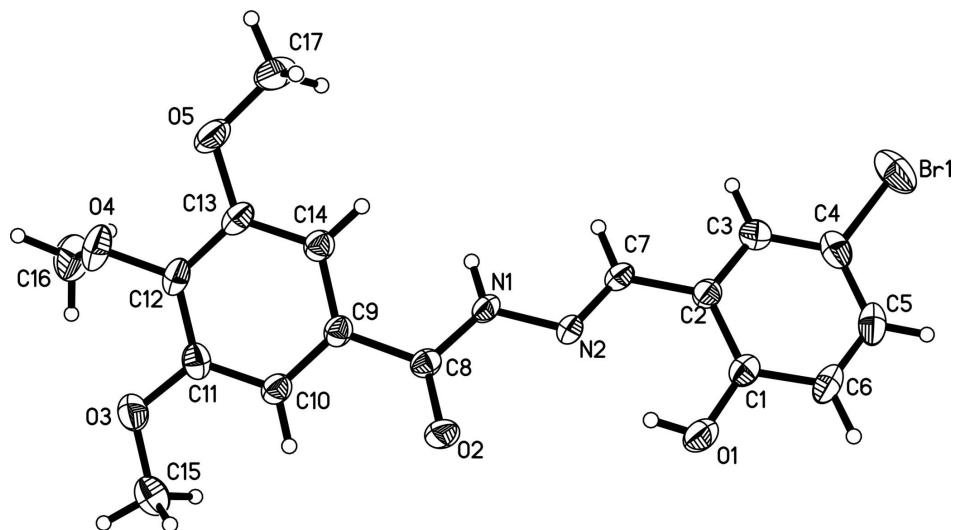


Figure 1

A view of the molecular structure of (I), showing displacement ellipsoids at the 50% probability level.

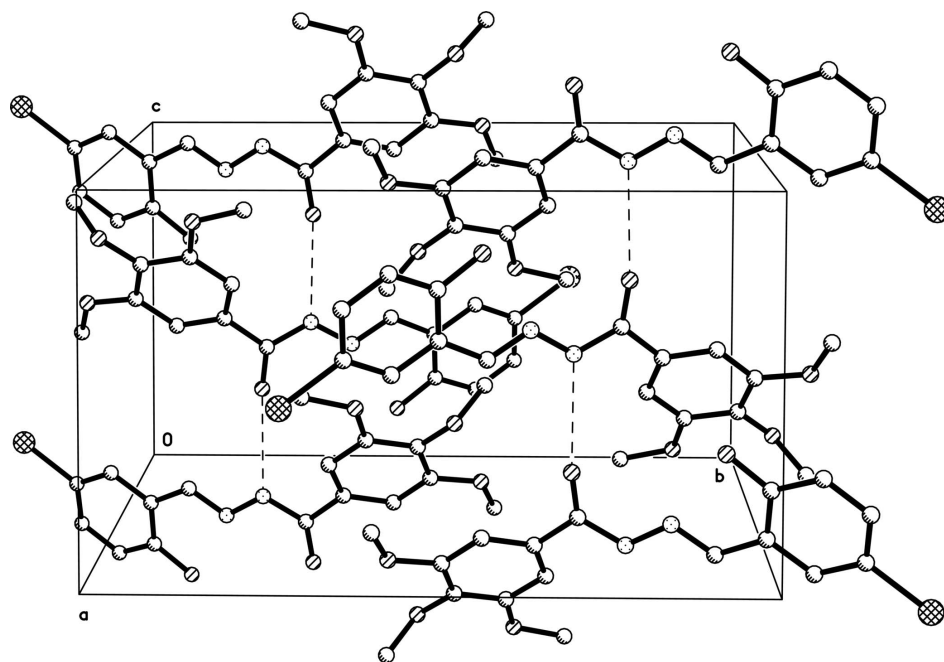


Figure 2

The packing of (I), showing the intermolecular hydrogen-bonded extended network.

(E)-N'-(5-Bromo-2-hydroxybenzylidene)-3,4,5-trimethoxybenzohydrazide

Crystal data

$C_{17}H_{17}BrN_2O_5$

$M_r = 409.24$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2_1/c$

$a = 11.4157(19)\ \text{\AA}$

$b = 16.279(3)\ \text{\AA}$

$c = 9.3738(16)\ \text{\AA}$

$\beta = 100.210(3)^\circ$

$V = 1714.4(5)\ \text{\AA}^3$

$Z = 4$

$F(000) = 832$

$D_x = 1.586\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 2452 reflections
 $\theta = 2.5\text{--}22.8^\circ$
 $\mu = 2.43 \text{ mm}^{-1}$

$T = 273 \text{ K}$
 Block, colourless
 $0.15 \times 0.10 \times 0.06 \text{ mm}$

Data collection

Bruker APEX CCD area-detector
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.712$, $T_{\max} = 0.868$

8954 measured reflections
 3037 independent reflections
 2065 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.033$
 $\theta_{\text{max}} = 25.1^\circ$, $\theta_{\text{min}} = 1.8^\circ$
 $h = -13 \rightarrow 12$
 $k = -19 \rightarrow 19$
 $l = -11 \rightarrow 8$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.097$
 $S = 1.01$
 3037 reflections
 228 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-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0439P)^2 + 0.6769P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.002$
 $\Delta\rho_{\text{max}} = 0.43 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.54 \text{ e \AA}^{-3}$
 Extinction correction: SHELXL97 (Sheldrick,
 2008), $F_c^* = kFc[1 + 0.001x\text{Fc}^2\lambda^3/\sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.0171 (11)

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.94048 (4)	1.21831 (2)	0.94475 (6)	0.0865 (2)
O1	0.9086 (2)	0.92631 (14)	1.3259 (2)	0.0609 (6)
H1	0.8677	0.8897	1.2824	0.091*
O2	0.74380 (19)	0.70776 (12)	1.2188 (2)	0.0493 (5)
O3	0.6562 (2)	0.42963 (12)	0.9513 (2)	0.0605 (6)
O4	0.4797 (2)	0.47058 (14)	0.7287 (2)	0.0661 (7)

O5	0.42036 (19)	0.62400 (14)	0.6698 (2)	0.0595 (6)
N1	0.7240 (2)	0.78966 (13)	1.0217 (2)	0.0413 (6)
H1A	0.6980	0.7944	0.9302	0.050*
N2	0.7787 (2)	0.85483 (14)	1.0984 (2)	0.0392 (6)
C1	0.9120 (3)	0.99080 (18)	1.2363 (3)	0.0446 (7)
C2	0.8549 (2)	0.98993 (16)	1.0913 (3)	0.0400 (7)
C3	0.8637 (3)	1.05886 (18)	1.0069 (3)	0.0485 (8)
H3	0.8252	1.0596	0.9107	0.058*
C4	0.9282 (3)	1.12578 (18)	1.0634 (4)	0.0542 (8)
C5	0.9863 (3)	1.1252 (2)	1.2051 (4)	0.0592 (9)
H5	1.0316	1.1702	1.2427	0.071*
C6	0.9772 (3)	1.0587 (2)	1.2899 (4)	0.0573 (9)
H6	1.0158	1.0591	1.3861	0.069*
C7	0.7914 (2)	0.91924 (17)	1.0253 (3)	0.0417 (7)
H7	0.7588	0.9206	0.9271	0.050*
C8	0.7105 (2)	0.71797 (17)	1.0892 (3)	0.0369 (6)
C9	0.6508 (2)	0.65335 (17)	0.9912 (3)	0.0381 (7)
C10	0.6826 (3)	0.57283 (17)	1.0217 (3)	0.0416 (7)
H10	0.7397	0.5600	1.1023	0.050*
C11	0.6292 (3)	0.51139 (17)	0.9317 (3)	0.0450 (7)
C12	0.5412 (3)	0.52992 (18)	0.8141 (3)	0.0460 (7)
C13	0.5089 (3)	0.61112 (19)	0.7852 (3)	0.0445 (7)
C14	0.5643 (2)	0.67246 (18)	0.8733 (3)	0.0413 (7)
H14	0.5434	0.7270	0.8534	0.050*
C15	0.7519 (3)	0.4079 (2)	1.0611 (4)	0.0713 (10)
H15A	0.8233	0.4340	1.0431	0.107*
H15B	0.7622	0.3493	1.0618	0.107*
H15C	0.7355	0.4255	1.1533	0.107*
C16	0.5452 (4)	0.4238 (2)	0.6446 (4)	0.0853 (12)
H16A	0.5791	0.4595	0.5812	0.128*
H16B	0.4935	0.3848	0.5880	0.128*
H16C	0.6077	0.3952	0.7071	0.128*
C17	0.3907 (3)	0.7059 (2)	0.6315 (4)	0.0697 (10)
H17A	0.3538	0.7308	0.7052	0.105*
H17B	0.3364	0.7070	0.5407	0.105*
H17C	0.4616	0.7357	0.6225	0.105*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0686 (3)	0.0506 (3)	0.1438 (5)	0.00267 (18)	0.0284 (3)	0.0265 (2)
O1	0.0746 (17)	0.0646 (15)	0.0377 (12)	-0.0140 (12)	-0.0059 (11)	-0.0017 (11)
O2	0.0660 (14)	0.0498 (12)	0.0287 (11)	0.0048 (10)	-0.0008 (10)	-0.0018 (9)
O3	0.0799 (16)	0.0411 (12)	0.0584 (14)	-0.0084 (11)	0.0066 (13)	-0.0024 (11)
O4	0.0701 (16)	0.0687 (15)	0.0606 (15)	-0.0312 (12)	0.0147 (13)	-0.0247 (12)
O5	0.0551 (14)	0.0737 (16)	0.0426 (13)	-0.0110 (12)	-0.0106 (11)	-0.0072 (11)
N1	0.0504 (14)	0.0429 (14)	0.0266 (12)	-0.0088 (11)	-0.0039 (11)	-0.0032 (11)
N2	0.0450 (14)	0.0359 (13)	0.0336 (13)	-0.0029 (11)	-0.0019 (11)	-0.0071 (11)

C1	0.0470 (17)	0.0485 (18)	0.0377 (17)	-0.0016 (14)	0.0053 (15)	-0.0063 (14)
C2	0.0383 (16)	0.0407 (16)	0.0394 (17)	0.0028 (13)	0.0032 (14)	-0.0047 (13)
C3	0.0459 (18)	0.0463 (18)	0.0512 (19)	0.0034 (14)	0.0031 (15)	0.0013 (15)
C4	0.0466 (19)	0.0399 (17)	0.079 (3)	0.0017 (14)	0.0184 (19)	0.0001 (17)
C5	0.051 (2)	0.0457 (19)	0.082 (3)	-0.0077 (15)	0.0138 (19)	-0.0220 (19)
C6	0.052 (2)	0.063 (2)	0.054 (2)	-0.0096 (16)	0.0002 (16)	-0.0239 (18)
C7	0.0454 (17)	0.0456 (17)	0.0304 (15)	-0.0010 (14)	-0.0032 (13)	-0.0056 (14)
C8	0.0372 (15)	0.0422 (16)	0.0308 (16)	0.0025 (13)	0.0051 (13)	-0.0042 (13)
C9	0.0416 (16)	0.0406 (16)	0.0325 (15)	-0.0050 (13)	0.0077 (13)	-0.0048 (13)
C10	0.0436 (17)	0.0463 (17)	0.0339 (16)	-0.0040 (14)	0.0043 (13)	-0.0037 (13)
C11	0.0531 (19)	0.0387 (16)	0.0471 (18)	-0.0075 (14)	0.0194 (16)	-0.0045 (14)
C12	0.0483 (18)	0.0507 (18)	0.0400 (18)	-0.0182 (15)	0.0107 (15)	-0.0130 (15)
C13	0.0396 (17)	0.060 (2)	0.0333 (17)	-0.0103 (14)	0.0033 (14)	-0.0069 (14)
C14	0.0429 (17)	0.0436 (17)	0.0371 (16)	-0.0028 (13)	0.0064 (14)	-0.0015 (13)
C15	0.084 (3)	0.0450 (19)	0.083 (3)	0.0038 (18)	0.010 (2)	0.0028 (18)
C16	0.113 (3)	0.066 (2)	0.075 (3)	-0.008 (2)	0.011 (3)	-0.035 (2)
C17	0.056 (2)	0.084 (3)	0.061 (2)	-0.0003 (19)	-0.0123 (18)	0.004 (2)

Geometric parameters (Å, °)

Br1—C4	1.892 (3)	C5—H5	0.9300
O1—C1	1.350 (4)	C6—H6	0.9300
O1—H1	0.8200	C7—H7	0.9300
O2—C8	1.219 (3)	C8—C9	1.481 (4)
O3—C11	1.371 (3)	C9—C10	1.377 (4)
O3—C15	1.407 (4)	C9—C14	1.381 (4)
O4—C12	1.366 (3)	C10—C11	1.379 (4)
O4—C16	1.403 (4)	C10—H10	0.9300
O5—C13	1.359 (3)	C11—C12	1.386 (4)
O5—C17	1.406 (4)	C12—C13	1.386 (4)
N1—C8	1.349 (3)	C13—C14	1.376 (4)
N1—N2	1.369 (3)	C14—H14	0.9300
N1—H1A	0.8600	C15—H15A	0.9600
N2—C7	1.274 (3)	C15—H15B	0.9600
C1—C6	1.378 (4)	C15—H15C	0.9600
C1—C2	1.399 (4)	C16—H16A	0.9600
C2—C3	1.387 (4)	C16—H16B	0.9600
C2—C7	1.440 (4)	C16—H16C	0.9600
C3—C4	1.368 (4)	C17—H17A	0.9600
C3—H3	0.9300	C17—H17B	0.9600
C4—C5	1.375 (5)	C17—H17C	0.9600
C5—C6	1.357 (5)		
C1—O1—H1	109.5	C14—C9—C8	121.4 (3)
C11—O3—C15	118.0 (2)	C9—C10—C11	119.4 (3)
C12—O4—C16	116.3 (3)	C9—C10—H10	120.3
C13—O5—C17	117.4 (2)	C11—C10—H10	120.3
C8—N1—N2	120.2 (2)	O3—C11—C10	123.9 (3)

C8—N1—H1A	119.9	O3—C11—C12	115.6 (3)
N2—N1—H1A	119.9	C10—C11—C12	120.5 (3)
C7—N2—N1	116.1 (2)	O4—C12—C13	117.9 (3)
O1—C1—C6	118.1 (3)	O4—C12—C11	122.4 (3)
O1—C1—C2	122.5 (3)	C13—C12—C11	119.5 (3)
C6—C1—C2	119.4 (3)	O5—C13—C14	124.3 (3)
C3—C2—C1	118.5 (3)	O5—C13—C12	115.9 (3)
C3—C2—C7	118.9 (3)	C14—C13—C12	119.8 (3)
C1—C2—C7	122.5 (3)	C13—C14—C9	120.3 (3)
C4—C3—C2	120.8 (3)	C13—C14—H14	119.9
C4—C3—H3	119.6	C9—C14—H14	119.9
C2—C3—H3	119.6	O3—C15—H15A	109.5
C3—C4—C5	120.2 (3)	O3—C15—H15B	109.5
C3—C4—Br1	119.8 (3)	H15A—C15—H15B	109.5
C5—C4—Br1	120.0 (3)	O3—C15—H15C	109.5
C6—C5—C4	119.9 (3)	H15A—C15—H15C	109.5
C6—C5—H5	120.1	H15B—C15—H15C	109.5
C4—C5—H5	120.1	O4—C16—H16A	109.5
C5—C6—C1	121.2 (3)	O4—C16—H16B	109.5
C5—C6—H6	119.4	H16A—C16—H16B	109.5
C1—C6—H6	119.4	O4—C16—H16C	109.5
N2—C7—C2	121.6 (3)	H16A—C16—H16C	109.5
N2—C7—H7	119.2	H16B—C16—H16C	109.5
C2—C7—H7	119.2	O5—C17—H17A	109.5
O2—C8—N1	123.0 (2)	O5—C17—H17B	109.5
O2—C8—C9	123.3 (3)	H17A—C17—H17B	109.5
N1—C8—C9	113.7 (2)	O5—C17—H17C	109.5
C10—C9—C14	120.4 (3)	H17A—C17—H17C	109.5
C10—C9—C8	118.2 (3)	H17B—C17—H17C	109.5

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1—H1 \cdots N2	0.82	1.92	2.642 (3)	145
N1—H1A \cdots O2 ⁱ	0.86	2.14	2.888 (3)	146

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