

(E)-N'-(3,5-Dibromo-2-hydroxybenzylidene)-2-nitrobenzohydrazide methanol solvate

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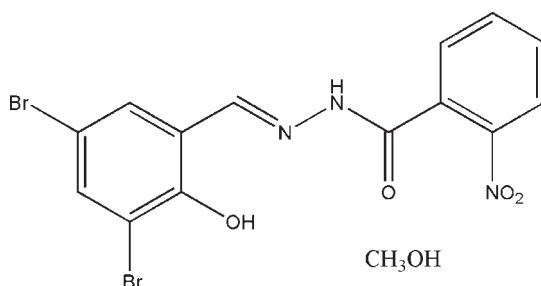
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Key indicators: single-crystal X-ray study; $T = 298\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; R factor = 0.036; wR factor = 0.094; data-to-parameter ratio = 16.3.

In the title compound, $\text{C}_{14}\text{H}_9\text{Br}_2\text{N}_3\text{O}_4\cdot\text{CH}_3\text{OH}$, the Schiff base molecule adopts an *E* geometry with respect to the $\text{C}=\text{N}$ bond and the benzene rings are oriented at a dihedral angle of $45.3(2)^\circ$. An intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond helps to establish the conformation. In the crystal, the methanol solvent molecule is linked to the Schiff base molecule through an $\text{O}-\text{H}\cdots\text{O}$ hydrogen bond and intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds link the components to form layers parallel to the *bc* direction.

Related literature

For our previous work in this area, see: Yin, Qian *et al.* (2007); Yin, Guo *et al.* (2007); Qian *et al.* (2009).



Experimental

Crystal data

$\text{C}_{14}\text{H}_9\text{Br}_2\text{N}_3\text{O}_4\cdot\text{CH}_3\text{OH}$
 $M_r = 475.10$
Monoclinic, $C2/c$
 $a = 18.981(1)\text{ \AA}$
 $b = 10.054(2)\text{ \AA}$
 $c = 19.746(2)\text{ \AA}$
 $\beta = 110.974(2)^\circ$

$V = 3518.6(8)\text{ \AA}^3$
 $Z = 8$
Mo $K\alpha$ radiation
 $\mu = 4.64\text{ mm}^{-1}$
 $T = 298\text{ K}$
 $0.18 \times 0.17 \times 0.16\text{ mm}$

Data collection

Bruker SMART CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2001)
 $T_{\min} = 0.490$, $T_{\max} = 0.524$

10461 measured reflections
3784 independent reflections
2670 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.094$
 $S = 1.02$
3784 reflections
232 parameters
1 restraint

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.36\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.66\text{ e \AA}^{-3}$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1 \cdots N1	0.82	1.87	2.587 (3)	146
O5—H5 \cdots O2	0.82	1.94	2.735 (4)	165
N2—H2 \cdots O5 ⁱ	0.893 (10)	1.958 (13)	2.840 (3)	169 (4)

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

Data collection: *SMART* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; 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*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5053).

References

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supporting information

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S1. Comment

As part of our ongoing studies of Schiff bases (Yin, Qian *et al.*, 2007; Yin, Guo *et al.*, 2007; Qian *et al.*, 2009), we now report the synthesis and structure of the title compound, (I), (Fig. 1).

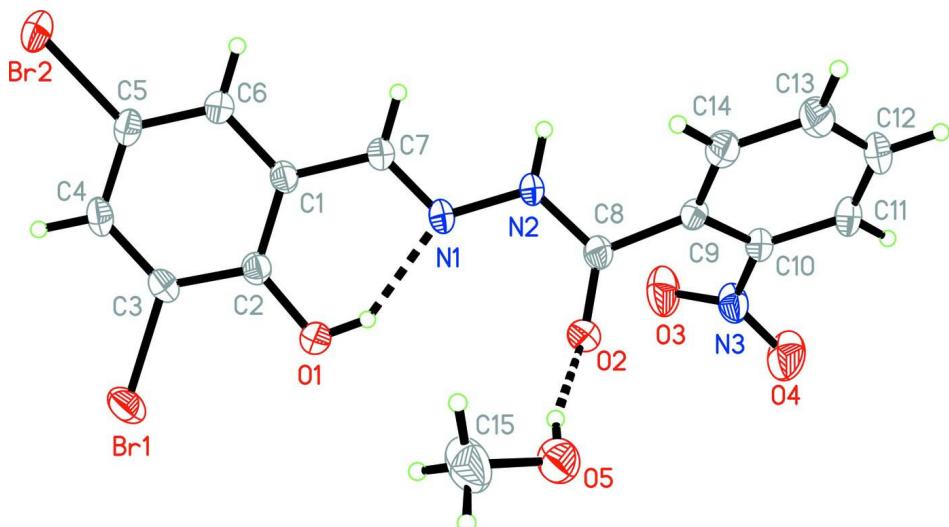
The Schiff base molecule adopts an *E* geometry with respect to the C=N bond, and there forms an intramolecular O—H···N hydrogen bond. The two benzene rings forms a dihedral angle of 45.3 (2)°. The dihedral angle between the O3/N3/O4 plane and the C9—C14 benzene ring is 37.1 (2)°. The methanol molecule is linked to the Schiff base molecule through the O—H···O hydrogen bond (Table 1). In the crystal structure, molecules are linked through intermolecular N—H···O hydrogen bonds (Table 1) to form layers parallel to the *bc* direction (Fig. 2).

S2. Experimental

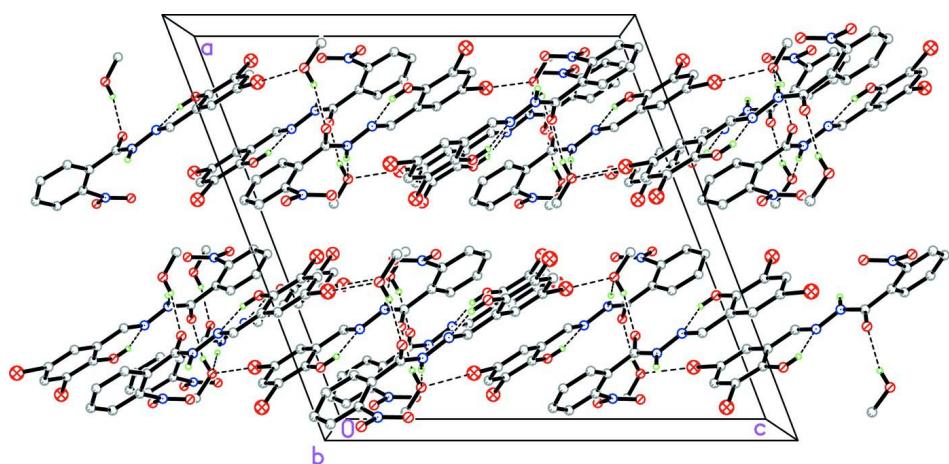
2-Nitrobenzohydrazide (1 mmol, 0.181 g) and 3,5-dibromosalicylaldehyde (1 mmol, 0.280 g) were dissolved in anhydrous methanol (15 ml). The mixture was stirred for several minutes at room temperature. The product was isolated and recrystallized from methanol, colourless blocks of (I) were obtained after five days.

S3. Refinement

The imino H atom was located in a difference map and its positional parameters were refined with a fixed isotropic thermal parameter of 0.08 Å². Other H atoms were positioned geometrically and refined as riding with C—H = 0.93 Å (aromatic) and 0.96 Å (methyl), O—H = 0.82 Å, and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{C}15 \text{ and O})$.

**Figure 1**

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level. Hydrogen bonding is shown as dashed lines.

**Figure 2**

The molecular packing of the title compound, viewed along the *b* axis. Hydrogen bonding is shown in dashed lines.

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Crystal data

$C_{14}H_9Br_2N_3O_4 \cdot CH_4O$

$M_r = 475.10$

Monoclinic, $C2/c$

Hall symbol: -C 2yc

$a = 18.981 (1) \text{ \AA}$

$b = 10.054 (2) \text{ \AA}$

$c = 19.746 (2) \text{ \AA}$

$\beta = 110.974 (2)^\circ$

$V = 3518.6 (8) \text{ \AA}^3$

$Z = 8$

$F(000) = 1872$

$D_x = 1.794 \text{ Mg m}^{-3}$

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

Cell parameters from 3081 reflections

$\theta = 2.7-25.0^\circ$

$\mu = 4.64 \text{ mm}^{-1}$

$T = 298 \text{ K}$

Block, colourless

$0.18 \times 0.17 \times 0.16 \text{ mm}$

Data collection

Bruker SMART CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2001)
 $T_{\min} = 0.490$, $T_{\max} = 0.524$

10461 measured reflections
3784 independent reflections
2670 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$
 $\theta_{\max} = 26.9^\circ$, $\theta_{\min} = 2.2^\circ$
 $h = -24 \rightarrow 20$
 $k = -12 \rightarrow 12$
 $l = -19 \rightarrow 25$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.094$
 $S = 1.02$
3784 reflections
232 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 atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0488P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.36 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.66 \text{ e } \text{\AA}^{-3}$

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	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.07056 (2)	0.15611 (4)	0.382792 (19)	0.05768 (14)
Br2	0.14655 (2)	0.69346 (4)	0.365230 (19)	0.06064 (15)
O1	0.16461 (14)	0.1962 (2)	0.53986 (11)	0.0507 (6)
H1	0.1901	0.2081	0.5828	0.076*
O2	0.26477 (13)	0.1296 (2)	0.74915 (11)	0.0556 (6)
O3	0.42343 (15)	0.0570 (3)	0.78985 (13)	0.0758 (8)
O4	0.43759 (18)	-0.0577 (3)	0.88592 (14)	0.0873 (9)
O5	0.12161 (14)	0.0782 (2)	0.74447 (13)	0.0640 (7)
H5	0.1605	0.1003	0.7384	0.096*
N1	0.25316 (14)	0.3244 (2)	0.65150 (13)	0.0403 (6)
N2	0.30067 (15)	0.3314 (2)	0.72253 (13)	0.0421 (6)
N3	0.42213 (15)	0.0446 (3)	0.85052 (15)	0.0526 (7)
C1	0.20261 (16)	0.4236 (3)	0.53512 (14)	0.0357 (6)
C2	0.16312 (16)	0.3082 (3)	0.50269 (15)	0.0369 (7)
C3	0.12163 (17)	0.3109 (3)	0.42817 (15)	0.0391 (7)

C4	0.11755 (16)	0.4237 (3)	0.38783 (15)	0.0429 (7)
H4	0.0893	0.4242	0.3384	0.052*
C5	0.15566 (17)	0.5361 (3)	0.42122 (15)	0.0422 (7)
C6	0.19789 (16)	0.5366 (3)	0.49391 (15)	0.0405 (7)
H6	0.2235	0.6133	0.5156	0.049*
C7	0.24867 (16)	0.4255 (3)	0.61175 (15)	0.0390 (7)
H7	0.2753	0.5021	0.6320	0.047*
C8	0.30280 (16)	0.2312 (3)	0.76752 (16)	0.0384 (7)
C9	0.35228 (16)	0.2548 (3)	0.84468 (15)	0.0365 (7)
C10	0.40432 (17)	0.1623 (3)	0.88545 (15)	0.0391 (7)
C11	0.4444 (2)	0.1791 (3)	0.95794 (16)	0.0517 (9)
H11	0.4787	0.1150	0.9840	0.062*
C12	0.4331 (2)	0.2913 (4)	0.99107 (18)	0.0608 (10)
H12	0.4601	0.3039	1.0402	0.073*
C13	0.3827 (2)	0.3856 (4)	0.95328 (18)	0.0622 (10)
H13	0.3756	0.4621	0.9766	0.075*
C14	0.34200 (19)	0.3670 (3)	0.87998 (16)	0.0494 (8)
H14	0.3074	0.4312	0.8544	0.059*
C15	0.0603 (2)	0.1292 (4)	0.6894 (2)	0.0898 (15)
H15A	0.0148	0.0934	0.6928	0.135*
H15B	0.0634	0.1056	0.6435	0.135*
H15C	0.0599	0.2243	0.6937	0.135*
H2	0.3299 (18)	0.403 (2)	0.7369 (19)	0.080*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0549 (2)	0.0620 (2)	0.0483 (2)	-0.01531 (16)	0.00887 (17)	-0.01568 (16)
Br2	0.0684 (3)	0.0597 (2)	0.0450 (2)	0.00128 (17)	0.00967 (18)	0.02143 (16)
O1	0.0630 (16)	0.0411 (12)	0.0378 (12)	-0.0092 (10)	0.0055 (11)	0.0045 (10)
O2	0.0565 (15)	0.0490 (13)	0.0457 (13)	-0.0159 (11)	-0.0007 (11)	0.0061 (10)
O3	0.095 (2)	0.091 (2)	0.0403 (15)	0.0193 (16)	0.0228 (14)	-0.0040 (13)
O4	0.131 (3)	0.0544 (16)	0.0632 (17)	0.0358 (16)	0.0182 (17)	0.0122 (14)
O5	0.0643 (17)	0.0531 (15)	0.0629 (15)	0.0092 (13)	0.0086 (13)	0.0013 (12)
N1	0.0377 (15)	0.0457 (15)	0.0276 (12)	-0.0015 (10)	-0.0005 (11)	0.0039 (10)
N2	0.0436 (16)	0.0422 (15)	0.0268 (12)	-0.0059 (11)	-0.0041 (11)	0.0071 (11)
N3	0.0528 (18)	0.0576 (18)	0.0380 (16)	0.0095 (13)	0.0051 (13)	0.0014 (13)
C1	0.0332 (17)	0.0404 (16)	0.0293 (14)	-0.0005 (12)	0.0062 (12)	0.0007 (12)
C2	0.0336 (17)	0.0408 (16)	0.0339 (15)	0.0000 (12)	0.0091 (13)	0.0012 (13)
C3	0.0345 (17)	0.0461 (17)	0.0333 (16)	-0.0041 (12)	0.0080 (13)	-0.0061 (13)
C4	0.0366 (18)	0.061 (2)	0.0263 (15)	0.0010 (14)	0.0054 (13)	0.0032 (14)
C5	0.0431 (19)	0.0491 (19)	0.0317 (15)	0.0051 (13)	0.0100 (13)	0.0102 (13)
C6	0.0414 (18)	0.0387 (17)	0.0362 (16)	-0.0008 (13)	0.0075 (13)	0.0038 (13)
C7	0.0398 (18)	0.0382 (16)	0.0313 (15)	-0.0013 (12)	0.0034 (13)	0.0032 (13)
C8	0.0325 (17)	0.0425 (17)	0.0343 (15)	0.0014 (13)	0.0049 (13)	0.0040 (13)
C9	0.0386 (17)	0.0378 (15)	0.0308 (14)	-0.0031 (12)	0.0097 (13)	0.0061 (12)
C10	0.0412 (19)	0.0422 (17)	0.0300 (15)	-0.0001 (12)	0.0078 (13)	0.0036 (12)
C11	0.055 (2)	0.058 (2)	0.0309 (17)	0.0027 (15)	0.0018 (15)	0.0105 (15)

C12	0.081 (3)	0.064 (2)	0.0268 (16)	-0.0072 (19)	0.0065 (17)	-0.0010 (16)
C13	0.093 (3)	0.051 (2)	0.041 (2)	-0.0017 (19)	0.022 (2)	-0.0061 (16)
C14	0.059 (2)	0.0438 (19)	0.0419 (19)	0.0029 (15)	0.0138 (17)	0.0044 (14)
C15	0.080 (3)	0.075 (3)	0.081 (3)	0.018 (2)	-0.013 (3)	-0.005 (2)

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

Br1—C3	1.882 (3)	C4—C5	1.377 (4)
Br2—C5	1.902 (3)	C4—H4	0.9300
O1—C2	1.338 (3)	C5—C6	1.371 (4)
O1—H1	0.8200	C6—H6	0.9300
O2—C8	1.228 (4)	C7—H7	0.9300
O3—N3	1.214 (3)	C8—C9	1.495 (4)
O4—N3	1.218 (4)	C9—C14	1.376 (5)
O5—C15	1.376 (4)	C9—C10	1.385 (4)
O5—H5	0.8200	C10—C11	1.370 (4)
N1—C7	1.269 (4)	C11—C12	1.359 (5)
N1—N2	1.371 (3)	C11—H11	0.9300
N2—C8	1.335 (4)	C12—C13	1.364 (5)
N2—H2	0.893 (10)	C12—H12	0.9300
N3—C10	1.469 (4)	C13—C14	1.388 (4)
C1—C6	1.382 (4)	C13—H13	0.9300
C1—C2	1.405 (4)	C14—H14	0.9300
C1—C7	1.452 (4)	C15—H15A	0.9600
C2—C3	1.399 (4)	C15—H15B	0.9600
C3—C4	1.372 (4)	C15—H15C	0.9600
C2—O1—H1	109.5	C1—C7—H7	119.5
C15—O5—H5	109.5	O2—C8—N2	123.8 (3)
C7—N1—N2	117.7 (2)	O2—C8—C9	121.4 (3)
C8—N2—N1	119.6 (2)	N2—C8—C9	114.6 (3)
C8—N2—H2	122 (2)	C14—C9—C10	117.2 (3)
N1—N2—H2	118 (2)	C14—C9—C8	119.7 (3)
O3—N3—O4	124.6 (3)	C10—C9—C8	122.9 (3)
O3—N3—C10	118.0 (3)	C11—C10—C9	122.6 (3)
O4—N3—C10	117.3 (3)	C11—C10—N3	117.0 (3)
C6—C1—C2	120.0 (3)	C9—C10—N3	120.3 (3)
C6—C1—C7	119.4 (3)	C12—C11—C10	118.7 (3)
C2—C1—C7	120.7 (3)	C12—C11—H11	120.6
O1—C2—C3	119.1 (3)	C10—C11—H11	120.6
O1—C2—C1	122.8 (2)	C11—C12—C13	121.0 (3)
C3—C2—C1	118.1 (3)	C11—C12—H12	119.5
C4—C3—C2	121.4 (3)	C13—C12—H12	119.5
C4—C3—Br1	119.6 (2)	C12—C13—C14	119.8 (3)
C2—C3—Br1	119.0 (2)	C12—C13—H13	120.1
C3—C4—C5	119.3 (3)	C14—C13—H13	120.1
C3—C4—H4	120.3	C9—C14—C13	120.8 (3)
C5—C4—H4	120.3	C9—C14—H14	119.6

C6—C5—C4	121.0 (3)	C13—C14—H14	119.6
C6—C5—Br2	120.2 (2)	O5—C15—H15A	109.5
C4—C5—Br2	118.8 (2)	O5—C15—H15B	109.5
C5—C6—C1	120.3 (3)	H15A—C15—H15B	109.5
C5—C6—H6	119.9	O5—C15—H15C	109.5
C1—C6—H6	119.9	H15A—C15—H15C	109.5
N1—C7—C1	121.1 (3)	H15B—C15—H15C	109.5
N1—C7—H7	119.5		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···N1	0.82	1.87	2.587 (3)	146
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N2—H2···O5 ⁱ	0.89 (1)	1.96 (1)	2.840 (3)	169 (4)

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