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## Structure Reports

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3-Bromo-*N'*-[(*E*)-4-hydroxybenzylidene]-benzohydrazide

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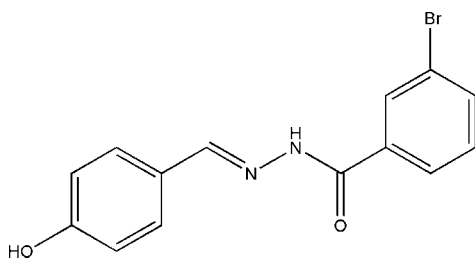
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Key indicators: single-crystal X-ray study;  $T = 298$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.031;  $wR$  factor = 0.060; data-to-parameter ratio = 15.7.

The title compound,  $\text{C}_{14}\text{H}_{11}\text{BrN}_2\text{O}_2$ , was synthesized by the reaction of 4-hydroxybenzaldehyde with an equimolar quantity of 3-bromobenzohydrazide in methanol. The dihedral angle between the two benzene rings is  $40.1(2)^\circ$ . In the crystal structure, molecules are linked through intermolecular  $\text{O}-\text{H}\cdots\text{O}$ ,  $\text{O}-\text{H}\cdots\text{N}$  and  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds to form a three-dimensional network.

## Related literature

For related structures, see: Cao (2007*a,b*).

## Experimental

## Crystal data

 $\text{C}_{14}\text{H}_{11}\text{BrN}_2\text{O}_2$  $M_r = 319.16$ Orthorhombic,  $P2_12_12_1$  $a = 7.5576(11)$  Å $b = 11.7337(18)$  Å $c = 15.021(2)$  Å $V = 1332.0(3)$  Å<sup>3</sup> $Z = 4$ Mo  $K\alpha$  radiation  
 $\mu = 3.08$  mm<sup>-1</sup> $T = 298(2)$  K  
 $0.20 \times 0.17 \times 0.16$  mm

## Data collection

Bruker SMART CCD area-detector diffractometer

Absorption correction: multi-scan

(SADABS; Bruker, 2001)

 $T_{\min} = 0.577$ ,  $T_{\max} = 0.638$ 7740 measured reflections  
2757 independent reflections  
2145 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.038$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.030$  $wR(F^2) = 0.059$  $S = 0.97$ 

2757 reflections

176 parameters

1 restraint

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{\max} = 0.17$  e Å<sup>-3</sup> $\Delta\rho_{\min} = -0.29$  e Å<sup>-3</sup>Absolute structure: Flack (1983), with 1154 Friedel pairs  
Flack parameter: 0.006(9)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O1}-\text{H1}\cdots\text{O2}^{\text{i}}$	0.82	1.95	2.750 (2)	166
$\text{O1}-\text{H1}\cdots\text{N1}^{\text{i}}$	0.82	2.56	3.003 (3)	116
$\text{N2}-\text{H2A}\cdots\text{O1}^{\text{ii}}$	0.904 (10)	2.136 (14)	3.007 (3)	162 (3)

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

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.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SJ2509).

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## supporting information

*Acta Cryst.* (2008). E64, o1186 [doi:10.1107/S1600536808015912]

### 3-Bromo-*N'*-[(*E*)-4-hydroxybenzylidene]benzohydrazide

Tao Yang, Guo-Biao Cao, Ji-Ming Xiang and Li-Hui Zhang

#### S1. Comment

We have recently reported some transition metal complexes with Schiff base ligands (Cao, 2007*a,b*). We report herein the crystal structure of the title compound, (I), derived from the reaction of 4-hydroxybenzaldehyde with an equimolar quantity of 3-bromobenzohydrazide in methanol.

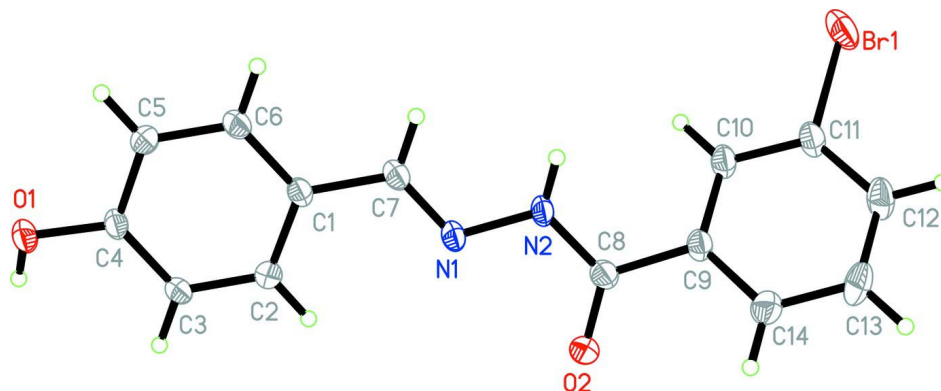
In compound (I), Fig. 1, the dihedral angle between the two benzene rings is 40.1 (2)°. In the crystal structure, molecules are linked through intermolecular O—H···O, O—H···N, and N—H···O hydrogen bonds, Table 1, to form a three-dimensional network, Figure 2.

#### S2. Experimental

The compound was prepared by refluxing equimolar quantities of 4-hydroxybenzaldehyde with 3-bromobenzohydrazide in methanol. Colourless block-like crystals were formed when the solution was evaporated in air for about a week.

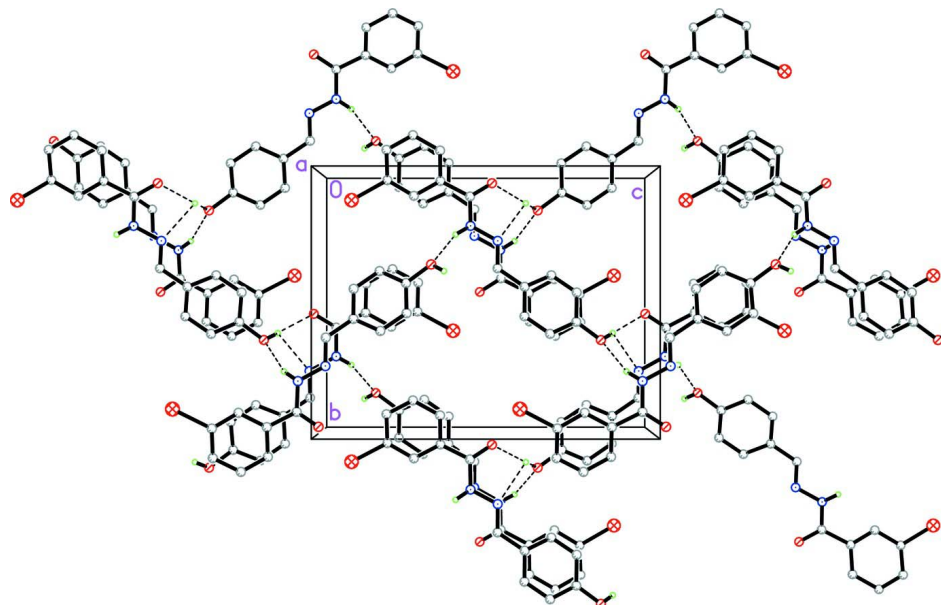
#### S3. Refinement

H2A was located in a difference Fourier map and refined isotropically, with the N—H distance restrained to 0.90 (1) Å. The other H atoms were placed in idealized positions and constrained to ride on their parent atoms, with C—H distances of 0.93 Å, the O—H distance 0.82 Å, and with  $U_{\text{iso}}(\text{H})$  set at  $1.2U_{\text{eq}}(\text{C})$  and  $1.5U_{\text{eq}}(\text{O})$ .



**Figure 1**

The molecular structure of (I) with ellipsoids drawn at the 30% probability level.

**Figure 2**

The molecular packing of (I), viewed along the *a* axis. Hydrogen bonds are drawn as dashed lines.

### 3-Bromo-*N'*-[(*E*)-4-hydroxybenzylidene]benzohydrazide

#### Crystal data

$C_{14}H_{11}BrN_2O_2$

$M_r = 319.16$

Orthorhombic,  $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 7.5576$  (11) Å

$b = 11.7337$  (18) Å

$c = 15.021$  (2) Å

$V = 1332.0$  (3) Å<sup>3</sup>

$Z = 4$

$F(000) = 640$

$D_x = 1.591$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å

Cell parameters from 1827 reflections

$\theta = 2.3$ – $24.5^\circ$

$\mu = 3.09$  mm<sup>-1</sup>

$T = 298$  K

Block, colourless

$0.20 \times 0.17 \times 0.16$  mm

#### Data collection

Bruker SMART CCD area-detector  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\omega$  scans

Absorption correction: multi-scan  
(*SADABS*; Bruker, 2001)

$T_{\min} = 0.577$ ,  $T_{\max} = 0.638$

7740 measured reflections

2757 independent reflections

2145 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.038$

$\theta_{\max} = 26.6^\circ$ ,  $\theta_{\min} = 2.2^\circ$

$h = -9 \rightarrow 9$

$k = -10 \rightarrow 14$

$l = -18 \rightarrow 18$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.031$

$wR(F^2) = 0.059$

$S = 0.97$

2757 reflections

176 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0151P)^2]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.17 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.29 \text{ e } \text{\AA}^{-3}$   
Absolute structure: Flack (1983), with 1154  
Friedel pairs  
Absolute structure parameter: 0.006 (9)

*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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.04594 (5)	0.39161 (3)	0.901456 (17)	0.06805 (13)
O1	0.1347 (3)	-0.14174 (14)	0.15827 (10)	0.0414 (5)
H1	0.0663	-0.1163	0.1208	0.062*
O2	0.0446 (3)	0.45672 (13)	0.48129 (10)	0.0435 (4)
N1	0.1134 (3)	0.23480 (18)	0.46451 (12)	0.0380 (6)
N2	0.1329 (3)	0.29110 (18)	0.54514 (12)	0.0372 (5)
C1	0.1401 (3)	0.0602 (2)	0.38450 (14)	0.0310 (6)
C2	0.0757 (3)	0.1044 (2)	0.30437 (14)	0.0343 (6)
H2	0.0339	0.1789	0.3024	0.041*
C3	0.0735 (3)	0.03889 (19)	0.22853 (14)	0.0331 (6)
H3	0.0315	0.0692	0.1754	0.040*
C4	0.1341 (3)	-0.0726 (2)	0.23148 (14)	0.0309 (6)
C5	0.1975 (3)	-0.1177 (2)	0.31015 (15)	0.0357 (6)
H5	0.2381	-0.1925	0.3120	0.043*
C6	0.2003 (3)	-0.0514 (2)	0.38585 (15)	0.0352 (6)
H6	0.2432	-0.0820	0.4387	0.042*
C7	0.1494 (3)	0.1299 (2)	0.46504 (15)	0.0348 (6)
H7	0.1829	0.0958	0.5183	0.042*
C8	0.0976 (3)	0.4039 (2)	0.54681 (14)	0.0339 (6)
C9	0.1227 (3)	0.4615 (2)	0.63473 (15)	0.0333 (6)
C10	0.0837 (3)	0.4069 (2)	0.71421 (14)	0.0373 (6)
H10	0.0457	0.3315	0.7143	0.045*
C11	0.1020 (4)	0.4657 (2)	0.79290 (15)	0.0405 (7)
C12	0.1571 (4)	0.5781 (2)	0.79396 (18)	0.0482 (8)
H12	0.1689	0.6169	0.8476	0.058*
C13	0.1942 (4)	0.6314 (2)	0.7149 (2)	0.0516 (8)
H13	0.2320	0.7068	0.7151	0.062*
C14	0.1760 (4)	0.5743 (2)	0.63504 (17)	0.0425 (7)
H14	0.1995	0.6116	0.5817	0.051*
H2A	0.195 (4)	0.257 (2)	0.5890 (15)	0.080*

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Br1	0.0960 (3)	0.0793 (2)	0.02888 (14)	0.0102 (2)	0.00631 (17)	-0.00726 (16)
O1	0.0589 (13)	0.0376 (10)	0.0276 (8)	0.0034 (9)	-0.0034 (9)	-0.0093 (8)
O2	0.0655 (12)	0.0350 (10)	0.0301 (8)	0.0023 (10)	-0.0072 (10)	0.0018 (8)
N1	0.0566 (16)	0.0338 (13)	0.0237 (10)	0.0039 (11)	-0.0060 (10)	-0.0063 (9)
N2	0.0581 (16)	0.0306 (13)	0.0230 (11)	0.0104 (11)	-0.0077 (10)	-0.0070 (9)
C1	0.0354 (14)	0.0340 (13)	0.0236 (13)	-0.0019 (11)	-0.0012 (10)	-0.0033 (10)
C2	0.0435 (16)	0.0295 (12)	0.0298 (11)	-0.0011 (13)	0.0013 (11)	-0.0003 (11)
C3	0.0424 (16)	0.0334 (14)	0.0235 (11)	0.0021 (12)	-0.0035 (11)	0.0015 (10)
C4	0.0343 (14)	0.0348 (15)	0.0238 (12)	-0.0046 (11)	0.0011 (11)	-0.0070 (10)
C5	0.0470 (17)	0.0258 (14)	0.0343 (12)	0.0036 (13)	-0.0010 (11)	-0.0011 (12)
C6	0.0452 (16)	0.0362 (14)	0.0243 (13)	0.0032 (11)	-0.0068 (11)	0.0013 (11)
C7	0.0414 (15)	0.0393 (17)	0.0238 (11)	0.0003 (13)	-0.0026 (11)	-0.0027 (11)
C8	0.0370 (16)	0.0361 (15)	0.0287 (12)	-0.0032 (13)	0.0000 (10)	-0.0013 (12)
C9	0.0362 (15)	0.0342 (15)	0.0297 (12)	0.0039 (12)	-0.0045 (11)	-0.0069 (11)
C10	0.0434 (17)	0.0381 (14)	0.0304 (12)	0.0049 (13)	-0.0038 (11)	-0.0097 (12)
C11	0.0450 (18)	0.0456 (17)	0.0308 (13)	0.0094 (13)	-0.0012 (11)	-0.0093 (12)
C12	0.0512 (19)	0.055 (2)	0.0388 (15)	0.0096 (15)	-0.0131 (14)	-0.0196 (14)
C13	0.054 (2)	0.0368 (17)	0.0644 (19)	0.0002 (14)	-0.0103 (15)	-0.0221 (15)
C14	0.0483 (18)	0.0370 (17)	0.0421 (15)	-0.0020 (13)	-0.0031 (13)	-0.0037 (12)

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

Br1—C11	1.896 (3)	C5—C6	1.378 (3)
O1—C4	1.367 (3)	C5—H5	0.9300
O1—H1	0.8200	C6—H6	0.9300
O2—C8	1.230 (3)	C7—H7	0.9300
N1—C7	1.260 (3)	C8—C9	1.495 (3)
N1—N2	1.387 (2)	C9—C14	1.384 (4)
N2—C8	1.351 (3)	C9—C10	1.387 (3)
N2—H2A	0.904 (10)	C10—C11	1.376 (3)
C1—C6	1.386 (3)	C10—H10	0.9300
C1—C2	1.398 (3)	C11—C12	1.382 (4)
C1—C7	1.462 (3)	C12—C13	1.371 (4)
C2—C3	1.374 (3)	C12—H12	0.9300
C2—H2	0.9300	C13—C14	1.381 (4)
C3—C4	1.386 (3)	C13—H13	0.9300
C3—H3	0.9300	C14—H14	0.9300
C4—C5	1.381 (3)		
C4—O1—H1	109.5	N1—C7—H7	119.0
C7—N1—N2	115.88 (19)	C1—C7—H7	119.0
C8—N2—N1	117.51 (19)	O2—C8—N2	122.9 (2)
C8—N2—H2A	121.7 (19)	O2—C8—C9	121.4 (2)
N1—N2—H2A	119 (2)	N2—C8—C9	115.7 (2)
C6—C1—C2	118.5 (2)	C14—C9—C10	120.1 (2)

C6—C1—C7	120.0 (2)	C14—C9—C8	118.2 (2)
C2—C1—C7	121.4 (2)	C10—C9—C8	121.7 (2)
C3—C2—C1	120.7 (2)	C11—C10—C9	119.1 (2)
C3—C2—H2	119.7	C11—C10—H10	120.5
C1—C2—H2	119.7	C9—C10—H10	120.5
C2—C3—C4	119.8 (2)	C10—C11—C12	121.3 (2)
C2—C3—H3	120.1	C10—C11—Br1	119.1 (2)
C4—C3—H3	120.1	C12—C11—Br1	119.65 (19)
O1—C4—C5	117.4 (2)	C13—C12—C11	119.1 (2)
O1—C4—C3	122.4 (2)	C13—C12—H12	120.4
C5—C4—C3	120.2 (2)	C11—C12—H12	120.4
C6—C5—C4	119.7 (2)	C12—C13—C14	120.7 (3)
C6—C5—H5	120.2	C12—C13—H13	119.6
C4—C5—H5	120.2	C14—C13—H13	119.6
C5—C6—C1	121.1 (2)	C13—C14—C9	119.7 (3)
C5—C6—H6	119.5	C13—C14—H14	120.1
C1—C6—H6	119.5	C9—C14—H14	120.1
N1—C7—C1	122.1 (2)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O1—H1...O2 <sup>i</sup>	0.82	1.95	2.750 (2)	166
O1—H1...N1 <sup>i</sup>	0.82	2.56	3.003 (3)	116
N2—H2A...O1 <sup>ii</sup>	0.90 (1)	2.14 (1)	3.007 (3)	162 (3)

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