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## Structure Reports

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## (E)-N'-(4-Methoxybenzylidene)benzohydrazide

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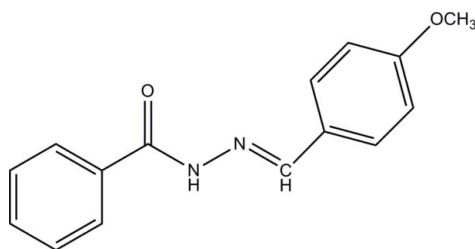
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 Key indicators: single-crystal X-ray study;  $T = 298$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.042;  $wR$  factor = 0.106; data-to-parameter ratio = 7.2.

 In the title molecule,  $\text{C}_{15}\text{H}_{14}\text{N}_2\text{O}_2$ , the dihedral angle between the benzene rings is  $5.93$  ( $17^\circ$ ). In the crystal, intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds link the molecules into chains propagating in  $[010]$ .

### Related literature

 For properties of Schiff base ligands, see: Cozzi *et al.* (2004). For related crystal structures, see: Fun *et al.* (2008); Cui *et al.* (2009); Nie (2008).


### Experimental

#### Crystal data

 $\text{C}_{15}\text{H}_{14}\text{N}_2\text{O}_2$ 
 $M_r = 254.28$ 

 Orthorhombic,  $Pca2_1$ 
 $a = 31.414$  (3) Å

 $b = 5.1067$  (5) Å

 $c = 8.1336$  (9) Å

 $V = 1304.8$  (2) Å<sup>3</sup>
 $Z = 4$ 

 Mo  $K\alpha$  radiation

 $\mu = 0.09$  mm<sup>-1</sup>
 $T = 298$  K

 $0.49 \times 0.48 \times 0.30$  mm

#### Data collection

Bruker SMART APEX CCD area-detector diffractometer

Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

 $T_{\min} = 0.958$ ,  $T_{\max} = 0.974$ 

2220 measured reflections

1239 independent reflections

 920 reflections with  $I > 2\sigma(I)$ 
 $R_{\text{int}} = 0.037$ 

#### Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.042$ 
 $wR(F^2) = 0.106$ 
 $S = 1.03$ 

1239 reflections

173 parameters

1 restraint

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.18$  e Å<sup>-3</sup>
 $\Delta\rho_{\text{min}} = -0.14$  e Å<sup>-3</sup>
**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{O1}^i$	0.86	2.17	2.961 (2)	152

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

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); 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: BQ2172).

### References

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## supporting information

*Acta Cryst.* (2009). E65, o3207 [doi:10.1107/S1600536809049988]

**(E)-N'-(4-Methoxybenzylidene)benzohydrazide**

Jian-Xia Gou, Ming-Zhi Song, Chuan-Gang Fan and Zhong-Nian Yang

**S1. Comment**

Schiff bases are popular ligands in coordination chemistry due to their ease of synthesis and their ability to be readily modified both electronically and sterically. Mixed-donor Schiff bases have been used extensively in catalysis (Cozzi, 2004).

In (I), (Fig. 1), the bond lengths and angles are normal and are comparable to the values observed in similar compounds (Nie *et al.*, 2008; Fun *et al.*, 2008; Cui *et al.*, 2009).

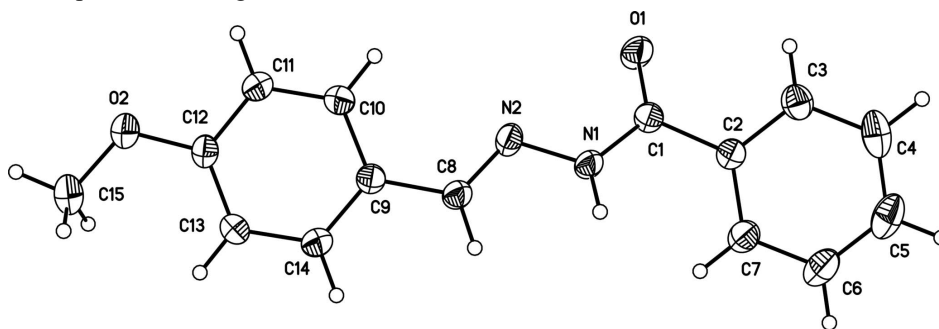
In the crystal structure, the C8=N2 bond length in the molecule is 1.269 (3) Å, showing the double-bond character. Meanwhile, the dihedral angle between the benzene ring (C2-C7) and the benzene ring (C9-C14) in the Schiff base molecule is 5.93 (17)°, indicating that the two aromatic ring planes are almost coplanar. Moreover, the crystal supramolecular structure was built from the connections of weak intermolecular N—H···O hydrogen bonds, as shown in table 1, and these hydrogen bonds link molecules into one-dimensional chains propagated in direction [010].

**S2. Experimental**

Benzohydrazide (5.0 mmol), 20 ml ethanol and 4-methoxybenzaldehyde (5.0 mmol) were mixed in 50 ml flask. After refluxing 3 h, the resulting mixture was cooled to room temperature, and recrystallized from ethanol, and afforded the title compound as a crystalline solid. Elemental analysis: calculated for C<sub>15</sub>H<sub>14</sub>N<sub>2</sub>O<sub>2</sub>: C 70.85, H 5.55, N 11.02%; found: C 70.78, H 5.64, N 11.13%.

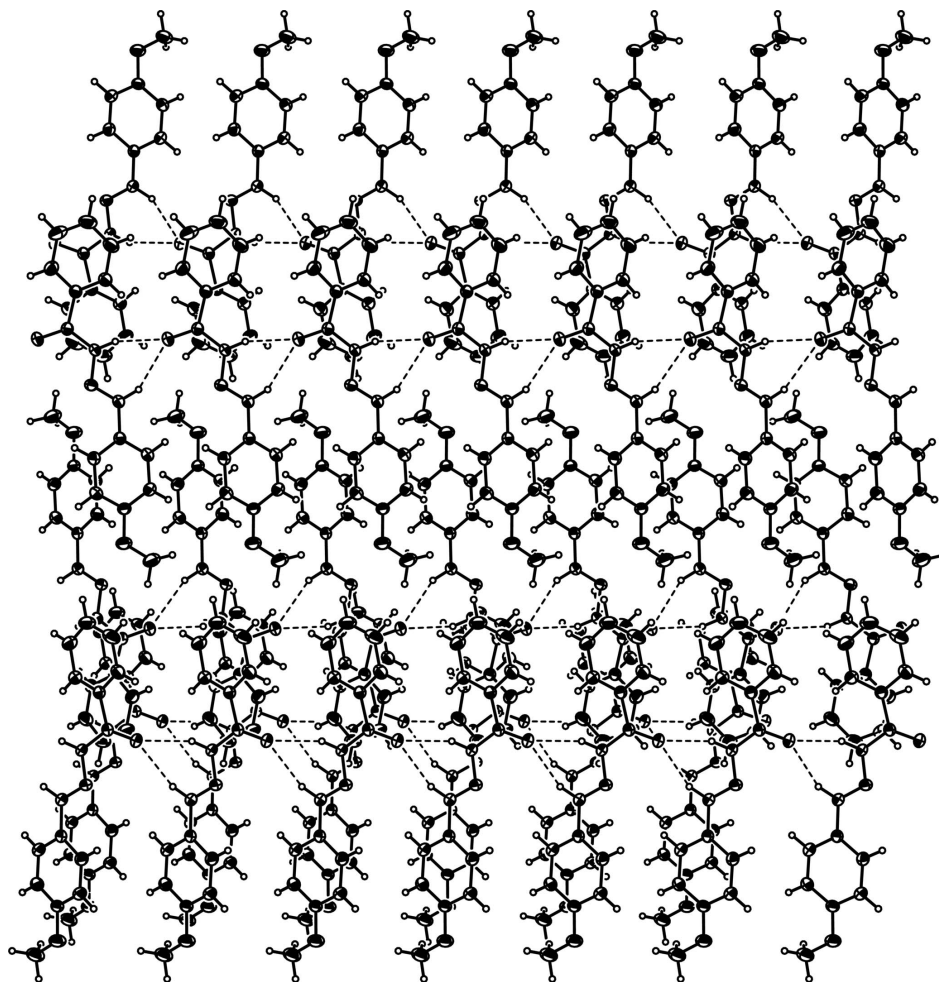
**S3. Refinement**

All H atoms were placed in geometrically idealized positions (N—H=0.86 and C—H=0.93–0.96 Å) and treated as riding on their parent atoms, with  $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5U_{\text{eq}}(\text{C})$  (C,N). Because of the meaningfulness of the absolute structure parameter, 981 Friedel-pairs were merged before final refinement.



**Figure 1**

A view of (I) showing the atomic numbering scheme and 30% probability displacement ellipsoids.

**Figure 2**

The packing of (I) built from the connections of weak intermolecular N—H...O hydrogen bonds with dashed lines.

### (*E*)-*N'*-(4-Methoxybenzylidene)benzohydrazide

#### Crystal data

$C_{15}H_{14}N_2O_2$

$M_r = 254.28$

Orthorhombic,  $Pca2_1$

Hall symbol:  $P\ 2c\ -2ac$

$a = 31.414\ (3)\ \text{\AA}$

$b = 5.1067\ (5)\ \text{\AA}$

$c = 8.1336\ (9)\ \text{\AA}$

$V = 1304.8\ (2)\ \text{\AA}^3$

$Z = 4$

$F(000) = 536$

$D_x = 1.294\ \text{Mg m}^{-3}$

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

Cell parameters from 1475 reflections

$\theta = 2.6\text{--}22.4^\circ$

$\mu = 0.09\ \text{mm}^{-1}$

$T = 298\ \text{K}$

Block, colourless

$0.49 \times 0.48 \times 0.30\ \text{mm}$

#### Data collection

Bruker SMART APEX CCD area-detector  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

phi and  $\omega$  scans

Absorption correction: multi-scan  
(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.958$ ,  $T_{\max} = 0.974$

2220 measured reflections

1239 independent reflections

920 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.037$   
 $\theta_{\text{max}} = 25.0^\circ$ ,  $\theta_{\text{min}} = 1.3^\circ$

$h = 0 \rightarrow 36$   
 $k = 0 \rightarrow 6$   
 $l = -9 \rightarrow 9$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.042$   
 $wR(F^2) = 0.106$   
 $S = 1.03$   
 1239 reflections  
 173 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  
 $w = 1/[\sigma^2(F_o^2) + (0.051P)^2 + 0.0327P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.18 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.14 \text{ e } \text{\AA}^{-3}$

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

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.32900 (6)	0.0513 (4)	0.8803 (3)	0.0446 (6)
H1	0.3182	0.2051	0.8691	0.053*
N2	0.37282 (6)	0.0199 (4)	0.8715 (3)	0.0448 (6)
O1	0.31604 (5)	-0.3787 (3)	0.9244 (3)	0.0612 (6)
O2	0.57114 (5)	0.2572 (4)	0.8020 (3)	0.0618 (6)
C1	0.30295 (7)	-0.1548 (5)	0.9062 (4)	0.0414 (6)
C2	0.25645 (8)	-0.0907 (4)	0.9049 (4)	0.0404 (6)
C3	0.22924 (9)	-0.2535 (6)	0.9909 (4)	0.0522 (8)
H3	0.2400	-0.3961	1.0487	0.063*
C4	0.18566 (10)	-0.2029 (7)	0.9905 (5)	0.0688 (11)
H4	0.1673	-0.3098	1.0501	0.083*
C5	0.16978 (9)	0.0027 (7)	0.9031 (6)	0.0707 (10)
H5	0.1407	0.0366	0.9043	0.085*
C6	0.19643 (9)	0.1599 (6)	0.8133 (5)	0.0653 (10)
H6	0.1853	0.2959	0.7505	0.078*
C7	0.23964 (8)	0.1159 (5)	0.8162 (4)	0.0506 (8)
H7	0.2577	0.2262	0.7579	0.061*
C8	0.39287 (8)	0.2202 (5)	0.8213 (4)	0.0437 (7)
H8	0.3775	0.3676	0.7898	0.052*
C9	0.43914 (7)	0.2271 (5)	0.8114 (4)	0.0399 (6)
C10	0.46450 (8)	0.0478 (5)	0.8944 (4)	0.0478 (7)
H10	0.4517	-0.0854	0.9547	0.057*

C11	0.50810 (8)	0.0635 (5)	0.8889 (4)	0.0479 (7)
H11	0.5245	-0.0577	0.9460	0.058*
C12	0.52766 (7)	0.2591 (5)	0.7988 (4)	0.0420 (6)
C13	0.50332 (9)	0.4391 (5)	0.7160 (4)	0.0472 (7)
H13	0.5163	0.5708	0.6550	0.057*
C14	0.45933 (8)	0.4235 (5)	0.7239 (4)	0.0485 (8)
H14	0.4430	0.5479	0.6691	0.058*
C15	0.59304 (9)	0.4528 (7)	0.7113 (5)	0.0735 (10)
H15A	0.5848	0.4435	0.5978	0.110*
H15B	0.6232	0.4249	0.7203	0.110*
H15C	0.5860	0.6224	0.7544	0.110*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
N1	0.0362 (12)	0.0317 (11)	0.0658 (16)	0.0055 (9)	-0.0014 (13)	0.0011 (12)
N2	0.0331 (12)	0.0412 (12)	0.0600 (16)	0.0020 (10)	-0.0012 (13)	-0.0067 (12)
O1	0.0481 (10)	0.0330 (10)	0.1026 (18)	0.0060 (8)	-0.0035 (13)	0.0002 (12)
O2	0.0379 (11)	0.0707 (13)	0.0766 (14)	-0.0043 (10)	-0.0008 (12)	0.0158 (12)
C1	0.0410 (13)	0.0339 (13)	0.0492 (17)	-0.0011 (12)	-0.0008 (16)	-0.0051 (15)
C2	0.0402 (14)	0.0369 (13)	0.0439 (17)	-0.0002 (11)	-0.0016 (15)	-0.0060 (16)
C3	0.0510 (18)	0.0496 (17)	0.056 (2)	-0.0085 (14)	0.0014 (16)	-0.0045 (16)
C4	0.050 (2)	0.082 (3)	0.075 (3)	-0.0216 (18)	0.0150 (19)	-0.018 (2)
C5	0.0399 (16)	0.079 (2)	0.093 (3)	0.0042 (16)	-0.011 (2)	-0.022 (3)
C6	0.0501 (19)	0.060 (2)	0.086 (3)	0.0124 (16)	-0.0138 (19)	-0.009 (2)
C7	0.0453 (17)	0.0448 (15)	0.062 (2)	0.0023 (13)	-0.0056 (16)	-0.0029 (17)
C8	0.0398 (14)	0.0357 (14)	0.0556 (19)	0.0035 (12)	-0.0036 (14)	0.0009 (14)
C9	0.0398 (14)	0.0342 (14)	0.0456 (16)	0.0011 (12)	0.0002 (15)	-0.0049 (14)
C10	0.0451 (15)	0.0416 (14)	0.0566 (19)	0.0012 (12)	0.0025 (17)	0.0087 (18)
C11	0.0428 (15)	0.0446 (15)	0.0563 (19)	0.0049 (12)	-0.0057 (17)	0.0068 (18)
C12	0.0336 (15)	0.0442 (15)	0.0482 (16)	-0.0017 (13)	0.0013 (15)	-0.0061 (14)
C13	0.0444 (17)	0.0428 (17)	0.0542 (19)	-0.0055 (13)	0.0028 (16)	0.0076 (15)
C14	0.0444 (18)	0.0408 (16)	0.060 (2)	0.0063 (13)	-0.0031 (16)	0.0051 (15)
C15	0.0471 (19)	0.081 (2)	0.092 (3)	-0.0165 (16)	-0.0041 (18)	0.013 (2)

*Geometric parameters (Å, °)*

N1—C1	1.350 (3)	C6—H6	0.9300
N1—N2	1.388 (2)	C7—H7	0.9300
N1—H1	0.8600	C8—C9	1.456 (3)
N2—C8	1.269 (3)	C8—H8	0.9300
O1—C1	1.224 (3)	C9—C14	1.384 (4)
O2—C12	1.366 (3)	C9—C10	1.389 (4)
O2—C15	1.420 (4)	C10—C11	1.373 (3)
C1—C2	1.497 (3)	C10—H10	0.9300
C2—C3	1.382 (4)	C11—C12	1.383 (4)
C2—C7	1.383 (3)	C11—H11	0.9300
C3—C4	1.393 (4)	C12—C13	1.372 (4)

C3—H3	0.9300	C13—C14	1.386 (4)
C4—C5	1.363 (5)	C13—H13	0.9300
C4—H4	0.9300	C14—H14	0.9300
C5—C6	1.370 (5)	C15—H15A	0.9600
C5—H5	0.9300	C15—H15B	0.9600
C6—C7	1.376 (4)	C15—H15C	0.9600
C1—N1—N2	121.29 (19)	N2—C8—H8	118.9
C1—N1—H1	119.4	C9—C8—H8	118.9
N2—N1—H1	119.4	C14—C9—C10	117.7 (2)
C8—N2—N1	114.6 (2)	C14—C9—C8	120.2 (2)
C12—O2—C15	118.0 (2)	C10—C9—C8	122.0 (3)
O1—C1—N1	122.9 (2)	C11—C10—C9	121.2 (3)
O1—C1—C2	122.2 (2)	C11—C10—H10	119.4
N1—C1—C2	114.8 (2)	C9—C10—H10	119.4
C3—C2—C7	119.1 (2)	C10—C11—C12	120.2 (3)
C3—C2—C1	117.9 (2)	C10—C11—H11	119.9
C7—C2—C1	122.9 (2)	C12—C11—H11	119.9
C2—C3—C4	119.7 (3)	O2—C12—C13	124.8 (3)
C2—C3—H3	120.2	O2—C12—C11	115.4 (2)
C4—C3—H3	120.2	C13—C12—C11	119.8 (2)
C5—C4—C3	120.3 (3)	C12—C13—C14	119.7 (3)
C5—C4—H4	119.9	C12—C13—H13	120.2
C3—C4—H4	119.9	C14—C13—H13	120.2
C4—C5—C6	120.4 (3)	C9—C14—C13	121.5 (3)
C4—C5—H5	119.8	C9—C14—H14	119.2
C6—C5—H5	119.8	C13—C14—H14	119.2
C5—C6—C7	119.9 (3)	O2—C15—H15A	109.5
C5—C6—H6	120.1	O2—C15—H15B	109.5
C7—C6—H6	120.1	H15A—C15—H15B	109.5
C6—C7—C2	120.7 (3)	O2—C15—H15C	109.5
C6—C7—H7	119.7	H15A—C15—H15C	109.5
C2—C7—H7	119.7	H15B—C15—H15C	109.5
N2—C8—C9	122.2 (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>
N1—H1...O1 <sup>i</sup>	0.86	2.17	2.961 (2)	152

Symmetry code: (i) *x*, *y*+1, *z*.