

4-Hydroxy-N'-[1-(2-hydroxyphenyl)-ethylidene]benzohydrazide

Xiao-Hui Ji and Jiu-Fu Lu*

School of Chemistry and Environmental Science, Shaanxi University of Technology, Hanzhong 723000, People's Republic of China
Correspondence e-mail: jiufulu@163.com

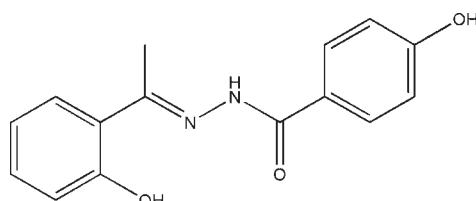
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Key indicators: single-crystal X-ray study; $T = 298\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; R factor = 0.069; wR factor = 0.172; data-to-parameter ratio = 14.7.

In the title compound, $\text{C}_{15}\text{H}_{14}\text{N}_2\text{O}_3$, there is an intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond and the dihedral angle between the two aromatic rings is $13.9(3)^\circ$. In the crystal structure, molecules are stabilized by intermolecular $\text{O}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For related structures, see: Lu *et al.* (2008a,b,c); Xiao & Wei (2009); He (2008); Shi *et al.* (2007). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{15}\text{H}_{14}\text{N}_2\text{O}_3$

$M_r = 270.28$

Monoclinic, $P2_1/c$

$a = 4.926(2)\text{ \AA}$

$b = 31.06(2)\text{ \AA}$

$c = 8.473(3)\text{ \AA}$

$\beta = 93.852(3)^\circ$

$V = 1293.5(11)\text{ \AA}^3$

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 0.10\text{ mm}^{-1}$

$T = 298\text{ K}$

$0.21 \times 0.20 \times 0.17\text{ mm}$

Data collection

Bruker APEXII CCD area-detector

diffractometer

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 2004)

$T_{\min} = 0.980$, $T_{\max} = 0.984$

8965 measured reflections

2770 independent reflections

1569 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.064$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.069$

$wR(F^2) = 0.172$

$S = 1.05$

2770 reflections

188 parameters

1 restraint

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\text{max}} = 0.26\text{ e \AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.21\text{ e \AA}^{-3}$

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	1.78	2.499 (3)	146
O3—H3 \cdots O1 ⁱ	0.82	1.97	2.786 (3)	179
N2—H2 \cdots O2 ⁱⁱ	0.90 (1)	2.09 (2)	2.961 (4)	163 (3)

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

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); 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: SU2181).

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supporting information

Acta Cryst. (2010). E66, o1514 [https://doi.org/10.1107/S1600536810019914]

4-Hydroxy-N'-[1-(2-hydroxyphenyl)ethylidene]benzohydrazide

Xiao-Hui Ji and Jiu-Fu Lu

S1. Comment

As part of our investigation of the crystal structures of Schiff bases derived from the condensation of aldehydes or ketones with benzohydrazides (Lu *et al.*, 2008a,b,c), we report herein on the crystal structure of the new title Schiff base compound.

In the title compound, illustrated in Fig. 1, the bond lengths have normal values (Allen *et al.*, 1987), and are comparable to those observed in similar compounds (Xiao & Wei, 2009; He, 2008; Shi *et al.*, 2007). The dihedral angle between the two aromatic rings is 13.9 (3) $^{\circ}$, indicating that the Schiff base molecule is slightly twisted. An intramolecular O—H \cdots N hydrogen bond is observed (Table 1).

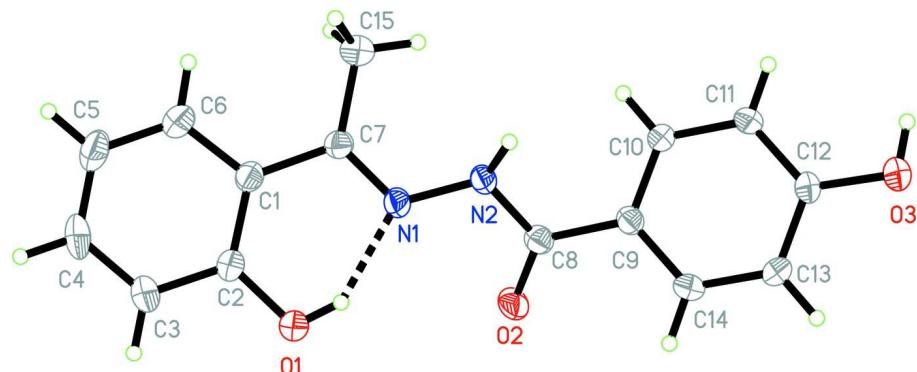
In the crystal structure, the molecules are stabilized by intermolecular O—H \cdots O and N—H \cdots O hydrogen bonds (Table 1 and Fig. 2).

S2. Experimental

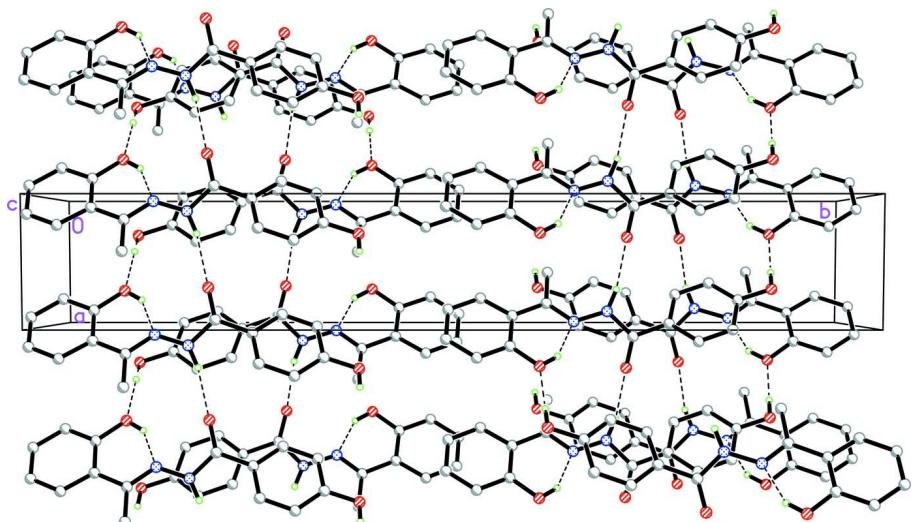
The title compound was prepared by the Schiff base condensation of 1-(2-hydroxyphenyl)ethanone (0.1 mol, 13.6 g) and 4-hydroxybenzohydrazide (0.1 mmol, 15.2 g) in 95% ethanol (70 ml). The excess ethanol was removed by distillation. The colourless solid obtained was filtered and washed with ethanol. Single crystals, suitable for X-ray diffraction analysis, were obtained by slow evaporation of a 95% ethanol solution at rt.

S3. Refinement

The N2 H-atom (H2) was located in a difference Fourier map and refined with a distance restraint: N—H = 0.90 (1) Å. The other H-atoms were positioned geometrically (C—H = 0.93–0.96 Å and O—H = 0.82 Å) and refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{C}_{\text{methyl}} \text{ and } \text{O})$. A rotating group model was used for the methyl and hydroxyl groups.

**Figure 1**

The molecular structure of the title compound, showing 30% probability displacement ellipsoids and the atom-numbering scheme. The dashed line indicates the intramolecular O-H \cdots N hydrogen bond (see Table 1 for details).

**Figure 2**

The crystal packing of the title compound, viewed along the c axis. H-atoms not involved in hydrogen bonding (dashed lines) have been omitted for clarity.

4-Hydroxy-N'-(1-(2-hydroxyphenyl)ethylidene)benzohydrazide

Crystal data

$C_{15}H_{14}N_2O_3$
 $M_r = 270.28$
Monoclinic, $P2_1/c$
Hall symbol: -P 2ybc
 $a = 4.926 (2)$ Å
 $b = 31.06 (2)$ Å
 $c = 8.473 (3)$ Å
 $\beta = 93.852 (3)^\circ$
 $V = 1293.5 (11)$ Å 3
 $Z = 4$

$F(000) = 568$
 $D_x = 1.388$ Mg m $^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 957 reflections
 $\theta = 2.5\text{--}24.5^\circ$
 $\mu = 0.10$ mm $^{-1}$
 $T = 298$ K
Block, colourless
 $0.21 \times 0.20 \times 0.17$ mm

Data collection

Bruker APEXII CCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 2004)
 $T_{\min} = 0.980$, $T_{\max} = 0.984$

8965 measured reflections
2770 independent reflections
1569 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.064$
 $\theta_{\max} = 27.0^\circ$, $\theta_{\min} = 2.5^\circ$
 $h = -6 \rightarrow 6$
 $k = -39 \rightarrow 34$
 $l = -10 \rightarrow 10$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.069$
 $wR(F^2) = 0.172$
 $S = 1.05$
2770 reflections
188 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
 $w = 1/[\sigma^2(F_o^2) + (0.0649P)^2 + 0.4306P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.26 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$
Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0057 (18)

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.9548 (5)	0.64659 (7)	0.6666 (3)	0.0384 (6)
N2	0.8786 (5)	0.68800 (7)	0.6186 (3)	0.0389 (6)
O1	1.2807 (5)	0.60332 (7)	0.8408 (3)	0.0581 (7)
H1	1.2112	0.6252	0.8021	0.087*
O2	1.3097 (4)	0.71011 (6)	0.6779 (3)	0.0510 (6)
O3	0.7088 (5)	0.89007 (6)	0.5716 (3)	0.0593 (7)
H3	0.5810	0.8920	0.5048	0.089*
C1	0.9088 (6)	0.57209 (9)	0.6782 (3)	0.0377 (7)
C2	1.1338 (6)	0.56842 (9)	0.7889 (3)	0.0413 (8)
C3	1.2167 (7)	0.52855 (10)	0.8469 (4)	0.0530 (9)
H3A	1.3668	0.5266	0.9192	0.064*
C4	1.0804 (8)	0.49197 (10)	0.7992 (4)	0.0603 (10)
H4	1.1375	0.4653	0.8389	0.072*
C5	0.8616 (8)	0.49468 (11)	0.6938 (5)	0.0644 (11)
H5	0.7681	0.4698	0.6618	0.077*

C6	0.7776 (7)	0.53373 (10)	0.6342 (4)	0.0544 (9)
H6	0.6274	0.5348	0.5617	0.065*
C7	0.8165 (6)	0.61399 (9)	0.6136 (3)	0.0366 (7)
C8	1.0701 (6)	0.71922 (9)	0.6405 (3)	0.0374 (7)
C9	0.9732 (6)	0.76376 (9)	0.6167 (3)	0.0346 (7)
C10	0.7529 (6)	0.77407 (9)	0.5133 (3)	0.0394 (7)
H10	0.6639	0.7523	0.4547	0.047*
C11	0.6634 (6)	0.81585 (9)	0.4958 (3)	0.0411 (8)
H11	0.5158	0.8222	0.4254	0.049*
C12	0.7927 (6)	0.84846 (9)	0.5827 (3)	0.0392 (7)
C13	1.0155 (6)	0.83870 (10)	0.6841 (4)	0.0477 (8)
H13	1.1056	0.8605	0.7415	0.057*
C14	1.1051 (6)	0.79695 (10)	0.7005 (4)	0.0446 (8)
H14	1.2560	0.7908	0.7687	0.054*
C15	0.5813 (7)	0.61695 (11)	0.4922 (4)	0.0541 (9)
H15A	0.4183	0.6075	0.5378	0.081*
H15B	0.6154	0.5990	0.4035	0.081*
H15C	0.5595	0.6463	0.4575	0.081*
H2	0.710 (3)	0.6991 (11)	0.622 (4)	0.080*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0324 (14)	0.0301 (14)	0.0522 (15)	0.0026 (11)	0.0000 (11)	0.0024 (11)
N2	0.0325 (14)	0.0302 (14)	0.0531 (15)	0.0022 (11)	-0.0030 (12)	0.0062 (11)
O1	0.0596 (15)	0.0364 (13)	0.0744 (16)	-0.0035 (11)	-0.0252 (12)	0.0046 (11)
O2	0.0294 (12)	0.0443 (13)	0.0784 (16)	0.0045 (10)	-0.0033 (11)	0.0088 (11)
O3	0.0686 (18)	0.0336 (13)	0.0721 (17)	0.0086 (11)	-0.0224 (13)	-0.0024 (10)
C1	0.0364 (17)	0.0327 (16)	0.0439 (16)	-0.0017 (13)	0.0030 (13)	-0.0025 (13)
C2	0.0458 (19)	0.0313 (17)	0.0464 (17)	-0.0017 (14)	0.0001 (14)	-0.0005 (13)
C3	0.062 (2)	0.0367 (19)	0.060 (2)	0.0050 (16)	-0.0040 (17)	0.0084 (15)
C4	0.077 (3)	0.0292 (19)	0.076 (2)	0.0094 (17)	0.013 (2)	0.0120 (16)
C5	0.072 (3)	0.0326 (19)	0.088 (3)	-0.0091 (18)	0.000 (2)	-0.0022 (17)
C6	0.055 (2)	0.041 (2)	0.066 (2)	-0.0074 (16)	-0.0070 (17)	-0.0049 (16)
C7	0.0340 (17)	0.0351 (16)	0.0405 (16)	0.0006 (13)	-0.0001 (13)	-0.0023 (12)
C8	0.0338 (17)	0.0386 (17)	0.0397 (16)	0.0010 (14)	0.0017 (13)	0.0043 (13)
C9	0.0314 (16)	0.0334 (16)	0.0388 (15)	0.0013 (13)	0.0017 (12)	0.0036 (12)
C10	0.0415 (18)	0.0313 (16)	0.0438 (17)	-0.0016 (13)	-0.0080 (13)	-0.0003 (13)
C11	0.0426 (18)	0.0367 (17)	0.0419 (17)	0.0008 (14)	-0.0125 (14)	0.0048 (13)
C12	0.0450 (18)	0.0277 (16)	0.0445 (17)	0.0026 (13)	-0.0013 (14)	0.0032 (13)
C13	0.046 (2)	0.0353 (18)	0.0588 (19)	-0.0043 (14)	-0.0146 (16)	-0.0076 (15)
C14	0.0380 (18)	0.0428 (18)	0.0509 (19)	0.0019 (15)	-0.0127 (14)	0.0019 (14)
C15	0.049 (2)	0.053 (2)	0.058 (2)	-0.0024 (17)	-0.0111 (16)	0.0030 (16)

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

N1—C7	1.284 (3)	C5—H5	0.9300
N1—N2	1.393 (3)	C6—H6	0.9300

N2—C8	1.357 (4)	C7—C15	1.500 (4)
N2—H2	0.900 (10)	C8—C9	1.473 (4)
O1—C2	1.360 (3)	C9—C10	1.385 (4)
O1—H1	0.8200	C9—C14	1.388 (4)
O2—C8	1.234 (3)	C10—C11	1.375 (4)
O3—C12	1.358 (3)	C10—H10	0.9300
O3—H3	0.8200	C11—C12	1.382 (4)
C1—C6	1.394 (4)	C11—H11	0.9300
C1—C2	1.407 (4)	C12—C13	1.381 (4)
C1—C7	1.472 (4)	C13—C14	1.374 (4)
C2—C3	1.384 (4)	C13—H13	0.9300
C3—C4	1.367 (4)	C14—H14	0.9300
C3—H3A	0.9300	C15—H15A	0.9600
C4—C5	1.356 (5)	C15—H15B	0.9600
C4—H4	0.9300	C15—H15C	0.9600
C5—C6	1.367 (5)		
C7—N1—N2	120.0 (2)	O2—C8—N2	121.0 (3)
C8—N2—N1	116.7 (2)	O2—C8—C9	123.1 (3)
C8—N2—H2	111 (2)	N2—C8—C9	115.9 (2)
N1—N2—H2	125 (2)	C10—C9—C14	118.1 (3)
C2—O1—H1	109.5	C10—C9—C8	122.5 (3)
C12—O3—H3	109.5	C14—C9—C8	119.4 (2)
C6—C1—C2	116.1 (3)	C11—C10—C9	121.2 (3)
C6—C1—C7	122.0 (3)	C11—C10—H10	119.4
C2—C1—C7	121.9 (2)	C9—C10—H10	119.4
O1—C2—C3	117.4 (3)	C10—C11—C12	120.1 (3)
O1—C2—C1	122.0 (2)	C10—C11—H11	120.0
C3—C2—C1	120.6 (3)	C12—C11—H11	120.0
C4—C3—C2	120.7 (3)	O3—C12—C13	118.5 (3)
C4—C3—H3A	119.6	O3—C12—C11	122.2 (3)
C2—C3—H3A	119.6	C13—C12—C11	119.3 (3)
C5—C4—C3	119.8 (3)	C14—C13—C12	120.4 (3)
C5—C4—H4	120.1	C14—C13—H13	119.8
C3—C4—H4	120.1	C12—C13—H13	119.8
C4—C5—C6	120.4 (3)	C13—C14—C9	120.9 (3)
C4—C5—H5	119.8	C13—C14—H14	119.6
C6—C5—H5	119.8	C9—C14—H14	119.6
C5—C6—C1	122.4 (3)	C7—C15—H15A	109.5
C5—C6—H6	118.8	C7—C15—H15B	109.5
C1—C6—H6	118.8	H15A—C15—H15B	109.5
N1—C7—C1	115.0 (2)	C7—C15—H15C	109.5
N1—C7—C15	124.0 (3)	H15A—C15—H15C	109.5
C1—C7—C15	121.0 (3)	H15B—C15—H15C	109.5

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···N1	0.82	1.78	2.499 (3)	146
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