Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

N′-(Furan-2-ylmethylene)-2-hydroxybenzohydrazide

Yan-Xia Zhang

Department of Materials Science and Chemical Engineering, Taishan University, 271021 Taian, Shandong, People's Republic of China
Correspondence e-mail: zhangyanxia@mail.sdu.edu.cn

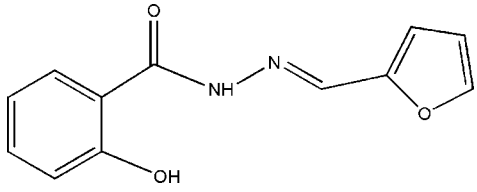
Received 20 October 2008; accepted 23 October 2008

Key indicators: single-crystal X-ray study; T = 295 K; mean σ(C–C) = 0.002 Å; R factor = 0.034; wR factor = 0.097; data-to-parameter ratio = 12.2.

In the title molecule, $C_{12}H_{10}N_2O_3$, the aromatic and furan rings form a dihedral angle of 8.89 (1)° and an intramolecular N—H···O hydrogen bond occurs. In the crystal structure, intermolecular O—H···O hydrogen bonds link the molecules into zigzag chains running along the c axis.

Related literature

For background on Schiff bases, see: Garnovskii *et al.* (1993); Anderson *et al.* (1997); Musie *et al.*, (2001); Paul *et al.* (2002); Yang, (2006). For reference bond distances, see: Allen *et al.* (1987).



Experimental

Crystal data

$C_{12}H_{10}N_2O_3$
M_r = 230.22
Monoclinic, $P2_1/n$
a = 4.9898 (5) Å
b = 20.662 (2) Å
c = 10.6994 (11) Å
β = 101.421 (2)°
V = 1081.24 (19) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.10 mm^{-1}
T = 295 (2) K
0.12 × 0.10 × 0.06 mm

Data collection

Bruker SMART APEXII area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
T_{min} = 0.98, T_{max} = 0.99
5631 measured reflections
1904 independent reflections
1451 reflections with $I > 2\sigma(I)$
R_{int} = 0.022

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.034
$wR(F^2)$ = 0.097
S = 1.02
1904 reflections
156 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.12 e Å^{-3}
$\Delta\rho_{min}$ = −0.13 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···O2^{i}	0.82	2.14	2.804 (2)	139
N1—H1A···O1	0.86	1.99	2.650 (2)	133

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

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); 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*.

This project was supported by the Postgraduate Foundation of Taishan University (No. Y07–2–16).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BG2220).

References

Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
Anderson, O. P., Cour, A. L., Findeisen, M., Hennig, L., Simonsen, O., Taylor, L. & Toflund, H. (1997). *J. Chem. Soc. Dalton Trans.* pp. 111–120.
Bruker (2005). *APEX2*, *SAINT* and *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
Garnovskii, A. D., Nivorozhkin, A. L. & Minki, V. I. (1993). *Coord. Chem. Rev.* **126**, 1–69.
Musie, G. T., Wei, M., Subramaniam, B. & Busch, D. H. (2001). *Inorg. Chem.* **40**, 3336–3341.
Paul, S., Barik, A. K., Peng, S. M. & Kar, S. K. (2002). *Inorg. Chem.* **41**, 5803–5809.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.
Yang, D.-S. (2006). *Acta Cryst.* E**62**, o1591–o1592.

supporting information

Acta Cryst. (2008). E**64**, o2208 [doi:10.1107/S1600536808034636]

N′-(Furan-2-ylmethylene)-2-hydroxybenzohydrazide

Yan-Xia Zhang

S1. Comment

Recently, a number of Schiff-bases have been investigated because of their coordination chemistry (Garnovskii *et al.*, 1993; Musie *et al.*, 2001; Paul *et al.*, 2002; Yang, 2006;) and biological systems (Anderson *et al.*, 1997). In order to search for new Schiff-bases with higher bioactivity, the title compound, (I), was synthesized and its crystal structure determined. In (I) (Fig. 1), the bond lengths and angles are in good agreement with the expected values (Allen *et al.*, 1987). In the crystal structure (Fig. 2), the molecules are linked into infinite chains by O—H···O hydrogen bonds. There is also an intramolecular N—H···O hydrogen bond.

S2. Experimental

The title compound was synthesized by the reaction of 2-hydroxy-benzoic acid hydrazide(1 mmol, 152.2 mg) with furan-2-carbaldehyde(1 mmol, 96.2 mg) in ethanol(20 ml) under reflux conditions (348 K) for 5 h. The solvent was removed and the solid product recrystallized from tetrahydrofuran. After six days orange crystals suitable for the X-ray diffraction study were obtained.

S3. Refinement

All H atoms were placed in idealized positions (C—H = 0.93— 0.97 Å, N—H = 0.86 Å) and refined as riding atoms. For those bound to C, U_{iso}(H) = 1.2 or 1.5U_{eq}(C). while for those bound to N, U_{iso}(H) = 1.2 U_{eq}(N).

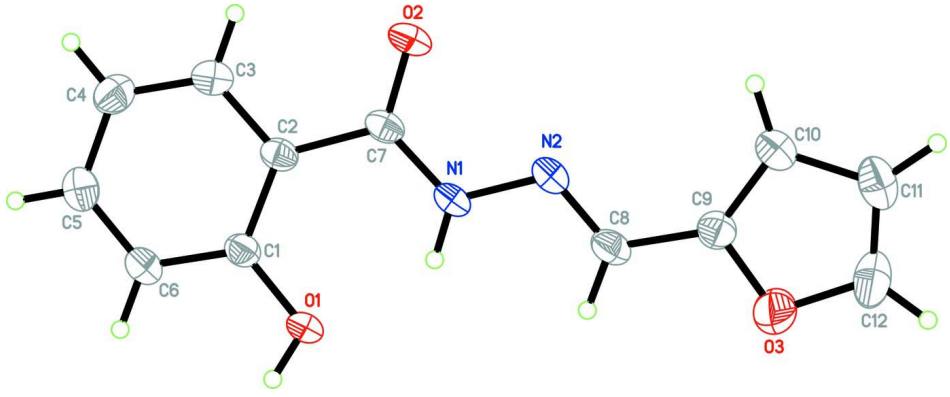


Figure 1

The molecular structure of (I), with displacement ellipsoids drawn at the 50% probability level.

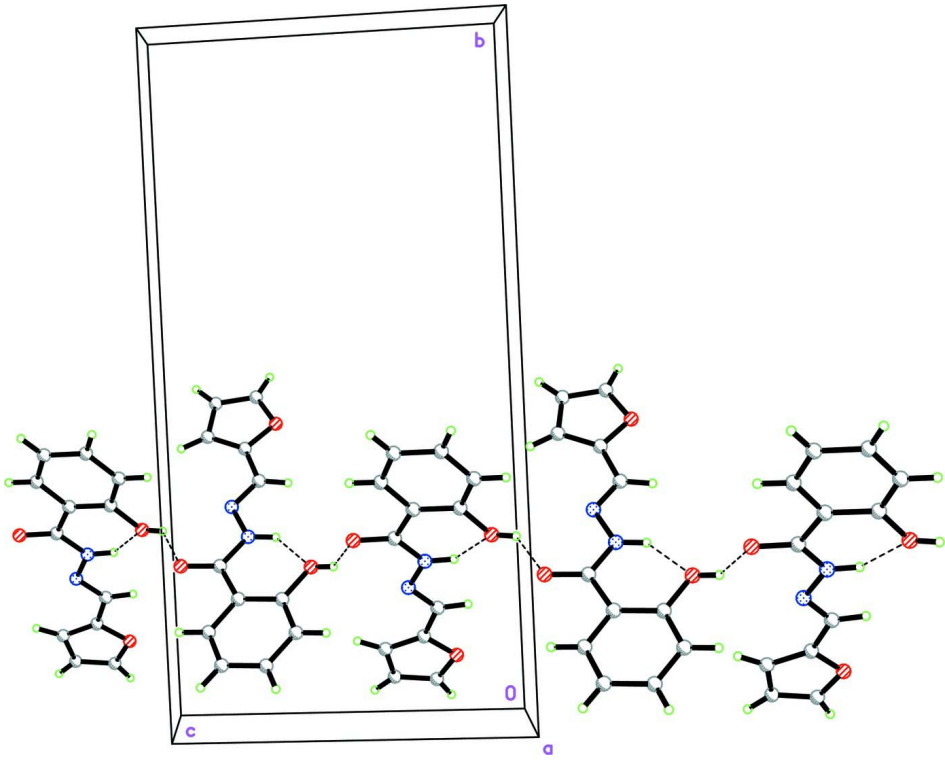


Figure 2

The structure of the infinite chains formed *via* hydrogen bonds, H atoms have been omitted for clarity. Dashed lines indicate hydrogen bonds.

N′-(Furan-2-ylmethylene)-2-hydroxybenzohydrazide

Crystal data

$C_{12}H_{10}N_2O_3$
$M_r = 230.22$
Monoclinic, $P2_1/n$
Hall symbol: -P 2yn
$a = 4.9898$ (5) Å
$b = 20.662$ (2) Å
$c = 10.6994$ (11) Å
$\beta = 101.421$ (2)°
$V = 1081.24$ (19) Å^3
$Z = 4$

$F(000) = 480$
$D_x = 1.414$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 1748 reflections
$\theta = 2.2$–$25.0°$
$\mu = 0.10$ mm^{-1}
$T = 295$ K
Block, orange
0.12 × 0.10 × 0.06 mm

Data collection

Bruker SMART APEXII area-detector diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
φ and ω scans
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
$T_{min} = 0.98$, $T_{max} = 0.99$

5631 measured reflections
1904 independent reflections
1451 reflections with $I > 2\sigma(I)$
$R_{int} = 0.023$
$\theta_{max} = 25.1°$, $\theta_{min} = 2.0°$
$h = -5 \rightarrow 5$
$k = -24 \rightarrow 21$
$l = -11 \rightarrow 12$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.034$
$wR(F^2) = 0.097$
$S = 1.02$
1904 reflections
156 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
$w = 1/[\sigma^2(F_o^2) + (0.0472P)^2 + 0.1418P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max} = 0.12$ e Å^{-3}
$\Delta\rho_{min} = -0.13$ e Å^{-3}
Extinction correction: *SHELXTL* (Sheldrick, 2008), $Fc^*=kFc[1+0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0107 (19)

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 (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
O1	0.2921 (2)	0.22463 (6)	0.59528 (10)	0.0611 (4)
H1	0.3615	0.2261	0.5320	0.092*
O2	0.2021 (2)	0.22907 (5)	0.97330 (9)	0.0541 (3)
O3	−0.6276 (3)	0.42331 (7)	0.68440 (12)	0.0815 (4)
N1	0.0370 (2)	0.26559 (6)	0.77544 (11)	0.0435 (3)
H1A	0.0365	0.2638	0.6951	0.052*
N2	−0.1342 (2)	0.30720 (6)	0.82213 (11)	0.0422 (3)
C1	0.4351 (3)	0.18305 (7)	0.68139 (14)	0.0415 (4)
C2	0.3961 (3)	0.18397 (7)	0.80733 (13)	0.0385 (3)
C3	0.5525 (3)	0.14165 (7)	0.89334 (15)	0.0493 (4)
H3	0.5312	0.1417	0.9778	0.059*
C4	0.7373 (3)	0.09972 (8)	0.85745 (16)	0.0560 (4)
H4	0.8409	0.0723	0.9171	0.067*
C5	0.7674 (3)	0.09876 (8)	0.73201 (16)	0.0549 (4)
H5	0.8892	0.0699	0.7065	0.066*
C6	0.6191 (3)	0.14000 (8)	0.64527 (15)	0.0508 (4)
H6	0.6417	0.1392	0.5611	0.061*
C7	0.2058 (3)	0.22765 (7)	0.85884 (13)	0.0397 (4)
C8	−0.2876 (3)	0.34231 (7)	0.74000 (15)	0.0483 (4)
H8	−0.2834	0.3378	0.6539	0.058*
C9	−0.4676 (3)	0.38898 (7)	0.77975 (14)	0.0460 (4)
C10	−0.5197 (3)	0.40836 (8)	0.89143 (17)	0.0582 (5)
H10	−0.4375	0.3928	0.9714	0.070*

C11	−0.7221 (4)	0.45682 (9)	0.8659 (2)	0.0668 (5)
H11	−0.7991	0.4793	0.9254	0.080*
C12	−0.7803 (4)	0.46390 (9)	0.7418 (2)	0.0767 (6)
H12	−0.9090	0.4928	0.6988	0.092*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0660 (8)	0.0871 (9)	0.0369 (6)	0.0281 (6)	0.0263 (5)	0.0161 (6)
O2	0.0647 (7)	0.0696 (8)	0.0325 (6)	0.0026 (6)	0.0206 (5)	0.0006 (5)
O3	0.0984 (10)	0.0848 (9)	0.0550 (8)	0.0339 (8)	0.0000 (7)	−0.0057 (7)
N1	0.0502 (7)	0.0512 (7)	0.0328 (7)	0.0022 (6)	0.0171 (6)	−0.0049 (6)
N2	0.0457 (7)	0.0457 (7)	0.0386 (7)	−0.0034 (6)	0.0167 (6)	−0.0075 (6)
C1	0.0397 (8)	0.0502 (9)	0.0365 (8)	−0.0013 (7)	0.0120 (6)	0.0019 (7)
C2	0.0386 (8)	0.0434 (8)	0.0350 (7)	−0.0083 (6)	0.0107 (6)	−0.0015 (6)
C3	0.0566 (9)	0.0529 (9)	0.0388 (8)	−0.0027 (8)	0.0103 (7)	0.0013 (7)
C4	0.0582 (10)	0.0533 (10)	0.0538 (10)	0.0060 (8)	0.0046 (8)	0.0067 (8)
C5	0.0525 (10)	0.0530 (10)	0.0612 (11)	0.0058 (8)	0.0162 (8)	−0.0053 (8)
C6	0.0512 (9)	0.0611 (10)	0.0439 (9)	0.0048 (8)	0.0185 (7)	−0.0016 (8)
C7	0.0427 (8)	0.0450 (8)	0.0343 (8)	−0.0099 (7)	0.0146 (6)	−0.0019 (7)
C8	0.0584 (10)	0.0526 (9)	0.0362 (8)	−0.0008 (8)	0.0147 (8)	−0.0076 (7)
C9	0.0475 (9)	0.0461 (9)	0.0439 (9)	−0.0030 (7)	0.0080 (7)	−0.0032 (7)
C10	0.0629 (11)	0.0656 (11)	0.0506 (10)	0.0079 (9)	0.0220 (8)	−0.0004 (8)
C11	0.0634 (12)	0.0592 (11)	0.0856 (15)	0.0003 (9)	0.0337 (11)	−0.0125 (10)
C12	0.0694 (13)	0.0596 (12)	0.0966 (17)	0.0202 (10)	0.0055 (12)	−0.0088 (11)

Geometric parameters (Å, º)

O1—C1	1.3549 (17)	C3—H3	0.9300
O1—H1	0.8200	C4—C5	1.380 (2)
O2—C7	1.2288 (16)	C4—H4	0.9300
O3—C12	1.359 (2)	C5—C6	1.365 (2)
O3—C9	1.3633 (19)	C5—H5	0.9300
N1—C7	1.3503 (18)	C6—H6	0.9300
N1—N2	1.3737 (16)	C8—C9	1.438 (2)
N1—H1A	0.8600	C8—H8	0.9300
N2—C8	1.2720 (19)	C9—C10	1.334 (2)
C1—C6	1.387 (2)	C10—C11	1.410 (2)
C1—C2	1.3991 (19)	C10—H10	0.9300
C2—C3	1.391 (2)	C11—C12	1.310 (3)
C2—C7	1.492 (2)	C11—H11	0.9300
C3—C4	1.374 (2)	C12—H12	0.9300
C1—O1—H1	109.5	C5—C6—C1	120.69 (15)
C12—O3—C9	106.28 (15)	C5—C6—H6	119.7
C7—N1—N2	118.33 (11)	C1—C6—H6	119.7
C7—N1—H1A	120.8	O2—C7—N1	120.95 (13)
N2—N1—H1A	120.8	O2—C7—C2	121.25 (13)

C8—N2—N1	115.98 (12)	N1—C7—C2	117.81 (12)
O1—C1—C6	120.38 (13)	N2—C8—C9	120.25 (13)
O1—C1—C2	119.43 (13)	N2—C8—H8	119.9
C6—C1—C2	120.19 (14)	C9—C8—H8	119.9
C3—C2—C1	117.51 (14)	C10—C9—O3	108.97 (14)
C3—C2—C7	116.76 (12)	C10—C9—C8	135.24 (16)
C1—C2—C7	125.71 (13)	O3—C9—C8	115.78 (14)
C4—C3—C2	122.05 (14)	C9—C10—C11	107.40 (17)
C4—C3—H3	119.0	C9—C10—H10	126.3
C2—C3—H3	119.0	C11—C10—H10	126.3
C3—C4—C5	119.29 (15)	C12—C11—C10	106.37 (17)
C3—C4—H4	120.4	C12—C11—H11	126.8
C5—C4—H4	120.4	C10—C11—H11	126.8
C6—C5—C4	120.24 (15)	C11—C12—O3	110.96 (17)
C6—C5—H5	119.9	C11—C12—H12	124.5
C4—C5—H5	119.9	O3—C12—H12	124.5
C7—N1—N2—C8	179.41 (13)	C3—C2—C7—O2	4.9 (2)
O1—C1—C2—C3	−178.33 (13)	C1—C2—C7—O2	−173.54 (14)
C6—C1—C2—C3	1.5 (2)	C3—C2—C7—N1	−175.00 (13)
O1—C1—C2—C7	0.1 (2)	C1—C2—C7—N1	6.6 (2)
C6—C1—C2—C7	179.90 (13)	N1—N2—C8—C9	−177.93 (12)
C1—C2—C3—C4	−0.6 (2)	C12—O3—C9—C10	−0.45 (19)
C7—C2—C3—C4	−179.16 (13)	C12—O3—C9—C8	179.81 (14)
C2—C3—C4—C5	−0.8 (2)	N2—C8—C9—C10	1.6 (3)
C3—C4—C5—C6	1.4 (3)	N2—C8—C9—O3	−178.74 (14)
C4—C5—C6—C1	−0.5 (2)	O3—C9—C10—C11	0.28 (19)
O1—C1—C6—C5	178.84 (14)	C8—C9—C10—C11	179.95 (17)
C2—C1—C6—C5	−1.0 (2)	C9—C10—C11—C12	0.0 (2)
N2—N1—C7—O2	1.8 (2)	C10—C11—C12—O3	−0.3 (2)
N2—N1—C7—C2	−178.36 (11)	C9—O3—C12—C11	0.5 (2)

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
O1—H1···O2^{i}	0.82	2.14	2.804 (2)	139
N1—H1*A*···O1	0.86	1.99	2.650 (2)	133

Symmetry code: (i) *x*+1/2, −*y*+1/2, *z*−1/2.