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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.004 Å R factor = 0.054 wR factor = 0.148 Data-to-parameter ratio = 12.6

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

The title compound, $C_{14}H_{11}FN_2O_2$, crystallizes with two molecules in the asymmetric unit. Both molecules are roughly planar and adopts a *trans* configuration with respect to the C=N double bond. In the crystal structure, intermolecular $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonds result in the formation of sheets parallel to the *ac* plane.

4-Fluorobenzaldehyde salicylhydrazone

Comment

Schiff base compounds have been of great interest for many years. These compounds play an important role in the development of coordination chemistry related to catalysis and enzymatic reactions (Brunner *et al.*, 1983), magnetism and molecular architectures (Miller & Epstein, 2000). The deprotonated hydroxyl group of the salicylhydrazone will sometimes cause the N atom of the hydrazone to become more electron-rich as a result of conjugation effects, and the N atom can thus coordinate more strongly (Bansse *et al.*, 1998). As an extension of work on the structural characterization of salicylhydrazone Schiff base compounds (Ma *et al.*, 2005), we report here the crystal structure of (I), a new salicylhydrazone with 4-fluorobenzaldehyde.



In the title compound, (I), which crystallizes with two molecules in the asymmetric unit (Fig. 1), C–N bonds in the hydrazone units are rather short (Table 1) owing to conjugation effects (Ma *et al.*, 2005). All other bond lengths are within normal ranges (Cambridge Structural Database, Version 5.26; Allen, 2002). The dihedral angles between the two benzene rings are 7.3 (2)° and 18.7 (1)° in the two molecules.

The occurrence of $O-H \cdots O$ hydrogen bonds results in the formation of infinite chains which are linked by $N-H \cdots O$ hydrogen bonds, forming sheets parallel to the *ac* plane (Table 2 and Fig. 2).

Experimental

The reagents were commercial products and were used without further purification. 4-Fluorobenzaldehyde (0.2 mmol, 28 mg) and 2-hydroxybenzhydrazide (0.2 mmol, 30.4 mg) were dissolved in methanol (10 ml). The mixture was stirred at room temperature for

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Figure 1

The structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme. Hydrogen bonds are represented by dashed lines.

about 10 min to give a clear yellow solution. The solution was set aside for 8 d to allow slow evaporation of the solvent. Large colourless prism-shaped crystals separated; these were collected and washed three times with water.

Crystal data

$C_{14}H_{11}FN_2O_2$	$D_x = 1.383 \text{ Mg m}^{-3}$
$M_r = 258.25$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 3063
a = 4.9532 (4) Å	reflections
b = 47.867 (4) Å	$ heta=4.7 extrm{-}41.4^\circ$
c = 10.7392 (9) Å	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 102.953 \ (2)^{\circ}$	T = 298 (2) K
V = 2481.4 (4) Å ³	Prism, colourless
Z = 8	0.35 \times 0.18 \times 0.12 mm
Data collection	
Bruker SMART APEX area-	4358 independent reflections
detector diffractometer	2137 reflections with $I > 2\sigma(I)$

 $R_{\rm int}=0.078$

 $\theta_{\rm max} = 25.0^{\circ}$

 $h = -5 \rightarrow 5$

 $k=-56\rightarrow 56$

 $l = -12 \rightarrow 12$

bruter SMART AT EX area detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2002) $T_{min} = 0.964, T_{max} = 0.988$ 23870 measured reflections

Refinement

Refinement on F^2	H-atom parameters constrained		
$R[F^2 > 2\sigma(F^2)] = 0.054$	$w = 1/[\sigma^2(F_o^2) + (0.086P)^2]$		
$wR(F^2) = 0.148$	where $P = (F_0^2 + 2F_c^2)/3$		
S = 0.87	$(\Delta/\sigma)_{\rm max} < 0.001$		
4358 reflections	$\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$		
345 parameters	$\Delta \rho_{\rm min} = -0.22 \ {\rm e} \ {\rm \AA}^{-3}$		

Table 1

Selected bond lengths (Å).

N1-C7	1.338 (3)	N3-C21	1.340 (3)
N1-N2	1.370 (3)	N3-N4	1.383 (3)
N2-C8	1.263 (3)	N4-C22	1.273 (3)



Figure 2

The crystal packing of (I), showing the formation of sheets. $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonds are represented by dashed lines. H atoms not involved in hydrogen bonding have been omitted for clarity.

Table 2Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O1−H1···O4	0.82	1.86	2.662 (2)	165
$O3 - H3 \cdot \cdot \cdot O2^{i}$	0.82	1.86	2.643 (2)	159
$N1 - H1A \cdots O1$	0.86	2.00	2.653 (3)	132
$N1 - H1A \cdots O4^{ii}$	0.86	2.53	3.109 (3)	126
N3−H3A···O3	0.86	1.94	2.638 (3)	137

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

All H atoms were placed in geometrically idealized positions (N–H 0.86, O–H 0.82 and C–H 0.93 Å) and treated as riding on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(C,O,N)$.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPIII* (Burnett & Johnson, 1996), *ORTEP-3 for Windows* (Farrugia, 1997) and *CAMERON* (Watkin *et al.*, 1993); software used to prepare material for publication: *SHELXL97*.

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