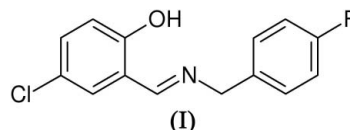


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Key indicators

Single-crystal X-ray study
 $T = 292$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.058
 wR factor = 0.171
Data-to-parameter ratio = 17.6For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.4-Chloro-2-[(*E*)-(4-fluorophenyl)methyl-
iminomethyl]phenolThe title compound, $\text{C}_{14}\text{H}_{11}\text{ClFNO}$, forms an intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond. The dihedral angle between the two benzene rings is $71.6(2)^\circ$.Received 4 May 2006
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Comment

Salicylaldehyde and its derivatives play an important role in developing molecular architectures in coordination chemistry (Mukhopadhyay *et al.*, 2003; Erxleben & Schumacher, 2001). As Lewis base ligands, the adducts of aldehydes with various primary amines are of interest in a large number of transition-metal complexes (Sreenivasulu *et al.*, 2005; Ranford *et al.*, 1999). Recently, we have reported the structural characterization of two Schiff base compounds derived from the condensation of salicylaldehyde and primary amines (Li *et al.*, 2005, 2006). As an extension of this work, we report here the crystal structure of the title compound, (I).In (I), all bond lengths are within normal ranges (Allen *et al.*, 1987) (Fig. 1). The $\text{C}7=\text{N}1$ bond length of $1.264(3)$ Å is shorter than the corresponding value of $1.283(3)$ Å observed in a similar Schiff base compound (Li *et al.*, 2006). The $\text{C}7-\text{N}1-\text{C}8-\text{C}9$ and $\text{N}1-\text{C}8-\text{C}9-\text{C}10$ torsion angles are $-124.1(2)$ and $98.2(3)^\circ$, respectively. The dihedral angle between the two benzene rings is $71.6(2)^\circ$. An intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond is formed between atoms O1 and N1 (Fig. 1 and Table 1).

Experimental

4-Fluorobenzylamine (125 mg, 1 mmol) and 5-chlorosalicylaldehyde (160 mg, 2 mmol) were dissolved in methanol (10 ml) at 323 K. The mixture was stirred for 10 min to give a clear yellow solution. After keeping the solution in air for 8 d, yellow block crystals were formed at the bottom of the vessel, in about 59% yield, on slow evaporation of the solvent. Elemental analysis found: C 63.76, H 4.23, N 5.45%; calculated: C 63.77, H 4.20, N 5.31%.

Crystal data

 $\text{C}_{14}\text{H}_{11}\text{ClFNO}$
 $M_r = 263.69$
Monoclinic, $P2_1/c$
 $a = 15.0408(17)$ Å
 $b = 6.0375(7)$ Å
 $c = 14.3943(17)$ Å
 $\beta = 105.068(2)^\circ$
 $V = 1262.2(3)$ Å³ $Z = 4$
 $D_x = 1.388$ Mg m⁻³
Mo $K\alpha$ radiation
 $\mu = 0.30$ mm⁻¹
 $T = 292(2)$ K
Block, yellow
 $0.36 \times 0.20 \times 0.20$ mm

Data collection

Bruker SMART APEX CCD
diffractometer
 φ and ω scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
 $T_{\min} = 0.930$, $T_{\max} = 0.942$

10448 measured reflections
2875 independent reflections
1922 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.075$
 $\theta_{\text{max}} = 27.5^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.058$
 $wR(F^2) = 0.171$
 $S = 1.06$
2875 reflections
163 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0827P)^2 + 0.0486P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.21 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.23 \text{ e } \text{\AA}^{-3}$

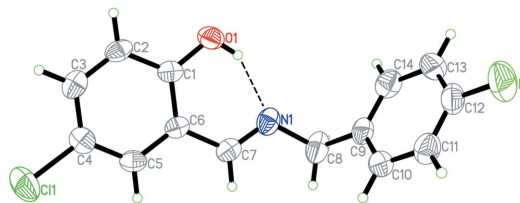


Figure 1

Molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are shown at the 30% probability level. H atoms are shown as spheres of arbitrary radii. The dashed line denotes the intramolecular O—H...N hydrogen bond.

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1—H1...N1	0.82	1.88	2.601 (2)	147

All H atoms were positioned geometrically and constrained to ride on their parent atoms, with C—H = 0.93 and 0.97 \AA for sp^2 and sp^3 C atoms, respectively, O—H = 0.82 \AA , and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{O})$. Atom H1 of the hydroxy group was placed so as to form the best hydrogen bond to N1.

Data collection: SMART (Bruker, 2003); cell refinement: SAINT (Bruker, 2003); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXL97.

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