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Kev indicators

Single-crystal X-ray study T = 160 K Mean σ (C–C) = 0.003 Å R factor = 0.043 wR factor = 0.112 Data-to-parameter ratio = 10.8

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

N-(4-Butylphenyl)salicylaldimine at 160 K

In the title compound, C₁₇H₁₉NO, which is a Schiff base, an intramolecular O-H···N hydrogen bond involving the hydroxy group forms a six-membered ring with a graph-set motif of S(6). Several intermolecular C-H··· π interactions occur between adjacent molecules which are arranged perpendicular to one another.

Comment

Schiff base compounds can be classified by their photochromic and thermochromic characteristics (Cohen et al., 1964; Hadjoudis et al., 1987). It is known that molecules exhibiting photochromism are non-planar (Moustakali-Mavridis et al., 1978). In view of this, an X-ray crystal structure determination of the title compound, (I), has been carried out to investigate the planarity of the molecule in the solid state.



Fig. 1 shows the asymmetric unit of (I) with the atomic numbering scheme. The corresponding bond lengths and angles in (I) are comparable with those in the related structures, N-salicylidene-2-aminopyridine, 3-methyl-N-(3-methoxysalicylidene)-2-aminopyridine, N-(5-bromosalicylidene)-2aminopyridine and 5-chloro-N-(5-methoxysalicylidene)-2aminopyridine (Moustakali-Mavridis et al., 1978), N-(5chlorosalicylidene)-2-hydroxy-5-chloroaniline (Elerman et al., 2002) and N-(2-fluoro-3-methoxy)salicylidene (Ünver et al., 2002). The butyl group projects in a planar zigzag fashion directly away from and perpendicular to the central aromatic ring plane $[C13-C12-C15-C16 = 88.4 (2)^{\circ}]$. The dihedral angle between the planes of the two aromatic rings is $11.46 (10)^{\circ}$, which shows that the conjugated part of the molecule is not entirely planar.



Figure 1

View of the asymmetric unit of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary radii.

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In the crystal structure, the hydroxy group forms an intramolecular hydrogen bond with the N atom, thereby completing a six-membered ring which has a graph-set motif of S(6) (Bernstein et al., 1995). Atom C11 (via H11) in the butyl-substituted phenyl ring acts as a donor for a weak intermolecular $C-H\cdots\pi$ interaction with the hydroxysubstituted phenyl ring (centroid Cg1) in the molecule at $(2 - x, \frac{1}{2} + y, \frac{3}{2} - z)$. Atom C14 (*via* H14), which is *para* to C11, is involved in a weak intermolecular $C-H\cdots\pi$ interaction with the hydroxy-substituted phenyl ring of another molecule lying in the opposite direction at $(1 - x, y - \frac{1}{2}, \frac{3}{2} - z)$. Atoms C3 (via H3) and C6 (via H6), which lie para with respect to each other in the hydroxy-substituted phenyl ring, are also involved in weak intermolecular $C-H\cdots\pi$ interactions with the butyl-substituted phenyl ring (centroid Cg2) in the molecules at $(1 - x, y - \frac{1}{2}, \frac{3}{2} - z)$ and $(2 - x, \frac{1}{2} + y, \frac{3}{2} - z)$, respectively (Table 1). These interactions result from the adjacent molecules being stacked perpendicular to one another along the [110] direction.

Experimental

The title Schiff base was synthesized by refluxing absolute ethanol solutions of salicylaldimine (15 mmol) and 4-butylaniline (15 mmol) for 2 h and leaving the solution overnight. The product was filtered off by suction, thoroughly washed with cold ethanol and recrystallized from absolute ethanol (yield 70%; m.p. 311–338 K).

Crystal data

C ₁₇ H ₁₉ NO	Mo $K\alpha$ radiation
$M_r = 253.33$	Cell parameters from 1931
Orthorhombic, $P2_12_12_1$	reflections
a = 5.5848 (1) Å	$\theta = 2.0-27.5^{\circ}$
b = 8.0987 (2) Å	$\mu = 0.07 \text{ mm}^{-1}$
c = 31.2943 (6) Å	T = 160 (2) K
V = 1415.43 (5) Å ³	Prism, colourless
Z = 4	$0.25 \times 0.23 \times 0.18 \text{ mm}$
$D_x = 1.189 \text{ Mg m}^{-3}$	

Data collection

Nonius KappaCCD diffractometer φ and ω scans with κ offsets 18 762 measured reflections 1917 independent reflections 1474 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.112$ S = 1.031917 reflections 177 parameters

H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0691P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} = 0.001$

 $\Delta \rho_{\text{max}} = 0.16 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\text{min}} = -0.13 \text{ e } \text{\AA}^{-3}$

 $R_{\rm int}=0.058$

 $\theta_{\rm max} = 27.5^\circ$

 $h = -7 \rightarrow 7$

 $k = -8 \rightarrow 10$

 $l = -40 \rightarrow 40$

Table 1

Hydrogen-bonding geometry (Å, °).

Cg1 is the centroid of ring C1-C6 and Cg2 is the centroid of ring C9-C14.

D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
0.90(2)	1.81 (2)	2.598 (2)	144 (2)
0.95	3.10	3.7998 (19)	132
0.95	2.97	3.7003 (18)	135
0.95	2.98	3.7585 (19)	140
0.95	2.81	3.5976 (19)	141
	<i>D</i> -H 0.90 (2) 0.95 0.95 0.95 0.95	$\begin{array}{c c} D-H & H\cdots A \\ \hline 0.90 (2) & 1.81 (2) \\ 0.95 & 3.10 \\ 0.95 & 2.97 \\ 0.95 & 2.98 \\ 0.95 & 2.81 \end{array}$	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$

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

The position of the hydroxy H atom was determined from a difference Fourier map and refined along with its isotropic displacement parameter, while restraining the O-H bond length to 0.82 (2) Å. The methyl H atoms were constrained to an ideal geometry (C-H = 0.98 Å), with $U_{iso}(H) = 1.5U_{eq}(C)$, but were allowed to rotate freely about the C-C bonds. All the remaining H atoms were placed in geometrically idealized positions (C-H = 0.95–0.99 Å) and constrained to ride on their parent atoms with $U_{iso} = 1.2U_{eq}(C)$. Although the molecule is achiral, the crystal structure is chiral. Because of the absence of any significant anomalous scatterers in the compound, attempts to refine the Flack (1983) parameter with 1320 Friedel pairs led to an inconclusive value (Flack & Bernardinelli, 2000) of -1.2 (16). Therefore, the absolute structure was assigned arbitrarily and the Friedel pairs were merged before the final refinement.

Data collection: *COLLECT* (Nonius, 2000); cell refinement: *DENZO–SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO–SMN* and *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL*97 and *PLATON* (Spek, 2003).

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