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Key indicators

Single-crystal X-ray study T = 273 K Mean σ (C–C) = 0.002 Å R factor = 0.038 wR factor = 0.114 Data-to-parameter ratio = 13.3

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

2-Hydroxy-*N*'-[(1*E*)-1-(2-hydroxy-5-methyl-phenyl)-2-phenylethylidene]benzohydrazide

The title compound, $C_{22}H_{20}N_2O_3$, displays a *trans* configuration with respect to the C=N double bond. The crystal structure is stabilized by intramolecular $O-H\cdots N$ and $N-H\cdots O$ and intermolecular $O-H\cdots O$ hydrogen bonds.

Comment

The chemistry of arylhydrazones continues to attract much attention due to their ability to coordinate to metal ions (Singh *et al.*, 1982; Salem, 1998) and their biological activity (Singh *et al.*, 1982; Carcelli *et al.*, 1995). As an extension of our work on the structural characterization of arylhydrazone derivatives, the title compound, (I), was synthesized.



The title molecule displays a *trans* configuration with respect to the C7=N1 double bond (Fig. 1). The three benzene rings, C1-C6 (*A*), C17-C22 (*B*) and C9-C14 (*C*) make dihedral angles of 19.02 (9) (*A*/*B*), 79.17 (5) (*A*/*C*) and 88.39 (5)° (*B*/*C*). Intramolecular N-H···O and O-H···N hydrogen bonds stabilize the conformation of the molecule, while O-H···O intermolecular hydrogen bonds lead to the formation of an infinite chain, graph set C(6) (Etter *et al.*, 1990), extending along the *a* axis (Table 1 and Fig. 2).

Experimental

2-Hydroxybenzoylhydrazine (0.02 mol, 3.04 g) was dissolved in anhydrous ethanol (50 ml), and 1-(2-hydroxy-5-methylphenyl)-2-phenylethanone (0.02 mol, 4.52 g) was added. The reaction mixture was refluxed for 6 h with stirring. The resulting precipitate was collected by filtration, washed several times with ethanol and dried *in vacuo* (yield 81%). The compound (2.0 mmol, 0.72 g) was dissolved in dimethylformamide (30 ml) and allowed to stand at room temperature for 30 d, yielding yellow single crystals suitable for X-ray diffraction.

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organic papers

Crystal data

 $\begin{array}{l} C_{22}H_{20}N_2O_3\\ M_r = 360.40\\ \text{Triclinic, }P\overline{1}\\ a = 6.7401 \ (11) \ \mathring{A}\\ b = 10.4033 \ (17) \ \mathring{A}\\ c = 14.065 \ (2) \ \mathring{A}\\ \alpha = 79.475 \ (6)^\circ\\ \beta = 80.193 \ (6)^\circ\\ \gamma = 78.639 \ (6)^\circ\end{array}$

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: none 10775 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0541P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.038$	+ 0.1919P]
$wR(F^2) = 0.114$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.03	$(\Delta/\sigma)_{\rm max} = 0.005$
3292 reflections	$\Delta \rho_{\rm max} = 0.26 \text{ e} \text{ Å}^{-3}$
247 parameters	$\Delta \rho_{\rm min} = -0.13 \text{ e} \text{ Å}^{-3}$
H-atom parameters constrained	

V = 941.4 (3) Å³

 $D_x = 1.271 \text{ Mg m}^{-3}$

 $0.35 \times 0.24 \times 0.13 \text{ mm}$

3292 independent reflections

2663 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

 $\mu = 0.09 \text{ mm}^{-1}$

T = 273 (2) K

Block, yellow

 $R_{\rm int} = 0.020$

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

Z = 2

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{matrix} O3-H3\cdots N2\\ N1-H1\cdots O1\\ O1-H1A\cdots O2^i \end{matrix}$	0.82	1.83	2.5468 (15)	145
	0.86	1.96	2.6338 (15)	134
	0.82	1.88	2.6850 (15)	169

Symmetry code: (i) x + 1, y, z.

All H atoms were positioned geometrically and treated as riding on their parent atoms, with methyl C–H = 0.96 Å, aromatic C–H = 0.93 Å, O–H = 0.82 Å and N–H = 0.86 Å, and with $U_{iso}(H) =$ 1.5 U_{eq} (methyl C,O) and 1.2 U_{eq} (aromatic C,N).

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXTL*.

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Figure 1

The molecular structure of compound (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented as small spheres of arbitrary radii.



Figure 2

Partial packing view showing $O-H \cdots N$, $N-H \cdots O$ and $O-H \cdots O$ hydrogen bonds and the formation of an infinite chain. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted for clarity. [Symmetry code: (i) 1 + x, y, z.]

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