

4-(4-Hydroxybenzylideneamino)-4H-1,2,4-triazole hemihydrate

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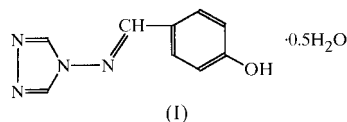
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In the title compound, 4-(4H-1,2,4-triazol-4-yliminomethyl)-phenol hemihydrate, $C_9H_8N_4O \cdot 0.5H_2O$ or (I)·0.5H₂O, molecules of (I) are arranged as layers running along the *b* axis through intermolecular O—H···N and C—H···O hydrogen bonds. These layers are stabilized by hydrogen-bonded water molecules to form three-dimensional networks.

Comment

The aroyl Schiff bases of 4-amino-1,2,4-triazole have received considerable attention over the past few decades (Kitaev *et al.*, 1971; Mazza *et al.*, 1976; Kargin *et al.*, 1988). It is of interest that some of them are anti-inflammatory agents (Gupta & Bhargava, 1978) and new coccidiostatic drugs (Colautti *et al.*, 1971). As a continuation of our study on the structure of Schiff-base-containing substituted 1,2,4-triazole (Shanmuga Sundara Raj *et al.*, 1999), we report here the title structure, (I).



An ORTEPII (Johnson, 1976) diagram of (I) with numbering scheme is shown in Fig. 1. Compared to 9-(4H-1,2,4-triazol-4-ylimino)-4,5-diazafluorene (Shanmuga Sundara Raj *et al.*, 1999), molecules of (I) are essentially planar [the maximum displacement from the least-squares mean plane through the whole molecule is 0.073 (2) Å for C9]. The bond lengths and angles observed in the structure are in the normal ranges. In contrast with 2-(2-hydroxybenzylidene)-1-(2-picoloyl)hydrazine hemihydrate (Wang *et al.*, 1998), molecules of (I) are arranged as layers running along the *b* axis through strong intermolecular O1—H11···N4($-\frac{1}{2} + x, -\frac{1}{2} - y, \frac{1}{2} + z$) hydrogen bonds and weak C9—H9···O1($\frac{1}{2} - x, -\frac{1}{2} + y, -\frac{1}{2} - z$) ones. These layers are stabilized by water molecules to form three-dimensional networks through strong O1W—H1W···N3 hydrogen bonds and weak C7—H7···O1W($x, -y, \frac{1}{2} + z$) ones. The geometry of these interactions are listed in

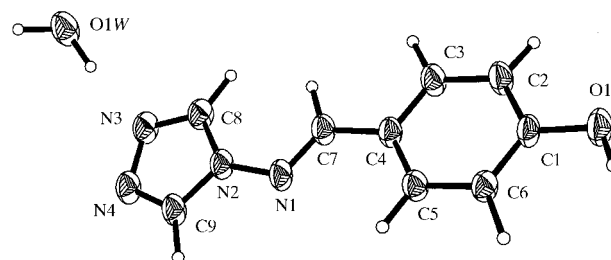


Figure 1
The structure of (I) showing 50% probability displacement ellipsoids with the numbering scheme.

Table 2. In the packing of molecules of the title compound, the water molecules (O1W) lie on crystallographic twofold axes.

Experimental

The title compound was prepared by condensation of equivalent amounts of *p*-hydroxybenzaldehyde and 4-amino-1,2,4-triazole in ethanol for 5 h (Kitaev *et al.*, 1971). Diffraction-quality crystals were obtained by recrystallization from ethanol.

Crystal data

$C_9H_8N_4O \cdot 0.5H_2O$
 $M_r = 197.20$
Monoclinic, C_2/c
 $a = 14.134$ (2) Å
 $b = 12.491$ (2) Å
 $c = 12.063$ (2) Å
 $\beta = 119.483$ (10)°
 $V = 1854.0$ (5) Å³
 $Z = 8$

$D_x = 1.413$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 36 reflections
 $\theta = 5.22$ – 9.01 °
 $\mu = 0.102$ mm⁻¹
 $T = 293$ (2) K
Block, colourless
0.30 × 0.26 × 0.20 mm

Data collection

Siemens P4 diffractometer
 $2\theta/\omega$ scans
Absorption correction: empirical (North *et al.*, 1968)
 $T_{\min} = 0.954$, $T_{\max} = 0.968$
2043 measured reflections
1642 independent reflections
1100 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.021$
 $\theta_{\text{max}} = 25$ °
 $h = -1 \rightarrow 16$
 $k = -1 \rightarrow 14$
 $l = -14 \rightarrow 12$
3 standard reflections every 97 reflections
intensity decay: 5.84%

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.057$
 $wR(F^2) = 0.155$
 $S = 1.166$
1637 reflections
133 parameters
H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0851P)^2 + 1.2289P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = -0.006$
 $\Delta\rho_{\text{max}} = 0.28$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³
Extinction correction: SHELXTL (Siemens, 1995)
Extinction coefficient: 0.0022 (6)

Table 1
Selected geometric parameters (Å, °).

N1—C7	1.267 (2)	N3—N4	1.383 (2)
N1—N2	1.409 (2)	N4—C9	1.302 (2)
N2—C9	1.357 (2)	C1—O1	1.350 (2)
N2—C8	1.362 (2)	C4—C7	1.456 (3)
N3—C8	1.304 (3)		
C7—N1—N2	115.4 (2)	C8—N2—N1	132.3 (2)
C9—N2—C8	105.7 (2)	O1—C1—C6	123.0 (2)
C9—N2—N1	122.0 (2)	N1—C7—C4	122.5 (2)

Table 2

Hydrogen-bonding geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O1—H11...N4 ⁱ	1.01	1.74	2.717 (2)	161
C9—H9...O1 ⁱⁱ	1.18	2.15	3.293 (3)	164
O1W—H1W...N3	1.02	1.89	2.898 (2)	169
C7—H7...O1W ⁱⁱⁱ	1.10	2.47	3.214 (2)	123

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

Data collection: *XSCANS* (Siemens, 1994); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Siemens, 1995); program(s) used to solve structure: *SHELXTL*; program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: NA1438). Services for accessing these data are described at the back of the journal.

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