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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.058 wR factor = 0.189 Data-to-parameter ratio = 17.2

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

3-[(2-Hydroxyphenyl)amino]-1-phenylbut-3-en-1-one ethanol solvate

The title compound, $C_{16}H_{15}NO_2 \cdot C_2H_5OH$, has been synthesized by the reaction of benzoylacetone with 2-aminophenol in ethanol. The enaminone structure is stabilized by a strong intramolecular hydrogen bond (N-H···O=C), while the one-dimensional infinite chain in the [010] direction is formed by the intermolecular O-H···O hydrogen bond between the enaminone and the ethanol solvent molecule.

Comment

Enaminones have been studied extensively, not only because of their very diverse reaction chemistry, but also owing to their applications in coordination chemistry (Kim *et al.*, 2001; Doherty *et al.*, 1999). As part of a systematic investigation of the chemistry of enaminones, the enaminone component of the title compound, (I), was obtained (Shi, Yang, Shen *et al.*, 2004; Shi, Yang, Song *et al.*, 2004) and its crystal structure is reported here.



The parent unsolvated enaminone has already been studied (Glowiak & Sobczak, 1992). It crystallizes as two polymorphs, the orange triclinic form with two molecules, (Ia) and (Ib), in the asymmetric unit, and the colorless monoclinic form, (Ic). Although the corresponding bond distances and angles in (I) are very similar to those in (Ia), (Ib) and (Ic), the most striking



Figure 1

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved The asymmetric unit of the title compound, showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 30% probability level. The intramolecular hydrogen bond is shown as a dashed line.

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 $D_x = 1.172 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 874 reflections $\theta = 2.8-24.6^{\circ}$ $\mu = 0.08 \text{ mm}^{-1}$ T = 293 (2) K Block, yellow $0.24 \times 0.22 \times 0.16 \text{ mm}$

3484 independent reflections 1948 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.033$

 $\begin{array}{l} \theta_{\rm max} = 26.5^{\circ} \\ h = -9 \rightarrow 6 \end{array}$

 $k = -11 \rightarrow 15$

 $l = -21 \rightarrow 23$





Packing diagram of the title compound. Dashed lines indicate hydrogen bonds. H atoms not involved in hydrogen bonding have been omitted.

difference between them lies in the dihedral angles between the O=C-C=C-N plane and the benzene and substituted benzene rings. The dihedral angles are 18.8 and 41.0° for (I*a*), 14.3 and 56.8° for (I*b*) and 28.1 and 85.5° for (I*c*); in (I), the values are 21.96 (13) and 46.68 (13)°. For each of (I), (I*a*), (I*b*) and (I*c*), the bond lengths in the O=C-C=C-N system indicate electron delocalization (Shi, Yang, Shen *et al.*, 2004; Shi, Yang, Song *et al.*, 2004; Gilli *et al.*, 2000; Arici *et al.*, 1999). Interestingly, the above dihedral angles suggest that neither the benzene nor the substituted benzene rings are involved in the conjugation of the O=C-C=C-N system.

As in (Ia), (Ib) and (Ic), the enamine N atom and carbonyl O atom in (I) form a strong intramolecular hydrogen bond which stabilizes the enaminone (Table 2). Strong intermolecular $O-H\cdots O$ hydrogen bonds are also present in the crystal structure. These lead to infinite one-dimensional chains in the [010] direction.

Experimental

The title compound was synthesized by refluxing a solution of benzoylacetone and 2-aminophenol (1:1) in ethanol for 6 h. It was recrystallized from ethanol as yellow crystals which are unstable and easily lose ethanol to form the yellow enaminone [90% yield; m.p. 443–444 K; literature m.p. 437–438 K (colorless crystals), 437–439 K (orange crystals) (Glowiak & Sobczak, 1992)]. IR (KBr, cm⁻¹): 3159.61 (*br, s*, NH, OH), 1604.55 (*vs*, C=O), 1574.13 (*s*, C=C). UV (λ_{max} , in DMF): 256.00 (B band, 1.36 × 10⁴), 359.5 (K band, 3.52 × 10⁴) nm. ¹H NMR (300 MHz, CDCl₃): δ 12.35 (1H, *s*, NH), 8.61 (1H, *s*, *br*, OH), 7.84–7.87, 7.41–7.43 (2H, *m*, 3H, *m*, C₆H₅), 6.82–7.03 (4H, 3*m*, C₆H₄), 5.66 (1H, *s*, CH), 1.61 (3H, *s*, CH₃).

Crystal data

$C_{16}H_{15}NO_2 \cdot C_2H_6O$
$M_r = 299.36$
Monoclinic, $P2_1/n$
$a = 7.663 (4) \text{ Å}_{1}$
b = 12.119(7) Å
c = 18.303 (9) Å
$\beta = 93.217 \ (14)^{\circ}$
$V = 1697.1 (16) \text{ Å}^3$
Z = 4

Data collection

Bruker SMART 1000 CCD areadetector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996; Blessing, 1995) $T_{\min} = 0.967, T_{\max} = 0.987$ 9539 measured reflections

Refinement

 $\begin{array}{ll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.098P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.058 & w + 0.1387P] \\ wR(F^2) = 0.189 & where \ P = (F_o^2 + 2F_c^2)/3 \\ S = 1.03 & (\Delta/\sigma)_{\rm max} < 0.001 \\ 3484 \ \mbox{reflections} & \Delta\rho_{\rm max} = 0.36 \ \mbox{e} \ {\rm \AA}^{-3} \\ 203 \ \mbox{parameters} & \Delta\rho_{\rm min} = -0.31 \ \mbox{e} \ {\rm \AA}^{-3} \\ \mbox{H-atom parameters constrained} \\ \end{array}$

Table 1

Selected geometric parameters (Å, °).

O1-C4 O2-C12 N1-C2 N1-C11	1.264 (3) 1.370 (3) 1.339 (3) 1.414 (3)	C2-C3 C3-C4 C4-C5	1.385 (4) 1.420 (3) 1.496 (3)
C2 - N1 - C11N1 - C2 - C3N1 - C2 - C1C3 - C2 - C1	130.3 (2) 120.6 (2) 120.1 (2) 119.4 (2)	C2-C3-C4 O1-C4-C3 O1-C4-C5 C3-C4-C5	124.1 (2) 121.9 (2) 118.5 (2) 119.6 (2)

able 2		
Iydrogen-bonding geometry	(Å,	°).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N1-H1···O1	0.86	1.92	2.628 (3)	139
O2−H2···O3	0.82	1.84	2.657 (3)	171
$O3-H3\cdots O1^i$	0.82	1.86	2.685 (3)	178

Symmetry code: (i) $\frac{3}{2} - x$, $y - \frac{1}{2}, \frac{1}{2} - z$.

All H atoms were located in difference maps and were subsequently treated as riding atoms, with C–H = 0.93 (olefinic and aromatic), 0.96 (methyl) or 0.97 Å (methylene), N–H = 0.86 Å and O–H = 0.82 Å. $U_{\rm iso}$ (H) values were set at $1.2U_{\rm eq}$ (C,N), $1.5U_{\rm eq}$ (C_{methyl}) or $1.5U_{\rm eq}$ (O).

Data collection: *SMART* (Bruker, 1998); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 1998); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2005); software used to prepare material for publication: *SHELXTL* (Sheldrick, 1997*b*).

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