

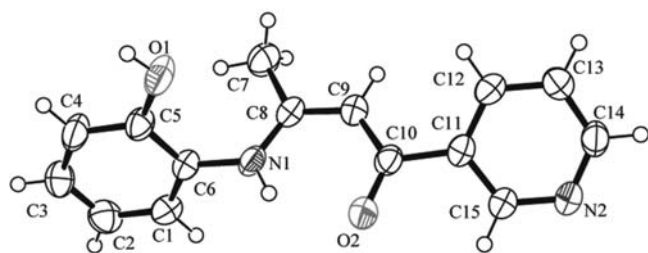
Crystal structure of (Z)-3-(2-hydroxyphenylamino)-1-(pyridin-3-yl)but-2-en-1-one, C₁₅H₁₄N₂O₂

Xin Xiao^{*,I}, Qi-Long Zhang^{II}, Yun-Qian Zhang^I, Bi-Xue Zhu^I and Zhu Tao^I

^I Key Laboratory of Macrocyclic and Supramolecular Chemistry of Guizhou Province, Guizhou University, Guiyang 550025, Guizhou, P. R. China

^{II} Guiyang Medical College, Department of Chemistry, Guiyang 550004, P. R. China

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Abstract

C₁₅H₁₄N₂O₂, monoclinic, *P*2₁/*n* (no. 14), *a* = 12.252(2) Å, *b* = 7.719(1) Å, *c* = 14.500(3) Å, β = 112.731(7)°, *V* = 1264.8 Å³, *Z* = 4, *R*_{gt}(*F*) = 0.050, *wR*_{ref}(*F*²) = 0.148, *T* = 293 K.

Source of material

Solution of 1-(pyridin-3-yl)butane-1,3-dione (0.02 mol) in ethanol (10 ml) was added to a stirred ethanol solution (10 ml) of 2-aminophenol (0.02 mol) in a three-necked flask. The reaction mixture was refluxed for 5 h, then the solution was cooled to room temperature. The yellow precipitated residue was removed from the solution by filtration and then washed with ethanol 3 times. Single crystals suitable for X-ray diffraction were obtained from an ethanol/chloroform mixture (1:1, *v:v*) by slow evaporation at room temperature.

Experimental details

Hydrogen atoms bound to nitrogen and oxygen atoms were found in the difference Fourier maps and refined freely. The other hydrogen atoms bound to carbon atoms were positioned geometrically with *d*(C—H) = 0.93 Å, *U*_{iso}(H) = 1.2 *U*_{eq}(C) for *Csp*² and *d*(C—H) = 0.96 Å, *U*_{iso}(H) = 1.5 *U*_{eq}(C) for *Csp*³.

Discussion

Pyridinone derivatives are of interest due to their biological and pharmacological activity. Recently, some of these compounds were studied as potential hypnotic drugs [1]. Aminophenols in their turn shows the ability to control the packing arrangement often producing the assembly of supramolecular ladders linked via hydrogen bonding and π - π stacking interaction [2,3].

The title crystal structure is built up of the C₁₅H₁₄N₂O₂ molecules with a benzene and a pyridine rings. The bond lengths and bond angles are in the normal ranges. The planes of pyridine and benzene rings form a dihedral angle of 31.0(1)°.

In the title molecule, an intramolecular N1–H1A...O2 hydrogen bond links the aniline NH group to the carbonyl oxygen atom, the

distance of the hydrogen bond is *d*(N1–H1A...O2) = 2.740(2) Å and $\angle$ N1–H1A...O2 = 139(2)°. The intermolecular hydrogen bond, which involves phenol OH-group and the pyridine N atom (*d*(O1–H1B...N2) = 2.771(3) Å and $\angle$ O1–H1B...N2 = 171(3)°), at the same time, there is also another intermolecular hydrogen bond between the NH-group and the carbonyl oxygen of adjacent molecules (*d*(N1–H1A...O2) = 3.169(3) Å and $\angle$ N1–H1A...O2 = 130(2)°). These hydrogen bonds interlink the molecules into a 2D net structure.

Table 1. Data collection and handling.

Crystal:	yellow block, size 0.21 × 0.22 × 0.25 mm
Wavelength:	Mo <i>K</i> _α radiation (0.71073 Å)
μ :	0.9 cm ^{−1}
Diffractometer, scan mode:	Bruker SMART CCD, ϕ/ω
2 θ _{max} :	50°
<i>N</i> (<i>hkl</i>) _{measured} , <i>N</i> (<i>hkl</i>) _{unique} :	12951, 2227
Criterion for <i>I</i> _{obs} , <i>N</i> (<i>hkl</i>) _{gt} :	<i>I</i> _{obs} > 2 σ (<i>I</i> _{obs}), 1489
<i>N</i> (<i>param</i>) _{refined} :	180
Programs:	SHELXS-97, SHELXL-97 [4]

Table 2. Atomic coordinates and displacement parameters (in Å²).

Atom	Site	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{iso}
H(1)	4e	0.5874	0.2684	1.2293	0.056
H(2)	4e	0.5975	0.3146	1.3893	0.065
H(3)	4e	0.4262	0.3642	1.4155	0.066
H(4)	4e	0.2453	0.3567	1.2831	0.065
H(7A)	4e	0.3203	0.0049	1.1061	0.083
H(7B)	4e	0.3555	−0.1248	1.0390	0.083
H(7C)	4e	0.2303	−0.0372	0.9975	0.083
H(9)	4e	0.3217	0.0219	0.8649	0.052
H(12)	4e	0.2437	0.0787	0.6988	0.054
H(13)	4e	0.2495	0.0159	0.5445	0.057
H(14)	4e	0.4147	0.0854	0.5154	0.057
H(15)	4e	0.5559	0.3151	0.7735	0.051
H(1A)	4e	0.443(2)	0.337(3)	1.033(2)	0.059(8)
H(1B)	4e	0.135(3)	0.294(4)	1.118(2)	0.10(1)

* Correspondence author (e-mail: gyhxiaoxin@163.com)

Table 3. Atomic coordinates and displacement parameters (in Å²).

Atom	Site	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> ₁₁	<i>U</i> ₂₂	<i>U</i> ₃₃	<i>U</i> ₁₂	<i>U</i> ₁₃	<i>U</i> ₂₃
C(1)	4e	0.5179(2)	0.2859(3)	1.2397(2)	0.040(1)	0.051(1)	0.051(2)	−0.002(1)	0.021(1)	0.002(1)
C(2)	4e	0.5246(2)	0.3144(3)	1.3358(2)	0.053(2)	0.061(2)	0.041(1)	−0.002(1)	0.010(1)	0.003(1)
C(3)	4e	0.4225(3)	0.3425(3)	1.3513(2)	0.066(2)	0.066(2)	0.036(1)	−0.003(1)	0.024(1)	0.000(1)
C(4)	4e	0.3141(2)	0.3385(4)	1.2718(2)	0.053(2)	0.078(2)	0.042(1)	0.003(1)	0.030(1)	0.000(1)
C(5)	4e	0.3066(2)	0.3077(3)	1.1747(2)	0.041(1)	0.059(1)	0.034(1)	0.004(1)	0.017(1)	0.007(1)
C(6)	4e	0.4112(2)	0.2824(3)	1.1587(2)	0.041(1)	0.043(1)	0.038(1)	0.001(1)	0.021(1)	0.004(1)
C(7)	4e	0.3124(2)	−0.0209(3)	1.0390(2)	0.066(2)	0.053(2)	0.055(2)	−0.008(1)	0.032(1)	0.006(1)
C(8)	4e	0.3610(2)	0.1264(3)	0.9992(2)	0.038(1)	0.044(1)	0.044(1)	0.002(1)	0.023(1)	0.004(1)
C(9)	4e	0.3586(2)	0.1174(3)	0.9034(2)	0.049(2)	0.048(1)	0.040(1)	−0.005(1)	0.024(1)	−0.004(1)
C(10)	4e	0.4080(2)	0.2418(3)	0.8590(2)	0.038(1)	0.045(1)	0.042(1)	0.001(1)	0.022(1)	−0.001(1)
C(11)	4e	0.4015(2)	0.2031(3)	0.7552(2)	0.042(1)	0.039(1)	0.040(1)	0.005(1)	0.022(1)	0.0028(9)
C(12)	4e	0.3084(2)	0.1123(3)	0.6845(2)	0.040(1)	0.051(1)	0.050(1)	−0.003(1)	0.025(1)	−0.001(1)
C(13)	4e	0.3125(2)	0.0724(3)	0.5934(2)	0.045(2)	0.054(1)	0.041(1)	−0.004(1)	0.015(1)	−0.006(1)
C(14)	4e	0.4110(2)	0.1174(3)	0.5759(2)	0.057(2)	0.054(1)	0.039(1)	0.000(1)	0.026(1)	−0.004(1)
N(2)	4e	0.5022(2)	0.2052(3)	0.6417(1)	0.049(1)	0.057(1)	0.042(1)	−0.003(1)	0.027(1)	−0.0009(9)
C(15)	4e	0.4948(2)	0.2494(3)	0.7285(2)	0.045(1)	0.047(1)	0.042(1)	−0.005(1)	0.024(1)	−0.002(1)
N(1)	4e	0.4086(2)	0.2614(3)	1.0598(1)	0.043(1)	0.048(1)	0.042(1)	−0.0022(9)	0.027(1)	0.0026(9)
O(1)	4e	0.2012(2)	0.3013(3)	1.0951(1)	0.039(1)	0.103(2)	0.040(1)	0.0122(9)	0.0205(9)	0.0077(9)
O(2)	4e	0.4602(2)	0.3759(2)	0.9024(1)	0.064(1)	0.053(1)	0.048(1)	−0.0141(8)	0.0324(9)	−0.0085(8)

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