

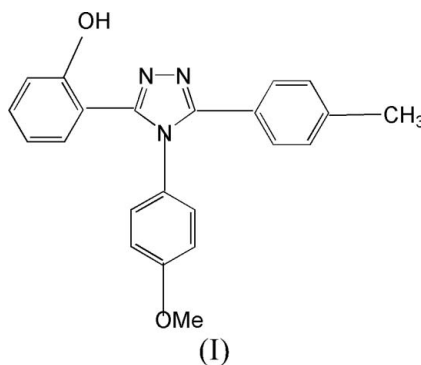
5-(2-Hydroxyphenyl)-4-(4-methoxyphenyl)-
3-(4-methylphenyl)-4*H*-1,2,4-triazoleShu-Ping Zhang, Song Yang, Ying
Zou, Hui-Quan Li and Si-Chang
Shao*Department of Chemistry, Fuyang Normal
College, Fuyang Anhui 236041, People's
Republic of China

Correspondence e-mail: shaosic@fync.edu.cn

Key indicators

Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.041
 wR factor = 0.129
Data-to-parameter ratio = 17.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.In the title compound, $\text{C}_{22}\text{H}_{19}\text{N}_3\text{O}_2$, there are two independent molecules in the asymmetric unit. Intramolecular O—H···N hydrogen bonds are observed in the crystal structure.Received 29 August 2006
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Comment

Recently, we reported the syntheses and structures of some 1,2,4-triazole ligands and their metal complexes (Zhang *et al.*, 2004; Zhang, Liu, Ma *et al.*, 2005; Zhang, Liu, Yang *et al.*, 2005; Zhang, Liu & Shao, 2006; Zhang, Liu, Chen *et al.*, 2006). As an extension of our work on the structural characterization of triazole derivatives, we report here the crystal structure of the title compound, (I).

The asymmetric unit of (I) consists of two independent molecules of 4-(4-methoxyphenyl)-3-(4-methylphenyl)-5-(2-hydroxyphenyl)-4*H*-1,2,4-triazole (Fig. 1). In both molecules, the bond lengths and angles are in normal ranges (Allen *et al.*, 1987). The dihedral angles formed by the triazole ring with the aromatic ring of the hydroxyphenyl, methylphenyl and methoxyphenyl groups are 15.13 (5), 31.00 (5) and 77.54 (4)°, respectively, in one molecule, and 5.25 (7), 58.71 (7) and 83.09 (6)°, respectively, in the other molecule.

The crystal structure is stabilized by intramolecular O—H···N hydrogen bonds (Table 1).

Experimental

The title compound was synthesized according to the literature method of Zhu *et al.* (2000). Equivalent amounts of *p*-methoxyphosphazoanilide (0.2743 g) and *N*-2-hydroxyphenyl-*N'*-4-methylphenylhydrazine (0.2703 g) were reacted in ethanol (10 ml) for 1 h. After allowing the resulting solution to stand in air for 15 d, colourless block-shaped crystals were formed on slow evaporation of the solvent.

Crystal data

$C_{22}H_{19}N_3O_2$
 $M_r = 357.40$
 Triclinic, $P\bar{1}$
 $a = 9.5954 (4) \text{ \AA}$
 $b = 12.4637 (5) \text{ \AA}$
 $c = 15.8750 (6) \text{ \AA}$
 $\alpha = 90.9566 (6)^\circ$
 $\beta = 103.2093 (6)^\circ$
 $\gamma = 90.4278 (6)^\circ$

$V = 1847.95 (13) \text{ \AA}^3$
 $Z = 4$
 $D_x = 1.285 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 $\mu = 0.08 \text{ mm}^{-1}$
 $T = 298 (2) \text{ K}$
 Block, colourless
 $0.18 \times 0.16 \times 0.11 \text{ mm}$

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.985, T_{\max} = 0.991$

22546 measured reflections
 8963 independent reflections
 4446 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$
 $\theta_{\text{max}} = 28.3^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.129$
 $S = 0.95$
 8963 reflections
 500 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0672P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.019$
 $\Delta\rho_{\text{max}} = 0.20 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.12 \text{ e \AA}^{-3}$
 Extinction correction: SHELXL97 (Sheldrick, 1997a)
 Extinction coefficient: 0.0081 (12)

Table 1

Hydrogen-bond geometry ($\text{\AA}, ^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1–H1 \cdots N2	0.946 (15)	1.716 (17)	2.5932 (18)	153 (2)
O3–H2 \cdots N5	0.902 (16)	1.745 (19)	2.547 (3)	147 (2)

The hydroxy H atoms were located in a difference map and refined freely. The remaining H atoms were placed in calculated positions and constrained to ride on their parent atoms, with $C-H = 0.93-0.96 \text{ \AA}$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$, or $1.5U_{\text{eq}}(\text{C})$ for methyl H atoms.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics:

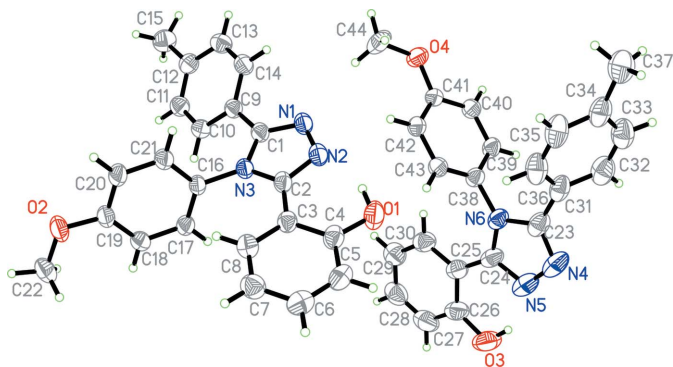


Figure 1

The asymmetric unit of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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