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Key indicators

Single-crystal X-ray study T = 566 KMean $\sigma(\text{C-C}) = 0.004 \text{ Å}$ R factor = 0.046 wR factor = 0.120 Data-to-parameter ratio = 12.7

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

4-(4-Ethoxyphenyl)-3,5-di-2-pyridyl-4*H*-1,2,4-triazole

In the title compound, $C_{20}H_{17}N_5O$, the two pyridyl rings form dihedral angles of 30.0 (1) and 21.0 (1)° with the triazole ring, and the dihedral angle between the triazole and benzene rings is 71.4 (1)°. The crystal packing is stabilized by $C-H\cdots N$ hydrogen bonds.

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Comment

1,2,4-Triazole and its derivatives constitute a promising class of ligands that are widely used in the synthesis of various complexes (Haasnoot, 2000). It is possible to produce complexes with different structures and properties, using the extensive coordinating capabilities of the N atoms of the triazole ring. We report here the crystal structure of the title compound, (I).



In (I), the pyridyl groups and the benzene ring lie in a propeller arrangement around the central 1,2,4-triazole ring (Fig. 1), thereby minimizing the steric effects among these rings. The dihedral angles between the 4-ethoxylphenyl ring and the two pyridyl rings (N5/C3–C7 and N4/C8–C12) are 70.9 (1) and 67.0 (1)°, respectively. The two pyridyl rings form dihedral angles of 30.0 (1) and 21.0 (1)°, respectively, with the triazole ring, and the dihedral angle between the triazole ring and the benzene ring is 71.4 (1)°. The crystal packing is stabilized by C–H···N hydrogen bonds (Table 1).

Experimental

Compound (I) was synthesized according to a literature method (Zhang *et al.*, 2004). Equivalent amounts of *p*-ethoxyphenyl-phosphazoanilide and *N*,*N'*-dipyridoylhydrazine were reacted in *N*,*N'*-dimethylaniline for 3 h at 463 K, with stirring. Colourless block-shaped crystals were obtained by slow evaporation of an acetone solution. The crystals were collected and dried in a vacuum desiccator using anhydrous CaCl₂ (yield 52%). Analysis found: C 69.89, H 5.02, N 20.44%; calculated for $C_{20}H_{17}N_5O$: C 69.96, H 4.99, N 20.40%

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Crystal data

 $\begin{array}{l} C_{20}H_{17}N_5O\\ M_r = 343.39\\ \text{Triclinic, } P\overline{1}\\ a = 8.094~(5)~\text{\AA}\\ b = 9.468~(6)~\text{\AA}\\ c = 13.152~(8)~\text{\AA}\\ \alpha = 104.657~(9)^{\circ}\\ \beta = 92.706~(8)^{\circ}\\ \gamma = 115.133~(8)^{\circ}\\ V = 868.9~(9)~\text{\AA}^{3} \end{array}$

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.983, T_{\max} = 0.992$ 4239 measured reflections

Refinement

Refinement on F^2
$R[F^2 > 2\sigma(F^2)] = 0.046$
$wR(F^2) = 0.120$
S = 0.87
2989 reflections
235 parameters

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C9-H9A\cdots N1^{i}$ $C18-H18A\cdots N2^{ii}$	0.93	2.46	3.308 (4)	152
	0.93	2.53	3.459 (4)	174

Z = 2

 $D_x = 1.312 \text{ Mg m}^{-3}$

Cell parameters from 1021

 $0.20 \times 0.18 \times 0.10 \text{ mm}$

2989 independent reflections

1790 reflections with $I > 2\sigma(I)$

H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0621P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$

Mo $K\alpha$ radiation

reflections

 $\theta = 4.6-27.3^{\circ}$

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

T = 293 (2) K

Block, white

 $R_{\rm int} = 0.033$

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

 $h = -9 \rightarrow 7$

 $k = -5 \rightarrow 11$

 $l = -15 \rightarrow 15$

 $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.23 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.25 \text{ e } \text{\AA}^{-3}$

Symmetry codes: (i) -x + 2, -y + 1, -z; (ii) -x + 2, -y, -z.

All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C–H distances of 0.93–0.97 Å, and with $U_{iso}(H) = 1.2U_{eq}(C)$ or 1.5(methyl C).

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



Figure 1

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

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

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