organic papers

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Key indicators

Single-crystal X-ray study T = 295 KMean σ (C–C) = 0.004 Å R factor = 0.054 wR factor = 0.130 Data-to-parameter ratio = 17.5

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

4-[(Z)-(2,4-Dimethylphenylamino)phenylmethylene]-3-methyl-1-phenyl-1H-pyrazol-5(4**H**)-one

The NH unit on the exocyclic C=C double bond in the title compound, C₂₅H₂₃N₃O, interacts with the carbonyl group through an intramolecular hydrogen bond.

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Comment

The title compound, (I) (Fig. 1) has two methyl substituents in the 2,4-dimethylphenylamino portion of the molecule; the general features are similar to those found in the analogous 2tolylamino (Bao et al., 2004) and 4-tolylamino (Ma, 2005) derivatives. The compound is used to chelate to a divalent transition metal (Ma et al., 2006).



Experimental

1-Phenyl-3-methyl-4-benzoyl-5-pyrazolone (1.60 g, 5.8 mm mol) and 2,4-dimethylaniline (0.73 g, 6.0 mm mol) were dissolved in ethanol (35 ml); formic acid (0.5 ml) was added to catalyse the reaction. The solution was heated under reflux for 8 h. The solvent was removed and the pure product obtained upon recrystallization from a 1:1 ethanol/n-heptane mixture in 80% yield. Crystals were grown from an ethanol solution of the compound. Elemental analysis calculated for C25H23N3O: C 79.29, H 9.15, N 11.56%; found: C 79.10, H 9.26, N 11.27%.

Crystal data	
C ₂₅ H ₂₃ N ₃ O	V = 1035.6 (2) Å ³
$M_r = 381.46$	Z = 2
Triclinic, P1	$D_x = 1.223 \text{ Mg m}^{-3}$
a = 7.909 (1) Å	Mo $K\alpha$ radiation
b = 11.200 (1) Å	$\mu = 0.08 \text{ mm}^{-1}$
c = 13.239 (1) Å	T = 295 (2) K
$\alpha = 106.410 \ (2)^{\circ}$	Block, yellow
$\beta = 106.327 \ (2)^{\circ}$	$0.10 \times 0.06 \times 0.04 \text{ mm}$
$\gamma = 100.390 \ (2)^{\circ}$	
Data collection	
Bruker APEX area-detector	4633 independent reflect

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diffractometer ω and φ scans Absorption correction: none 10060 measured reflections

4633 independent reflections 2018 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.078$ $\theta_{\rm max} = 27.5^\circ$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.054$	$w = 1/[\sigma^2(F_o^2) + (0.047P)^2]$
$wR(F^2) = 0.130$	where $P = (F_0^2 + 2F_c^2)/3$
S = 0.83	$(\Delta/\sigma)_{\rm max} = 0.001$
4633 reflections	$\Delta \rho_{\rm max} = 0.18 \ {\rm e} \ {\rm \AA}^{-3}$
265 parameters	$\Delta \rho_{\rm min} = -0.15 \text{ e } \text{\AA}^{-3}$

H atoms were positioned geometrically and were included in the refinement in the riding-model approximation [phenyl C–H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$; methyl C–H = 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$]; the methyl groups were rotated to fit the electron density. The amino H atom was similarly constrained [N–H = 0.86 Å and $U_{iso}H = 1.2U_{eq}(N)$].

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *SHELXL97*.

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Figure 1

The molecular structure of (I) with displacement ellipsoids drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radii. The dashed line denotes the intramolecular hydrogen bond.

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