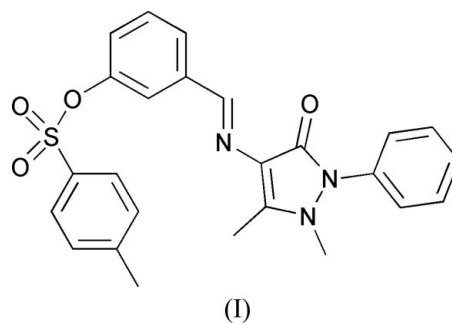


**(E)-3-[(1,5-Dimethyl-3-oxo-2-phenyl-1H-pyrazol-4-ylimino)methylene]phenyl 4-methylbenzenesulfonate****Xin Chen\* and Ming Yu**College of Sciences, Tianjin University of  
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People's Republic of ChinaCorrespondence e-mail:  
chen\_xin9999@163.com**Key indicators**Single-crystal X-ray study  
 $T = 294$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.048  
 $wR$  factor = 0.130  
Data-to-parameter ratio = 15.6For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

In the title compound,  $\text{C}_{25}\text{H}_{23}\text{N}_3\text{O}_4\text{S}$ , the central benzene ring makes dihedral angles of  $8.73$  (3),  $68.16$  (7) and  $41.67$  (7) $^\circ$  with the pyrazolone ring, the sulfur-bound benzene ring and the phenyl ring, respectively. The crystal packing is stabilized by a weak non-classical intermolecular  $\text{C}-\text{H}\cdots\text{O}=\text{C}$  hydrogen bond that forms a centrosymmetric dimer.

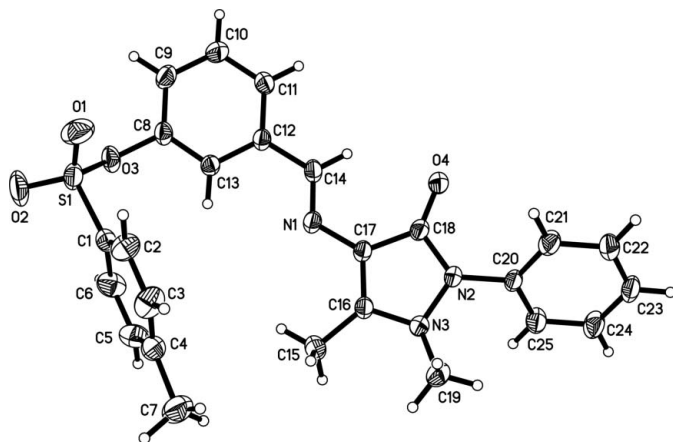
Received 16 October 2006  
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Schiff base ligands have received considerable attention in biology and chemistry (Kahwa *et al.*, 1986). Many Schiff base derivatives have been synthesized and employed to develop protein and enzyme mimics (Santos *et al.*, 2001). Among a large number of compounds, 4-amino-1,5-dimethyl-2-phenylpyrazol-3-one forms a variety of Schiff bases with aldehydes, and the synthesis and crystal structures of some of them, such as (*E*)-1,5-dimethyl-4-[2-[2-(2-nitrophenoxy)-ethoxy]benzylideneamino]-2-phenyl-1H-pyrazol-3(2H)-one (Diao & Chen, 2006) and (*E*)-4-[4-(4-chlorobenzoyloxy)benzylideneamino]-1,5-dimethyl-2-phenyl-1H-pyrazol-3(2H)-one (Hu, 2006) have been reported. Structural investigations may provide useful information concerning the coordination properties of Schiff bases functioning as ligands. We report here the synthesis and crystal structure of the title compound, (I) (Fig. 1).



The bond lengths and angles of (I) are within normal ranges (Allen *et al.*, 1987). The pyrazolone ring (C16–C18/N1/N2/N3/O4) is almost planar, with an r.m.s. deviation for fitted atoms of  $0.0369$  Å. It makes a dihedral angle of  $48.81$  (6) $^\circ$  with the attached phenyl ring (C20–C25). The central benzene ring (C8–C14/O3) is nearly planar, with an r.m.s. deviation for fitted atoms of  $0.0308$  Å. This group makes dihedral angles of  $8.73$  (3),  $68.16$  (7) and  $41.67$  (7) $^\circ$  with the the pyrazolone ring, the sulfur-bound benzene ring (C1–C6) and the phenyl ring, respectively.

The crystal packing of (I) is stabilized by a weak non-classical intermolecular  $\text{C}-\text{H}\cdots\text{O}=\text{C}$  hydrogen bond (Table 1) that forms a centrosymmetric dimer (Fig. 2).



**Figure 1**  
The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level.

## Experimental

An anhydrous ethanol solution (50 ml) of 3-formylphenyl 4-methylbenzenesulfonate (2.76 g, 10 mmol) was added to an anhydrous ethanol solution (50 ml) of 4-amino-1,5-dimethyl-2-phenylpyrazol-3-one (2.03 g, 10 mmol) and the mixture was stirred at 350 K for 3 h under  $N_2$ , giving a yellow precipitate. The product was isolated, recrystallized from acetonitrile and dried in a vacuum to give pure (I) in 85% yield. Yellow single crystals suitable for X-ray crystallographic analysis were obtained by slow evaporation of an acetonitrile solution.

### Crystal data

$C_{25}H_{25}N_3O_4S$   
 $M_r = 461.53$   
 Monoclinic,  $P2_1/c$   
 $a = 9.2661$  (18) Å  
 $b = 9.5978$  (18) Å  
 $c = 26.195$  (5) Å  
 $\beta = 97.693$  (3)°  
 $V = 2308.7$  (8) Å<sup>3</sup>

$Z = 4$   
 $D_x = 1.328$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 $\mu = 0.18$  mm<sup>-1</sup>  
 $T = 294$  (2) K  
 Block, yellow  
 $0.40 \times 0.28 \times 0.18$  mm

### Data collection

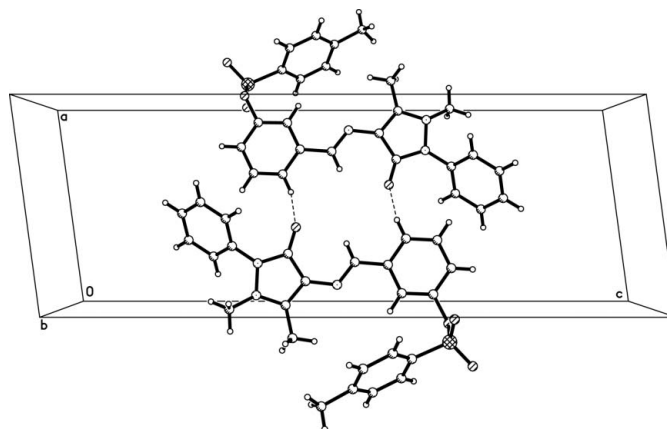
Bruker SMART APEX CCD area detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.921$ ,  $T_{\max} = 0.969$

12673 measured reflections  
 4685 independent reflections  
 2743 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.039$   
 $\theta_{\text{max}} = 26.3^\circ$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.048$   
 $wR(F^2) = 0.130$   
 $S = 1.00$   
 4685 reflections  
 301 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.06P)^2 + 0.3197P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.002$   
 $\Delta\rho_{\text{max}} = 0.20$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.32$  e Å<sup>-3</sup>



**Figure 2**  
A partial packing diagram of (I), viewed along the  $b$  axis, showing C—H...O hydrogen bonds (dashed lines).

**Table 1**  
Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C11-H11\cdots O4^i$	0.93	2.43	3.227 (3)	143

Symmetry code: (i)  $-x + 1, -y + 1, -z + 1$ .

H atoms were included in calculated positions, with C—H = 0.93–0.96 Å, and refined as riding, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{methyl C})$ .

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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