### organic papers

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# Ming Yu,<sup>a</sup> Jin-Ling Wang<sup>b\*</sup> and Fang-Ming Miao<sup>b</sup>

<sup>a</sup>Department of Scientific Technology, The Scientific and Technology University of Tianjin, Tianjin 300222, People's Republic of China, and <sup>b</sup>College of Chemical and Life Science, Tianjin Normal University, Tianjin 300074, People's Republic of China

Correspondence e-mail: wangjinling43@eyou.com

#### Key indicators

Single-crystal X-ray study T = 297 K Mean  $\sigma$ (C–C) = 0.005 Å Disorder in main residue R factor = 0.056 wR factor = 0.172 Data-to-parameter ratio = 11.3

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

## 1,5-Dimethyl-4-{[(*E*)-3-oxo-3-(2-thienyl)-1-(trifluoromethyl)-1-propenyl]amino}-2-phenyl-1,2-dihydro-3*H*-pyrazol-3-one

In the title compound,  $C_{19}H_{16}F_3N_3O_2S$ , the side-chain carbonyl group, the adjacent double bond and the amine N atom are essentially coplanar, with the largest deviation from the mean plane being 0.010 (2) Å. A strong intramolecular N-H···O hydrogen bond [N···O 2.702 (3) Å and N-H···O 139 (3)°] is observed, leading to an enamine tautomer in the solid state.

#### Comment

Schiff base ligands derived from 4-aminoantipyrine, such as salicylidene-4-aminoantipyrine, 5-chlorosalicylidene-4-aminoantipyrine, 2-hydroxy-1-naphthylidene-4-aminoantipyrine and 2-hydroxy-acetophenonylidene-4-aminoantipyrine, have been reported by McLendon & Martell (1976) and by Radhakrishnan & Joseph (1976). In order to study the coordination properties of Schiff base compounds, we have synthesized some of these compounds, and we report here the structure of the title compound, (I).



A view of the molecule of (I) is shown in Fig. 1. Atoms C6, C7, C8 and O1 of the thenoyltrifluoroacetone moiety and atom N3 of 4-aminoantipyrine are essentially coplanar, the largest deviation from the mean plane being 0.010 (2) Å for atom C6; the dihedral angle between this plane and the thienyl plane is 22.48 (3)°. The bond lengths within this part of the molecule [N3–C6 = 1.340 (4) Å, C6–C7 = 1.401 (4) Å, C7–C8 = 1.396 (4) Å and C8–O1 = 1.235 (4) Å] lie between the classical double- and single-bond lengths. These results clearly indicate that there is electron delocalization over this segment.

An intramolecular hydrogen bond is observed (Table 1), which indicates that the molecule exists as the enamine tautomer. This situation is completely different from that of 3-(2,3-dihydro-1,5-dimethyl-3-oxo-2-phenylpyrazol-4-ylimino)-4,4,4-trifluoro-1-(2-thienyl)butane-1,2-dione (Wang *et al.*, 2002), but similar to that of pyrazolinehydrazide (Liu *et al.*, 2001). The displacements of atoms C5 and C21 from the

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pyrazolinone ring are 0.454(7) and -0.479(5) Å, respectively, showing that the methyl group bonded to N1 and the phenyl group bonded to N2 are on opposite sides of the ring. Intermolecular contact distances of interest are:  $O2 \cdot \cdot \cdot S1^{i} =$ 3.040(4) Å, C4-H41···O1<sup>i</sup> = 3.380(4) Å [angle of 135.01 (4)° at H41], C4-H41···F2<sup>i</sup> = 2.930 (4) Å [angle of 138.96 (4)° at H41], C5-H53···F2 $A^{i}$  = 3.418 (5) Å [angle of 163.71 (5)° at H53], C4–H42···O1<sup>ii</sup> = 3.380 (4) Å [angle of 135.01 (5)° at H42], and C4-H43···F2 $A^{iii}$  = 3.366 (5) Å [angle of 116.32 (5)° at H43] [symmetry codes: (i) 1 - x, -y, 1 - z; (ii) -1 + x, y, z; (iii) -x, -y, 1 - z].

The torsion angle C21-N2-N1-C5 is 51.2 (4)°, close to the values of 55.6 (3) and 55.7 (3) $^{\circ}$  for the compounds 4-(salicylideneamino)-2-3-dimethyl-1-phenyl-3-pyrazolin-5-one (Chumakov et al., 2000) and 4-{[(1E)-(2-hydroxyphenyl) methylidene]amino]-1,5-dimethyl-2-phenyl-2,3-dihydro-1Hpyrazol-3-one (Hökelek et al., 2001). By contrast, conjugation leads to a small torsion angle of 7.75° in 4-(antipyrin-4-yliminomethyl)benzoic acid (Zhang et al., 2002).

The mean plane through the side-chain carbonyl group, the adjacent double bond and the amino N atom is nearly perpendicular to the pyrazolinone ring, with a dihedral angle of 84.10 (6)°, close to the value of 82.43 (8)° reported by Wang et al. (2002). However, the dihedral angle between the pyrazolinone ring and the phenyl ring is  $115.26(5)^\circ$ , quite different from the value of  $41.7 (5)^{\circ}$  in 4-(salicylideneamino)-2,3-dimethyl-1-phenyl-3-pyrazolin-5-one (Chumakov et al., 2000). These results indicate that the compounds may have quite different coordination properties as ligands to metal ions.

#### **Experimental**

Ethanol solutions of 0.1 mol of 4-aminoantipyrine and 0.1 mol of thenovltrifluoroacetone were refluxed together for 4 h over a steam bath. The excess solvent was removed by evaporation and the concentrated solution was cooled in an ice bath, with stirring. The title compound separated out as a cream-colored powder, which was collected and dried in air (m.p. 434 K). It was recrystallized from

Z = 2

 $D_x = 1.460 \text{ Mg m}^{-3}$ 

Cell parameters from 3101

Mo Ka radiation

reflections

 $\theta = 1.5 - 25.1^{\circ}$  $\mu = 0.22~\mathrm{mm}^{-1}$ 

T = 297 (2) K

Prism, yellow

 $R_{\rm int} = 0.018$ 

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

 $h = -5 \rightarrow 6$  $k = -12 \rightarrow 8$ 

 $l = -18 \rightarrow 17$ 

 $0.35 \times 0.30 \times 0.25 \text{ mm}$ 

3245 independent reflections 2229 reflections with  $I > 2\sigma(I)$ 

Crystal data

C19H16F3N3O2S  $M_{\rm m} = 407.42$ Triclinic, P1 a = 5.8063 (10) Å $b = 10.9254 (17) \text{ \AA}$ c = 15.288 (2) Å $\alpha = 74.790 \ (3)^{\circ}$  $\beta = 97.463 \ (3)^{\circ}$  $\gamma = 94.731 \ (4)^{\circ}$  $V = 926.7 (2) \text{ Å}^3$ 

#### Data collection

Bruker SMART 1000 CCD
diffractometer
$\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 1999)
$T_{\rm min} = 0.926, \ T_{\rm max} = 0.946$
3860 measured reflections
Refinement
Refinement on $F^2$

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0983P)^2$
$R[F^2 > 2\sigma(F^2)] = 0.057$	+ 0.1991P]
$wR(F^2) = 0.173$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} = 0.002$
3245 reflections	$\Delta \rho_{\rm max} = 0.33 \ {\rm e} \ {\rm \AA}^{-3}$
286 parameters	$\Delta \rho_{\rm min} = -0.45 \ {\rm e} \ {\rm \AA}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
-	Extinction coefficient: $0.011(4)$

#### Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
N3−H3···O1	0.81 (4)	2.04 (4)	2.702 (3)	139 (3)

Some rather high residual electron densities close to the CF<sub>3</sub> group and staggered with respect to the F atoms appeared in a difference Fourier map. A disordered distribution of the F atoms was refined with geometrical restraints; this led to much lower residual electron density and to lower R and wR2 values.

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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