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Key indicators

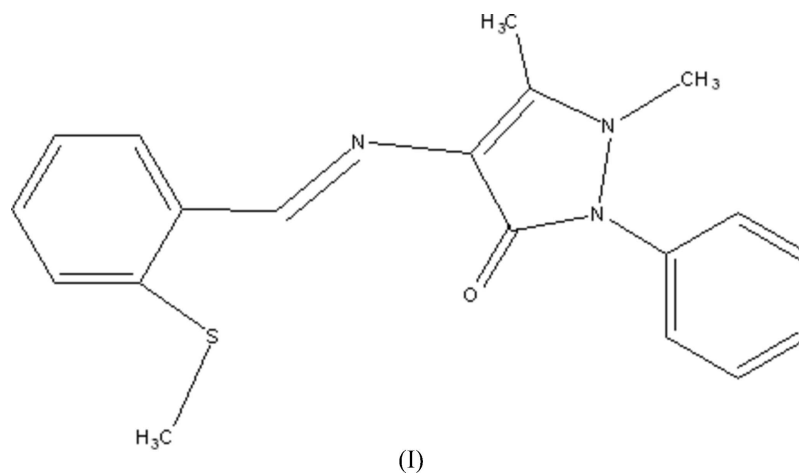
Single-crystal X-ray study
 $T = 273$ K
Mean $\sigma(\text{C}-\text{C}) = 0.050$ Å
 R factor = 0.040
 wR factor = 0.111
Data-to-parameter ratio = 20.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.1,5-Dimethyl-4-[2-(methylsulfanyl)benzyl-
ideneamino]-2-phenyl-1*H*-pyrazol-3(2*H*)-one

In the title compound, $\text{C}_{19}\text{H}_{19}\text{N}_3\text{OS}$, a Schiff base, there are three different intramolecular hydrogen bonds ($\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{S}$), which stabilize the structure; this is further extended into one-dimensional chains parallel to the a axis via intermolecular hydrogen bonds.

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Comment

Schiff base ligands and their metal complexes are interesting. For example, Schiff bases have exhibited solvatochromicity, and are suitable NLO (non-linear optical) materials (Alemi & Shaabani, 2000). They are also useful in asymmetric oxidation of methyl phenyl sulfide and enantioselective epoxidation (Kim & Shin, 1999). In this paper, we report the synthesis and crystal structure of the title Schiff base, (I).



The molecular structure of (I) (Fig. 1) contains three intramolecular hydrogen bonds ($\text{C8}-\text{H8}\cdots\text{O1}$ and $\text{C8}-\text{H8}\cdots\text{S1}$; Table 2). The $\text{C8}-\text{N1}$ distance of 1.255 (4) Å is indicative of a normal $\text{C}=\text{N}$ bond. The other $\text{C}-\text{N}$, $\text{C}-\text{S}$ and $\text{C}-\text{C}$ distances show no remarkable features.

In addition, intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 2) between the molecules lead to one-dimensional chains parallel to the a axis (Fig. 2).

Experimental

Under nitrogen, a mixture of 4-amino-1,5-dimethyl-2-phenyl-1,2-dihydropyrazol-3-one (2.03 g, 10 mmol), Na_2SO_4 (3.0 g) and 2-methylthiobenzaldehyde (1.52 g, 10 mmol) in absolute ethanol (20 ml) was refluxed for about 12 h, yielding a yellow precipitate. The product was collected by vacuum filtration and washed with ethanol. The crude solid was redissolved in CH_2Cl_2 (100 ml) and washed with water (2×10 ml) and brine (10 ml). After drying over Na_2SO_4 , the

solvent was removed under vacuum, and a yellow solid was isolated in 92% yield (3.1 g). Yellow single crystals of the Schiff base (I) suitable for X-ray analysis were grown from CH₂Cl₂ and absolute ethanol (4:1) by slow evaporation of the solvents at room temperature over a period of about a week.

Crystal data

C₁₉H₁₉N₃OS
M_r = 337.43
 Orthorhombic, *P*2₁2₁2₁
a = 6.9535 (9) Å
b = 12.9078 (17) Å
c = 19.800 (3) Å
V = 1777.1 (4) Å³
Z = 4
 Mo *K*α radiation
 μ = 0.19 mm⁻¹
T = 273 (2) K
 0.43 × 0.35 × 0.34 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.922, *T_{max}* = 0.938
 13405 measured reflections
 4287 independent reflections
 2166 reflections with *I* > 2σ(*I*)
R_{int} = 0.020

Refinement

R[*F*² > 2σ(*F*²)] = 0.040
wR(*F*²) = 0.111
S = 0.91
 4287 reflections
 208 parameters
 1 restraint
 H-atom parameters constrained
 Δρ_{max} = 0.22 e Å⁻³
 Δρ_{min} = -0.34 e Å⁻³
 Absolute structure: Flack (1983),
 1759 Friedel pairs
 Flack parameter: -0.02 (9)

Table 1

Selected geometric parameters (Å, °).

C18—S1	1.78 (2)	C12—N1	1.26 (4)
C11—O1	1.24 (4)	C19—S1	1.73 (6)
C11—N2	1.40 (4)	N2—N3	1.40 (4)
N1—C10—C11	130 (3)	C19—S1—C18	105 (2)
C12—N1—C10	121 (3)		

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C12—H12...S1	0.93	2.62	3.03 (4)	107
C12—H12...O1	0.93	2.41	3.07 (4)	128
C7—H7A...O1 ⁱ	0.96	2.55	3.37 (5)	142
C8—H8C...O1 ⁱ	0.96	2.54	3.49 (6)	170

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

All H atoms were placed in calculated positions [C—H = 0.93 (aromatic) or 0.96 Å (methyl)] and refined using a riding model, with *U*_{iso}(H) = 1.2*U*_{eq}(aromatic C) or 1.5*U*_{eq}(methyl C).

Data collection: SMART (Bruker, 1999); cell refinement: SAINT; data reduction: SAINT (Bruker, 1999); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1998); software used to prepare material for publication: SHELXTL.

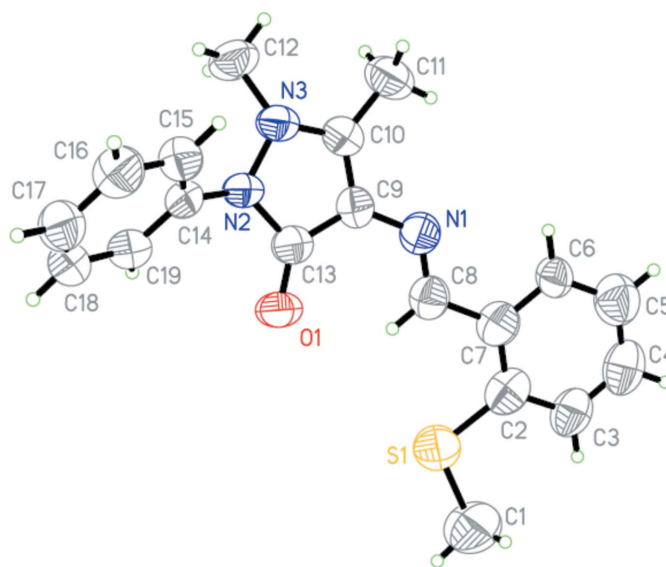


Figure 1

The molecular structure of (I), showing the atomic numbering scheme. Non-H atoms are shown as 50% probability displacement ellipsoids.

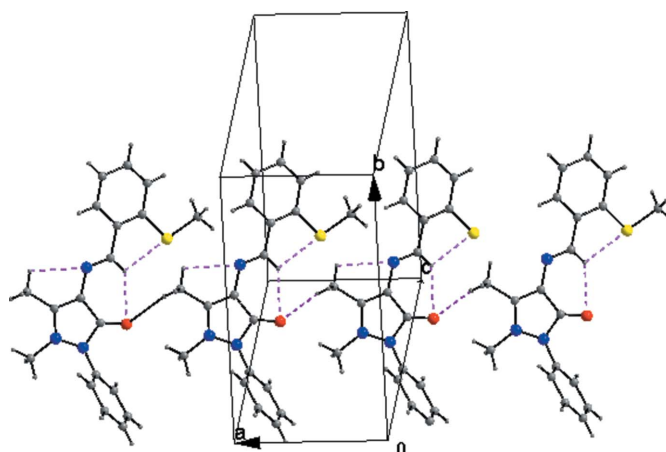


Figure 2

A chain of molecules linked by hydrogen bonds (dashed lines).

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