

**(E)-4-(4-Ethoxy-3-methoxybenzylideneamino)-  
1,5-dimethyl-2-phenylpyrazolidin-3-one****Yun Yang,\* Shu-Lin Ma, Lan-Li  
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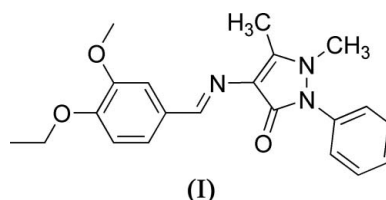
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**Key indicators**Single-crystal X-ray study  
 $T = 294$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.040  
 $wR$  factor = 0.115  
Data-to-parameter ratio = 15.8For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

In the title compound,  $\text{C}_{21}\text{H}_{23}\text{N}_3\text{O}_3$ , the substituted vanillin group makes dihedral angles of  $48.16$  (5) and  $4.01$  (4) $^\circ$  with the terminal phenyl ring and the pyrazolone ring, respectively. An intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bond helps to consolidate the crystal packing.

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Schiff base ligands have played an important role in the development of coordination chemistry since the late 19th century. Consequently, a large number of these species have been reported to be superior reagents in biological, pharmacological, clinical and analytical applications (Wang *et al.*, 2005; Yang *et al.*, 2005). As part of an investigation of their crystal structures, which will provide useful information for the coordination properties of Schiff bases functioning as ligands, we report here the synthesis and molecular structure of the title compound, (I).



In (I), the bond lengths and angles (Table 1) are within normal ranges. The substituted vanillin group (C3–C8/C10/O1/O2) is planar, with an r.m.s. deviation of  $0.0090$  (17) Å (Fig. 1). The pyrazolone ring (C11–C13/N1–N3/O3) is also essentially planar, with an r.m.s. deviation of  $0.0329$  (19) Å. The vanillin group makes dihedral angles of  $4.01$  (4) and  $48.16$  (5) $^\circ$  with the pyrazolone ring and the terminal phenyl ring (C16–C21), respectively. The pyrazolone ring makes a dihedral angle of  $44.43$  (6) $^\circ$  with the phenyl ring.

The crystal packing is stabilized by a weak intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bond (Table 2), forming an inversion-related dimer (Fig. 2).

**Experimental**

An anhydrous ethanol solution (50 ml) of 4-ethoxy-3-methoxybenzaldehyde (1.80 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 stirred at 350 K for 5 h under nitrogen, whereupon a yellow precipitate appeared. The product was then isolated and recrystallized from ethanol, and then dried *in vacuo* to give compound (I) in 83% yield. Yellow single crystals suitable for X-ray analysis were obtained by slow evaporation of an ethanol solution.

Crystal data

$C_{21}H_{23}N_3O_3$   
 $M_r = 365.42$   
 Monoclinic,  $P2_1/c$   
 $a = 16.214 (4) \text{ \AA}$   
 $b = 8.4979 (18) \text{ \AA}$   
 $c = 14.888 (3) \text{ \AA}$   
 $\beta = 109.204 (4)^\circ$   
 $V = 1937.1 (7) \text{ \AA}^3$

$Z = 4$   
 $D_x = 1.253 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.09 \text{ mm}^{-1}$   
 $T = 294 (2) \text{ K}$   
 Block, yellow  
 $0.22 \times 0.20 \times 0.18 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.962, T_{\max} = 0.983$

10562 measured reflections  
 3944 independent reflections  
 2373 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.030$   
 $\theta_{\text{max}} = 26.4^\circ$

Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.040$   
 $wR(F^2) = 0.115$   
 $S = 1.00$   
 3944 reflections  
 249 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0497P)^2 + 0.3272P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.003$   
 $\Delta\rho_{\text{max}} = 0.17 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.13 \text{ e \AA}^{-3}$   
 Extinction correction: SHELXL97  
 Extinction coefficient: 0.0083 (10)

Table 1

Selected geometric parameters ( $\text{\AA}, ^\circ$ ).

O1—C3	1.365 (2)	N1—C11	1.391 (2)
O1—C2	1.4330 (19)	N2—C12	1.368 (2)
O2—C8	1.3668 (18)	N2—N3	1.4047 (19)
O2—C9	1.414 (2)	N2—C15	1.462 (2)
O3—C13	1.235 (2)	N3—C13	1.392 (2)
N1—C10	1.276 (2)	N3—C16	1.422 (2)
C3—O1—C2	117.53 (13)	O1—C3—C8	115.12 (14)
C8—O2—C9	117.70 (13)	O2—C8—C7	125.38 (15)
C10—N1—C11	121.67 (16)	O2—C8—C3	114.41 (14)
C12—N2—N3	106.19 (14)	C12—C11—N1	122.30 (16)
C12—N2—C15	121.84 (15)	N1—C11—C13	129.79 (17)
N3—N2—C15	115.86 (13)	C11—C12—N2	110.76 (15)
C13—N3—N2	110.13 (13)	O3—C13—N3	124.20 (16)
C13—N3—C16	126.13 (15)	O3—C13—C11	131.17 (18)
N2—N3—C16	119.93 (14)	N3—C13—C11	104.61 (15)
O1—C3—C4	125.86 (15)		

Table 2

Hydrogen-bond geometry ( $\text{\AA}, ^\circ$ ).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C15—H15C $\cdots$ O1 <sup>i</sup>	0.96	2.56	3.310 (2)	135

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

H atoms were included in calculated positions ( $C-H = 0.93-0.96 \text{ \AA}$ ) and refined using the riding-model approximation, with  $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$  or  $1.5U_{\text{eq}}(\text{methyl } C)$ .

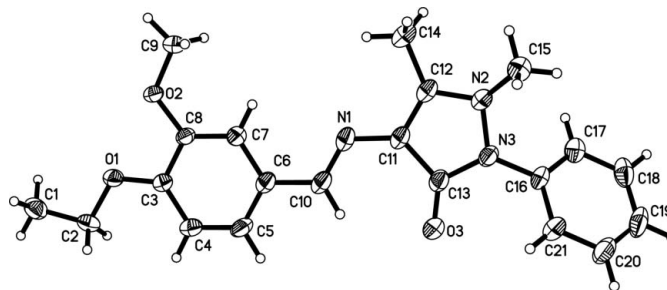


Figure 1

The structure of (I), with displacement ellipsoids for non-H atoms drawn at the 30% probability level.

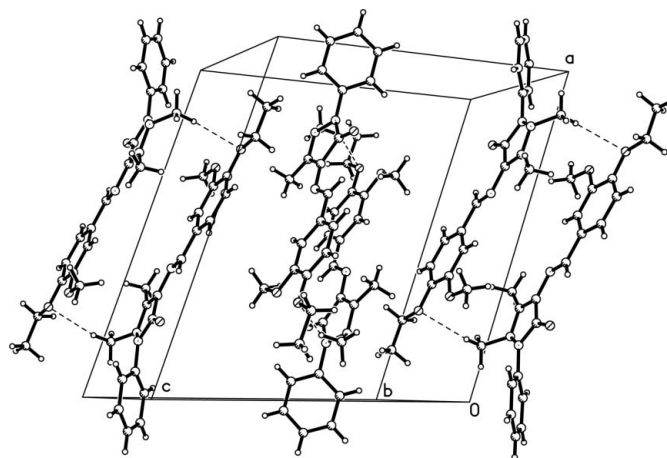


Figure 2

A packing view of (I), showing  $C-H\cdots O$  hydrogen bonds as dashed lines.

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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