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## Structure Reports

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## Wen-Jun Zhang,* Zhong-Yu Duan and Xin Zhao

Hebei University of Technology, Tianjin 300130, People's Republic of China

Correspondence e-mail:
zhang_wenjun99@163.com

## Key indicators

Single-crystal X-ray study
$T=294 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.045$
$w R$ factor $=0.130$
Data-to-parameter ratio $=13.6$

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

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## (E)-4-[3-Ethoxy-4-(2-phenoxyethoxy)benzyl-ideneamino]-1,5-dimethyl-2-phenyl-1H-pyrazol-3(2H)-one

The title compound, $\mathrm{C}_{28} \mathrm{H}_{29} \mathrm{~N}_{3} \mathrm{O}_{4}$, was prepared by the reaction of 4-amino-1,5-dimethyl-2-phenylpyrazol-3-one and 3-ethoxy-4-(2-phenoxyethoxy)benzaldehyde. The vanillin group makes dihedral angles of 73.08 (8) and $66.54(6)^{\circ}$ with the planes of the two terminal phenyl rings, and an angle of 30.49 (6) ${ }^{\circ}$ with the pyrazolone ring plane. The crystal structure is stabilized by intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds that form centrosymmetric dimers.

## Comment

Metal complexes based on Schiff bases have received a good deal of attention in the development of coordination chemistry because of their significant biological activity (Kahwa et al., 1986). Consequently, a significant effort has been devoted to the synthesis of Schiff base derivatives to develop protein and enzyme mimics (Santos et al., 2001). Among the large number of compounds, 4-amino-1,5-dimethyl-2-phenyl-pyrazol-3-one forms a variety of Schiff bases with aldehydes, and the synthesis and crystal structures of some them, such as ( $E$ )-(4)-(3-ethoxy-4-hydroxybenzylideneamino)-1,5-dimethyl-2-phenyl-1 $H$-pyrazol-3( $2 H$ )-one (Han \& Zhen, 2005) and (E)-4-[4-(4-chlorobenzyloxy)-3-ethoxybenzylideneamino]-1,5-di-methyl-2-phenyl-1H-pyrazol-3(2H)-one (Zhang et al., 2006) have been reported. We report here the synthesis and structure of the title compound, (I)

(I)

In the title molecule (Fig. 1), bond lengths and angles are within normal ranges (Allen et al., 1987). The pyrazolone ring (C18-C20/N2/N3/O4) is almost planar, with an r.m.s. deviation for fitted atoms of $0.036 \AA$. It makes a dihedral angle of 46.35 (7) ${ }^{\circ}$ with the attached phenyl ring (C23-C28). The vanillin group ( $\mathrm{C} 9 / \mathrm{C} 10 / \mathrm{C} 13-\mathrm{C} 17 / \mathrm{O} 2 / \mathrm{O} 3$ ) is planar, with an r.m.s. deviation for fitted atoms of $0.014 \AA$. This group makes dihedral angles of 30.49 (6), 73.08 (8) and 66.54 (6) ${ }^{\circ}$, respectively, with the pyrazolone ring ( $\mathrm{C} 18-\mathrm{C} 20 / \mathrm{N} 2 / \mathrm{N} 3 / \mathrm{O} 4$ ) and the terminal C1-C6 and C23-C28 phenyl rings, respectively.

The crystal structure is stabilized by weak non-classical intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}=\mathrm{C}$ hydrogen bonds (Table 1) that form centrosymmetric dimers (Fig. 2).

## Experimental

An anhydrous ethanol solution ( 50 ml ) of 3-ethoxy-4-(2-phenoxyethoxy)benzaldehyde ( $2.86 \mathrm{~g}, 10 \mathrm{mmol}$ ) was added to an anhydrous ethanol solution ( 50 ml ) of 4-amino-1,5-dimethyl-2-phenylpyrazol-3one $(2.03 \mathrm{~g}, 10 \mathrm{mmol})$ and the mixture stirred at 350 K for 3 h under $\mathrm{N}_{2}$, giving a yellow precipitate. The product was isolated, recrystallized from ethanol, and then dried in a vacuum to give pure compound (I) in $84 \%$ yield. Yellow single crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of an ethanol solution.

Crystal data

| $\mathrm{C}_{28} \mathrm{H}_{29} \mathrm{~N}_{3} \mathrm{O}_{4}$ | $V=1235.8(13) \AA^{3}$ |
| :--- | :--- |
| $M_{r}=471.54$ | $Z=2$ |
| Triclinic, $P \overline{1}$ | $D_{x}=1.267 \mathrm{Mg} \mathrm{m}^{-3}$ |
| $a=9.804(6) \AA$ | Mo $\mathrm{A} \alpha$ radiation |
| $b=10.226(7) \AA$ | $\mu=0.09 \mathrm{~mm}^{-1}$ |
| $c=13.055(8) \AA$ | $T=294(2) \mathrm{K}$ |
| $\alpha=99.985(11)^{\circ}$ | Block, yellow |
| $\beta=90.178(11)^{\circ}$ | $0.30 \times 0.24 \times 0.18 \mathrm{~mm}$ |

$\gamma=106.223(10)^{\circ}$

## Data collection

Bruker SMART APEX CCD areadetector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.961, T_{\text {max }}=0.985$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.045$
$w R\left(F^{2}\right)=0.130$
$S=1.01$
4325 reflections
319 parameters
H-atom parameters constrained

Table 1
Hydrogen-bond geometry ( $\left({ }^{\circ},{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| C17-H17 $\cdots$ O4 | 0.93 | 2.42 | $3.049(3)$ | 125 |
| C27-H27 $\mathrm{O}^{\mathrm{i}}$ |  | 0.93 | 2.56 | $3.341(3)$ |

Symmetry code: (i) $-x+1,-y-1,-z$.
H atoms were included at calculated positions and refined using a riding model. Constrained bond lengths and $U_{\text {iso }}(\mathrm{H})$ parameters: $0.93 \AA$ and $1.2 U_{\text {eq }}(\mathrm{C})$ for aromatic, $0.97 \AA$ and $1.2 U_{\text {eq }}(\mathrm{C})$ for methylene, $0.96 \AA$ and $1.5 U_{\mathrm{eq}}(\mathrm{C})$ for methyl H atoms. The $U^{i j}$ components of atom C12 were restrained to approximate isotropic behaviour.

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve


Figure 1
The structure of (I), with displacement ellipsoids for non-H atoms drawn at the $30 \%$ probability level.


Figure 2
A partial packing diagram for (I), with hydrogen bonds shown as dashed lines.
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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