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5-(4-Methoxyphenyl)-1-phenylpyrazolidin-3-one

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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(\text{C-C}) = 0.003 \text{ Å}$ R factor = 0.048 wR factor = 0.162Data-to-parameter ratio = 14.4

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

The title compound, $C_{16}H_{16}N_2O_2$, was synthesized by the reaction of ethyl 3-(4-methylphenyl)acrylate and phenylhydrazine. There are intermolecular $N-H\cdots O$ hydrogen bonds and $C-H\cdots \pi$ interactions.

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Comment

Pyrazolidin-3-one derivatives are effective medicines used in the treatment of inflammation and related disorders (Reddy & Bell, 2003), as liphoxygenase enzyme inhibitors (Brooks *et al.*, 1990), and in pesticides and herbicides (Tanaka *et al.*, 1999). We report here the crystal structure of the title compound, (I).

The molecular structure of (I) is shown in Fig. 1, and selected bond lengths and angles are given in Table 1. The pyrazolidine ring (N1/N2/C10/C9/C8) adopts a twisted conformation (Low *et al.*, 2003). The dihedral angle between the C8/C9/C10 and C8/N1/N2/C10 planes is 15.69 (19)°. In the crystal structure, molecules are linked by N-H \cdots O hydrogen bonds and C-H $\cdots\pi$ interactions (Table 2), forming a three-

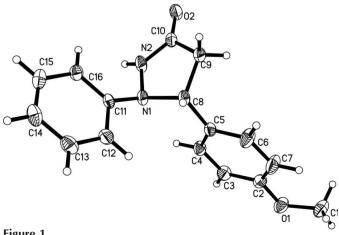


Figure 1

The molecule of (I). Displacement ellipsoids are drawn at the 30% probability level.

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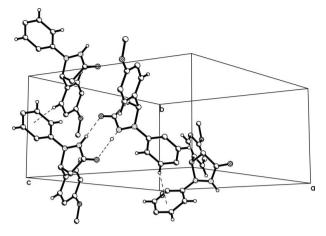


Figure 2 The crystal structure of (I). Dashed lines indicate $N-H\cdots O$ and $C-H\cdots \pi$ hydrogen bonds. H atoms not involved the hydrogen bonding are omitted for clarity.

dimensional network (Fig. 2). The phenyl ring acts as a double $C-H\cdots\pi$ hydrogen-bond acceptor, accepting one $H\cdots\pi$ bond from C6 of a methoxyphenyl group on one side and another $H\cdots\pi$ bond from C15 of a phenyl group on the other side. The dihedral angle between the two aromatic rings within the molecule is 79.5 (1)°. Owing to the presence of the carbonyl group in the pyrazolidine ring, the acidity of the H atom on C9 may be increased (Zhu *et al.*, 2004).

Experimental

To a solution of sodium (40 mmol) in anhydrous methanol (9 mol) were added ethanolamine (4 ml) and *n*-butanol (20 ml). The methanol was then removed by distillation and 3-(4-methylphenyl)-acrylic acid was added. The mixture was refluxed for 1 h at a temperature above 373 K, after which phenylhydrazine (4 ml) was added. The reactants were refluxed for a further 7 h, left to cool to room temperature, acidified with 36% acetic acid and then allowed to stand. After filtration, the filter cake was crystallized from ethyl acetate to give pure compound (I) (m.p. 435–437 K). The compound was crystallized twice by slow evaporation of an ethyl acetate solution and crystals suitable for X-ray diffraction analysis were obtained.

Crystal data

•	
$C_{16}H_{16}N_2O_2$	$D_x = 1.298 \text{ Mg m}^{-3}$
$M_r = 268.31$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 25
a = 13.151 (3) Å	reflections
b = 7.1660 (14) Å	$\theta = 1013^{\circ}$
c = 14.700 (3) Å	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 97.53 \ (3)^{\circ}$	T = 293 (2) K
$V = 1373.4 (5) \text{ Å}^3$	Block, colourless
Z = 4	$0.4 \times 0.3 \times 0.3 \text{ mm}$

Data collection

Duna concentori	
Entaf–Nonius CAD-4	$\theta_{\rm max} = 26.0^{\circ}$
diffractometer	$h = 0 \rightarrow 15$
$\omega/2\theta$ scans	$k = 0 \rightarrow 8$
2792 measured reflections	$l = -17 \rightarrow 17$
2675 independent reflections	3 standard reflections
1884 reflections with $I > 2\sigma(I)$	every 200 reflections
$R_{\rm int} = 0.026$	intensity decay: none

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.1P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.048$	where $P = (F_0^2 + 2F_c^2)/3$
$wR(F^2) = 0.162$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 1.06	$\Delta \rho_{\text{max}} = 0.22 \text{ e Å}^{-3}$
2675 reflections	$\Delta \rho_{\min} = -0.19 \text{ e Å}^{-3}$
186 parameters	Extinction correction: SHELXL97
H atoms treated by a mixture of	Extinction coefficient: 0.091 (8)
independent and constrained	
refinement	

Table 1Selected geometric parameters (Å, °).

O1-C2	1.373 (2)	N1-C11	1.425 (2)
O1-C1	1.424 (3)	N1-C8	1.500 (2)
O2-C10	1.233 (2)	N2-C10	1.332 (3)
N1-N2	1.416 (2)		
C2-O1-C1	117.39 (17)	C10-N2-N1	114.88 (16)
N2-N1-C11	113.72 (15)	C10-N2-H2A	123.8 (16)
N2-N1-C8	105.61 (14)	N1-N2-H2A	119.2 (15)
C11-N1-C8	116.17 (15)		

Table 2
Hydrogen-bond geometry (Å, °).

Cg3 is the centroid of the six-membered ring C11-C16.

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
$ \begin{array}{c} N2-H2A\cdots O2^{i} \\ C6-H6A\cdots Cg3^{ii} \\ C15-H15A\cdots Cg3^{iii} \end{array} $	0.90 (3)	1.94 (3)	2.825 (2)	169 (2)
	0.93	2.65	3.558 (3)	167
	0.93	2.86	3.737 (2)	157

Symmetry codes: (i) -x, -y + 1, -z + 1; (ii) x, y - 1, z; (iii) $-x - \frac{1}{2}$, $y + \frac{1}{2}$, $-z + \frac{3}{2}$.

The H atom on N was located in a difference electron-density map and refined isotropically. H atoms on C atoms were positioned geometrically and distances to these H atoms were set at 0.93–0.98 Å, with $U_{\rm iso}$ values constrained to be 1.5 $U_{\rm eq}$ of the parent atoms.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXL97*; software used to prepare material for publication: *SHELXL97*.

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