organic papers

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Key indicators

Single-crystal X-ray study T = 293 KMean σ (C–C) = 0.008 Å R factor = 0.090 wR factor = 0.223 Data-to-parameter ratio = 14.4

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

1-Acetyl-3,3-bis[3-(3-methoxyphenyl)-1,2,4oxadiazol-5-ylmethyl]indolin-2-one

In the title compound, $C_{30}H_{25}N_5O_6$, a derivative of oxindole, intramolecular $C-H\cdots O$ and $C-H\cdots N$ interactions help to establish the molecular conformation.

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Comment

Oxindole derivatives show many important biologial effects, such as anti-inflammatory (Kadin *et al.*, 1986) and anticonvulsant (Valenta *et al.*, 1990) properties. As part of our studies of these compounds, we have recently reported the synthesis and structure of 1-acetyl-3,3-bis[(3-(2-methylphenyl)-1,2,4-oxadiazol-5-ylmethyl]indolin-2-one, (II) (Yan *et al.*, 2006). We report here the structure of the title compound, (I) (Fig. 1), a close analogue of (II), in which the 2-methylphenyl group is replaced by a 3-methoxyphenyl group.



The indanone ring system in (I) is planar and the acetyl group at N3 is twisted from it by $1.4 (2)^{\circ}$. The dihedral angle between the N4/C22/O6/N5/C23 and C24–C29 planes is 6.7 (1)° and that between the N2/C8/N1/O2/C9 and C2–C7 planes is 2.8 (2)°. Three acute intramolecular C–H···N and C–H···O hydrogen bonds (Table 1) may help to establish the molecular conformation of (I).

Experimental

N-Acetyl-2-indolinone (20 mmol) was dissolved in acetone (40 ml) and potassium carbonate (60 mmol) was added in one portion. 5-Chloromethyl-3-(3-methoxyphenyl)-1,2,4-oxadiazole (40 mmol) in acetone (40 ml) was added to this mixture. The resulting mixture was refluxed for 72 h. After cooling and filtering, the crude title compound was obtained. The pure compound was obstained by recrystallization from a mixture of ethyl acetate (4 ml) and

© 2006 International Union of Crystallography All rights reserved petrolum ether (8 ml). Crystals of (I) suitable for X-ray diffraction analysis were obstained by slow evaporation of an ethanol solution.

V = 1349.3 (6) Å³

 $D_x = 1.358 \text{ Mg m}^{-3}$ Mo *K* α radiation

 $\mu = 0.10 \text{ mm}^{-1}$

T = 293 (2) K

 $R_{\rm int} = 0.017$

 $\theta_{\rm max} = 26.0^\circ$

+ 2P]

 $(\Delta/\sigma)_{\rm max} = 0.001$

 $\Delta \rho_{\rm max} = 0.44 \text{ e } \text{\AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.60 \text{ e } \text{\AA}^{-3}$

Block, colourless

 $0.30 \times 0.20 \times 0.10 \text{ mm}$

3 standard reflections

every 200 reflections

intensity decay: none

 $w = 1/[\sigma^2(F_o^2) + (0.06P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

Extinction correction: SHELXL97

Extinction coefficient: 0.016 (2)

5289 independent reflections

2900 reflections with $I > 2\sigma(I)$

Z = 2

Crystal data

 $\begin{array}{l} C_{30}H_{25}N_5O_6\\ M_r=551.55\\ \text{Triclinic, }P\overline{1}\\ a=8.6300\ (17)\ \mathring{A}\\ b=11.866\ (2)\ \mathring{A}\\ c=14.616\ (3)\ \mathring{A}\\ \alpha=108.95\ (3)^\circ\\ \beta=99.02\ (3)^\circ\\ \gamma=101.17\ (3)^\circ\end{array}$

Data collection

Enraf–Nonius CAD-4 diffractometer $\omega/2\theta$ scans Absorption correction: ψ scan (North *et al.*, 1968) $T_{\min} = 0.977, T_{\max} = 0.990$ 5644 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.090$ $wR(F^2) = 0.223$ S = 1.035289 reflections 365 parameters H-atom parameters constrained

Table 1

Hydrogen-bond geometry (Å, $^\circ).$

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
C7—H7A···N2	0.93	2.59	2.910 (7)	101
C16—H16A···O4	0.93	2.27	2.805 (7)	116
C25—H25A···N4	0.93	2.52	2.858 (6)	102



Figure 1

A view of the molecular structure of (I), showing displacement ellipsoids at the 30% probability level (arbitrary spheres for the H atoms).

All H atoms were positioned geometrically (C-H = 0.93–0.97 Å) and refined as riding, with $U_{iso}(H) = 1.2$ or $1.5U_{eq}(C)$.

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, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Siemens, 1996); software used to prepare material for publication: *SHELXL97*.

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