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Key indicators

Single-crystal X-ray study
 $T = 296$ K
 Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.041
 wR factor = 0.121
 Data-to-parameter ratio = 13.9

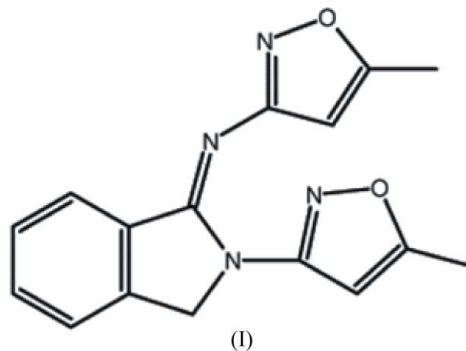
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

(5-Methylisoxazol-3-yl)[2-(5-methylisoxazol-3-yl)-2,3-dihydro-1H-isoindol-1-ylidene]amine

The title compound, $\text{C}_{16}\text{H}_{14}\text{N}_4\text{O}_2$, was synthesized by the reaction of *o*-phthalaldehyde and 5-methyl-4,5-dihydroisoxazol-3-ylamine in dichloromethane. The conformation of the molecular structure may be influenced by a weak intramolecular $\text{C}-\text{H}\cdots\text{N}$ interaction and weak $\text{C}-\text{H}\cdots\pi$ interactions, and $\pi-\pi$ stacking interactions may influence the crystal packing.

Comment

A large number of isoindoline derivatives such as staurosporine, indoprofen, and pazinaclone have been reported to possess biological activities (Takahashi & Hatanaka, 1997; Kundu *et al.*, 2001; Olmo *et al.*, 2003; Takahashi *et al.*, 2004; Cul *et al.*, 2004). Compound (I) shows antifungal activity against four species: *Chrysosporium tropicum*, *Fusarium oxysporum*, *Geotrichum candidum* and *Trichophyton rubrum* (Al-Shihry, 2005).



The molecular structure of (I) is shown in Fig. 1. All geometric parameters are in normal ranges (Allen *et al.*, 1987). Each of the individual rings is essentially planar. The isoindole system *A* (N1/C1–C8) is almost planar, with maximum deviations of -0.020 (2), 0.027 (2) and -0.024 (1) Å for atoms C5, C7 and C8, respectively. The five-membered isoxazole rings *B* (O1/N2/C9–C11) and *C* (O2/N4/C13–C15) make a dihedral angle of 75.35 (9)° with one another. The dihedral angles between rings *A/B* and *A/C* are 2.72 (7) and 76.59 (7)°, respectively. A weak intramolecular $\text{C}-\text{H}\cdots\text{N}$ hydrogen bond (Table 1) may influence the conformation of the molecular structure.

A packing diagram of (I) is shown in Fig. 2. In the crystal structure, significant $\pi-\pi$ stacking interactions exist between five-membered rings, where $\text{Cg}1\cdots\text{Cg}3(1-x, 1-y, 1-z) = 3.6285$ (9) Å (Cg1 and Cg3 are the centroids of atoms O1/N2/C9–C11 and N1/C1/C6–C8) and the perpendicular distance is 3.49 (9) Å. These and three weak intermolecular $\text{C}-\text{H}\cdots\pi(\text{arene})$ interactions may influence the crystal packing.

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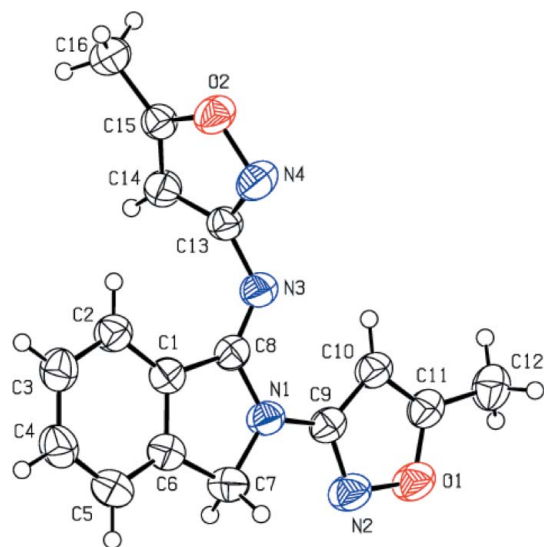


Figure 1
The molecular structure of (I), showing 50% probability displacement ellipsoids and H atoms drawn as small spheres.

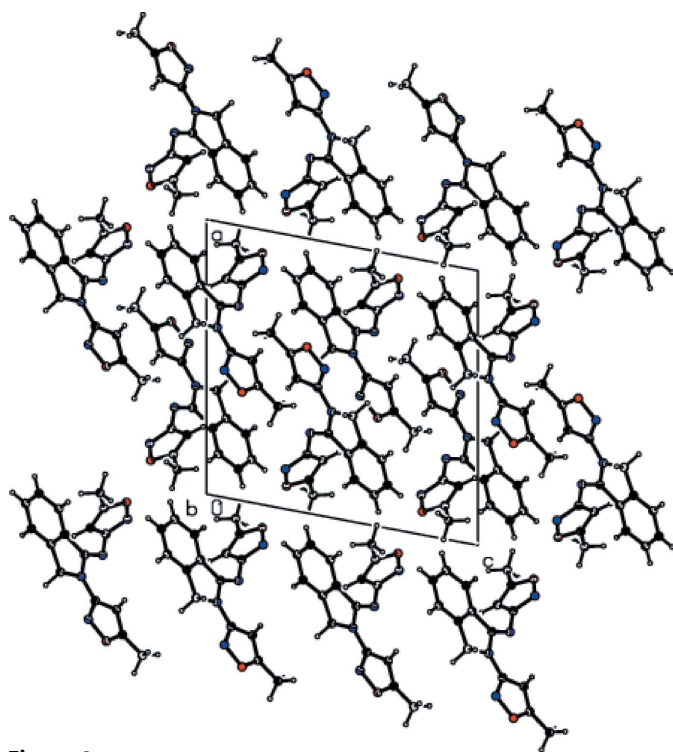


Figure 2
A view of the packing of (I).

Experimental

Compound (I) was obtained by the reaction of a stirred solution of *o*-phthalaldehyde (0.5 g, 3.73 mmol) in dichloromethane (20 ml) with a solution of 5-methyl-4,5-dihydroisoxazol-3-ylamine (0.73 g, 7.44 mmol) in dichloromethane (20 ml) (Al-Shihry, 2005). Yellow single crystals of (I) suitable for X-ray crystallographic analysis were obtained by slow evaporation of the reaction mixture (yield: 0.75 g, 68%; m.p. 464 K).

Crystal data

$C_{16}H_{14}N_4O_2$
 $M_r = 294.31$
Monoclinic, $P2_1/c$
 $a = 13.5698$ (8) Å
 $b = 7.8257$ (6) Å
 $c = 13.6527$ (8) Å
 $\beta = 100.690$ (4)°
 $V = 1424.66$ (16) Å³

$Z = 4$
 $D_x = 1.372$ Mg m⁻³
Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 296$ K
Prism with corners rounded,
colorless
 $0.44 \times 0.36 \times 0.29$ mm

Data collection

Stoe IPDS-2 diffractometer
 ω scans
Absorption correction: integration
(*X-RED32*; Stoe & Cie, 2002)
 $T_{\min} = 0.960$, $T_{\max} = 0.973$

11079 measured reflections
2807 independent reflections
2318 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.095$
 $\theta_{\max} = 26.0^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.121$
 $S = 1.07$
2807 reflections
202 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0573P)^2 + 0.1431P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.18$ e Å⁻³
 $\Delta\rho_{\min} = -0.16$ e Å⁻³
Extinction correction: *SHELXL97*
Extinction coefficient: 0.026 (3)

Table 1

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>
C10—H10...N3	0.93	2.48	2.920 (2)	109
C2—H2...Cg2	0.93	2.88	3.6348 (17)	139
C12—H12A...Cg4 ⁱ	0.96	2.96	3.5516 (18)	121
C16—H16B...Cg4 ⁱⁱ	0.96	2.74	3.5597 (18)	144

Symmetry codes: (i) $-x+1, -y+1, -z+1$; (ii) $x, y-1, z$. Cg2 and Cg4 are the centroids of atoms O2/N4/C13–C15 and C1–C6, respectively.

H atoms were placed in calculated positions and refined as riding on their parent atoms, with C—H = 0.96 (CH₃), 0.97 (CH₂) or 0.93 Å (CH), and with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{CH}_3)$ or $1.2U_{\text{eq}}(\text{CH}_2, \text{CH})$.

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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