

1-[*(E*)-(3,4-Dimethylisoxazol-5-yl)imino-methyl]-2-naphthol

Hoong-Kun Fun,^{a*}‡ Madhukar Hemamalini,^a Abdullah M. Asiri^b§ and Salman A. Khan^b

^aX-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, and ^bDepartment of Chemistry, Faculty of Science, King Abdul Aziz University, Jeddah, Saudi Arabia

Correspondence e-mail: hkfun@usm.my

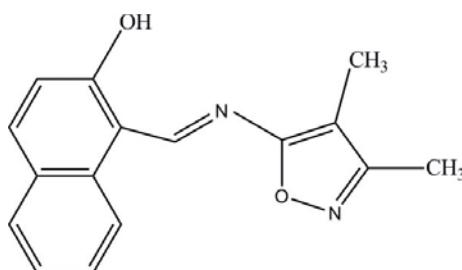
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(C-C) = 0.002$ Å; R factor = 0.046; wR factor = 0.134; data-to-parameter ratio = 15.6.

The title Schiff base compound, $C_{16}H_{14}N_2O_2$, has been synthesized by the reaction of 5-amino-3,4-dimethylisoxazole and 2-hydroxy-1-naphthaldehyde. The dihedral angle between the isoxazole ring and the naphthyl ring system is $3.29(7)^\circ$. The molecule adopts an *E* configuration about the central $C\equiv N$ double bond. Intramolecular O—H···N hydrogen bonding generates an *S*(6) ring motif. In the crystal structure, $\pi-\pi$ interactions are observed involving the isoxazole ring and the substituted benzene ring of the naphthyl unit, with centroid–centroid distances of $3.5200(10)$ Å.

Related literature

For related background and the biological activity of isoxazol, see: Howell & Kimmel (2008); Bartlett & Schleyerbach (1985); Lamani *et al.* (2009); Jayashankar *et al.* (2009). For related structures, see: Alvarez-Thon *et al.* (2006); Tahir *et al.* (2008); Shad *et al.* (2008); Fun *et al.* (2010). For details of hydrogen-bond motifs, see: Bernstein *et al.* (1995). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986).



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Experimental

Crystal data

$C_{16}H_{14}N_2O_2$	$V = 1281.17(16)$ Å ³
$M_r = 266.29$	$Z = 4$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 7.5250(6)$ Å	$\mu = 0.09$ mm ⁻¹
$b = 15.4643(12)$ Å	$T = 100$ K
$c = 12.3982(7)$ Å	$0.79 \times 0.06 \times 0.05$ mm
$\beta = 117.377(4)^\circ$	

Data collection

Bruker APEX DUO CCD area-detector diffractometer	16577 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2009)	3704 independent reflections
$T_{\min} = 0.930$, $T_{\max} = 0.996$	2843 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.134$	$\Delta\rho_{\max} = 0.45$ e Å ⁻³
$S = 1.05$	$\Delta\rho_{\min} = -0.23$ e Å ⁻³
3704 reflections	
237 parameters	

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1O1···N1	0.97 (2)	1.66 (3)	2.5471 (15)	150 (2)

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXTL* and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SJ2761).

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