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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.056 wR factor = 0.144 Data-to-parameter ratio = 16.3

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

The title compound, C₂₀H₁₆O₃, was synthesized from 3hvdroxy-4-methoxybenzaldehvde and 2-(prop-1-en-2-yl)naphthalene in aqueous ethanol. The naphthalene ring system makes a dihedral angle of 15.8 (4) Å with the plane of the benzene ring. In the crystal structure, molecules are linked into centrosymmetric dimers by $O-H \cdots O$ hydrogen bonds.

(E)-3-(3-Hydroxy-4-methoxyphenyl)-1-(2-naphthyl)-

Comment

prop-2-en-1-one

(*E*)-3-(3-Hydroxy-4-methoxyphenyl1-(2-naphthyl)prop-2-en-1-one, (I), is a valuable pharmaceutical intermediate (Jardine et al., 1980; Issell et al., 1984; Stahelin et al., 1989), which was synthesized by a new method from 3-hydroxy-4-methoxybenzaldehyde and 2-(prop-1-en-2-yl)naphthalene. Its structure is reported here (Fig. 1).



The dihedral angle formed between the naphthalene ring system and the plane of the benzene ring is $15.8 (4)^{\circ}$. The methoxyl group lies slightly out of the plane of the benzene ring, with atoms O2 and C7 deviating from the C1-C6 ring plane by 0.061 (2) and 0.295 (3) Å, respectively. In the crystal structure, molecules are linked into centrosymmetric dimers by $O-H \cdots O$ hydrogen bonds (Fig. 2 and Table 1).

Experimental

A mixture of 3-hydroxy-4-methoxybenzaldehyde (5 mmol) and 2-(prop-1-en-2-yl)naphthalene (10 mmol) was dissolved in aqueous ethanol (100 ml, 60% v/v). A solution of KOH (4.0 ml, 7.1 mmol) was added slowly with stirring at 323 K for 5 h. The resulting mixture was neutralized with 2 M HCl, the precipitate filtered off, washed with water, and crystallized from dichloromethane-petroleum ether (4:1 v/



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The molecular structure of (I), shown with 30% probability displacement ellipsoids (arbitrary spheres for H atoms).

Received 10 November 2006 Accepted 15 November 2006 v) to obtain (I) (m.p. 442–443 K). Crystals suitable for X-ray analysis were grown by slow evaporation of a petroleum ether–dichloromethane (4:1 v/v) solution at room temperature over a period of 17 d.

Z = 8

 $D_{\rm v} = 1.360 {\rm Mg m}^{-3}$

Mo $K\alpha$ radiation

 $\mu = 0.09 \text{ mm}^{-1}$

T = 293 (2) K

 $\begin{aligned} R_{\rm int} &= 0.058\\ \theta_{\rm max} &= 27.8^\circ \end{aligned}$

Prism, colorless

 $0.22 \times 0.16 \times 0.14 \text{ mm}$

25516 measured reflections

3509 independent reflections

3105 reflections with $I > 2\sigma(I)$

Crystal data

 $\begin{array}{l} C_{20}H_{16}O_{3} \\ M_{r} = 304.33 \\ \text{Orthorhombic, } Pbca \\ a = 14.7417 \ (11) \\ \text{\AA} \\ b = 6.1208 \ (4) \\ \text{\AA} \\ c = 32.941 \ (3) \\ \text{\AA} \\ V = 2972.3 \ (4) \\ \text{\AA}^{3} \end{array}$

Data collection

Rigaku Saturn-70 diffractometer ω scans Absorption correction: multi-scan (Jacobson, 1998) $T_{min} = 0.980, T_{max} = 0.987$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0656P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.056$ + 0.9426P] $wR(F^2) = 0.144$ where $P = (F_0^2 + 2F_c^2)/3$ S = 1.16 $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.24 \text{ e} \text{ Å}^{-3}$ 3509 reflections $\Delta \rho_{\rm min} = -0.18 \text{ e } \text{\AA}^{-3}$ 215 parameters H atoms treated by a mixture of Extinction correction: SHELXL97 Extinction coefficient: 0.0075 (13) independent and constrained refinement

Table 1

TT		(Å	0)
Hydrogen-bond	geometry	(A,	·).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$O1 - H1 \cdots O3^i$	0.94 (3)	1.74 (3)	2.6795 (18)	176 (2)
Commentation and as (i)		1.1		

Symmetry code: (i) -x, -y + 1, -z + 1.

The H atom of the OH group was located in a difference Fourier map and refined freely with an isotropic displacement parameter. Other H atoms were included using the riding-model approximation,





with C-H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic, and C-H = 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms.

Data collection: *CrystalClear* (Rigaku/MSC, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *CrystalStructure* (Rigaku/MSC, 2005).

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