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Key indicators

Single-crystal X-ray study T = 296 KMean $\sigma(C-C) = 0.004 \text{ Å}$ R factor = 0.060 wR factor = 0.171 Data-to-parameter ratio = 13.1

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. 1-(1-Hydroxy-2-naphthyl)ethanone

The molecule of the title compound, $C_{12}H_{10}O_2$, is planar, with an intramolecular $O-H\cdots O$ hydrogen bond between the hydroxy group and the carbonyl O atom. The structure is stabilized by intermolecular $C-H\cdots \pi$ and offset $\pi-\pi$ stacking interactions.

Comment

Some acetylnaphthalene derivatives have potential as pharmacologically active compounds (Walker *et al.*, 1981; Çalış *et al.*, 1988; Gupta & Singh, 1991; Karakurt *et al.*, 2001). In this study, we aimed to investigate the intra- and/or intermolecular interactions and the conformation of the title compound, (I), by X-ray crystallography.



In (I), the hydrogen-bonded ring A (O1/C2/C3/C12/O2/H2), and aromatic rings B (C3/C4/C5/C6/C11/C12) and C (C6/C7/ C8/C9/C10/C11), are planar, with dihedral angles of 1.30 (1)° for A/B, 1.18 (1)° for A/C and 1.15 (1)° for B/C (Fig. 1). A significant intramolecular interaction is noted, involving hydroxy atom H1 and carbonyl atom O1, such that a sixmembered ring is formed (Table 1). The bond lengths and angles are in normal ranges, and comparable with those in 2bromo-1-(1-hydroxynaphthalen-2-yl)ethanone (Köysal *et al.*, 2004) and in our previous work on *o*-hydroxyaldehydes (Odabaşoğlu *et al.*, 2006; Odabaşoğlu & Büyükgüngör, 2006). The C2–O1 bond distance in (I) is also consistent with the value of the C=O double bond in carbonyl compounds (Loudon, 2002).

The structure of (I) is stabilized by $C-H\cdots\pi$ intermolecular interactions (Table 1, Fig. 2). There are also weak offset $\pi-\pi$ stacking interactions between the *B* and *C* rings of symmetry-related molecules, with a plane-to-plane separation of 3.481 Å and a centroid-to-centroid distance of 3.821 Å, resulting in an angle between the ring normal and the centroid-to-centroid vector of 24° (Fig. 2).

Experimental

1-(1-Hydroxynaphthalen-2-yl)ethanone, purchased from Avocado Research Chemicals Ltd. (99%), was dissolved in dimethylsulfoxide (393 K), and well shaped crystals of (I) were obtained from this hot solution (m.p. 372-373 K).

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Crystal data

 $C_{12}H_{10}O_2$ $M_r = 186.20$ Monoclinic, $P2_1/c$ a = 7.6259 (10) Å b = 7.0348 (7) Å c = 18.755 (3) Å $\beta = 111.502 (10)^{\circ}$ $V = 936.1 (2) \text{ Å}^3$ Z = 4

Data collection

Stoe IPDS-2 diffractometer
ω scans
Absorption correction: integration
(X-RED32; Stoe & Cie, 2002)
$T_{\min} = 0.548, T_{\max} = 0.946$
10250 measured reflections
1692 independent reflections

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.060$	$w = 1/[\sigma^2(F_o^2) + (0.0964P)^2]$
$wR(F^2) = 0.172$	where $P = (F_0^2 + 2F_c^2)/3$
S = 0.99	$(\Delta/\sigma)_{\rm max} < 0.001$
1692 reflections	$\Delta \rho_{\rm max} = 0.11 \text{ e } \text{\AA}^{-3}$
129 parameters	$\Delta \rho_{\rm min} = -0.14 \text{ e } \text{\AA}^{-3}$

 $D_x = 1.321 \text{ Mg m}^{-3}$

Cell parameters from 10250

Prismatic stick, pale yellow

1015 reflections with $I > 2\sigma(I)$

 $0.72 \times 0.34 \times 0.14$ mm

Mo $K\alpha$ radiation

reflections

 $\theta = 2.3-27.1^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$

T = 296 (2) K

 $R_{\rm int} = 0.100$

 $\theta_{\rm max} = 25.3^\circ$

 $h = -9 \rightarrow 9$

 $k = -8 \rightarrow 8$

 $l = -22 \rightarrow 22$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots$
$C1-H1A\cdots Cg2^i$	0.96	2.79	3.593 (3)	142
O2-H2···O1	0.82	1.80	2.524 (3)	146

Symmetry code: (i) -x + 2, -y + 1, -z + 1. Cg2 is the centroid of atoms C6–C11

All H atoms were introduced in calculated positions and treated as riding on their parent atoms, with O–H = 0.82 Å, and C–H = 0.93 Å (C_{aromatic}) and 0.96 Å (CH₃), and with U_{iso} (H) = $1.2U_{eq}$ (C_{aromatic},O) or U_{iso} (H) = $1.5U_{eq}$ (CH₃).

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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Figure 1

A view of (I), with the atomic numbering scheme. The intramolecular hydrogen bond is shown as a dashed line. Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

A partial packing diagram for (I). The intramolecular hydrogen bonds are shown as short dashed lines, and the $C-H\cdots\pi$ and $\pi-\pi$ stacking interactions are shown as long dashed lines. H atoms not involved in hydrogen bonding have been omitted for clarity.

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