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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.006 Å R factor = 0.039 wR factor = 0.113 Data-to-parameter ratio = 10.5

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

N-(2-Hydroxy-1-naphthylmethyl)-(S)-proline

The title compound [systematic name: (2S)-1-(2-hydroxy-1naphthylmethyl)pyrrolidinio-2-carboxylate], C₁₆H₁₇NO₃, a derivative of L-proline, was synthesized *via* the Mannich reaction. The X-ray crystal analysis reveals that the pyrrolidine ring adopts an envelope conformation, with the flap C atom deviating by 0.631 (9) Å from the plane defined by the other four atoms. Received 5 January 2005 Accepted 4 March 2005 Online 18 March 2005

Comment

In order to synthesize chiral amino alcohols, which can be used for asymmetric Michael addition (Yamakoshi *et al.*, 1999; Sasai *et al.*, 1993), the title compound, (I), was prepared *via* a Mannich reaction with L-proline, 2-naphthol and formaldehyde. The original *S* configuration of the chiral centre at C2 is assumed not to change during the preparation, as shown in Fig. 1. Selected bond distances and angles are given in Table 1. The pyrrolidine ring may adopt different conformations (Stavropoulos *et al.*, 1990; Puliti *et al.*, 2000; Zheng & Li, 2003). Here it adopts an envelope conformation, with atom C4 deviating by 0.631 (9) Å from the plane defined by atoms N1, C2, C3 and C5.



Conjugation between bonds O1-C1 and O2-C1 is observed, with bond distances of 1.222 (3) and 1.239 (3) Å, respectively. The conformation and crystal structure of the title compound are stabilized by N-H···O and O-H···O hydrogen bonds (see Table 2).

Experimental

The title compound was prepared according to the procedure of Zheng & Li (2003). Suitable crystals were obtained by slow evaporation of a mixture of chloroform and cyclohexane (m.p. 455–457 K). ¹H NMR (CD₃OD, p.p.m.): δ 1.88–1.93 (*m*, 1H), 2.07–2.19 (*m*, 2H), 2.47–2.52 (*m*, 1H), 3.29–3.38 (*m*, 1H), 3.46–3.51 (*m*, 1H), 4.05–4.09 (*m*, 1H), 4.80–4.91 (*m*, 4H), 7.19 (*d*, 1H, *J* = 8 Hz), 7.35 (*t*, 1H, *J* = 8 Hz), 7.57 (*t*, 1H, *J* = 8 Hz), 7.79–7.85 (*m*, 2H), 8.20 (*d*, 1H, J = 8 Hz).

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Figure 1

A view of the molecule of (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.

Mo Ka radiation

Cell parameters from 1714 reflections $\theta = 2.6-22.2^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 293 (2) K Block, colourless $0.24 \times 0.16 \times 0.14 \text{ mm}$

1980 independent reflections

 $\begin{aligned} R_{\rm int} &= 0.029 \\ \theta_{\rm max} &= 28.0^\circ \end{aligned}$

 $h = -7 \rightarrow 7$

 $k=-15\rightarrow 15$

 $l = -26 \rightarrow 19$

1329 reflections with $I > 2\sigma(I)$

Crystal data

$C_{16}H_{17}NO_3$
$M_r = 271.31$
Orthorhombic, P2 ₁ 2 ₁ 2 ₁
a = 5.8037 (7) Å
b = 12.0508 (15) Å
c = 20.160 (3) Å
V = 1410.0 (3) Å ³
Z = 4
$D_x = 1.278 \text{ Mg m}^{-3}$
c .

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 1997) $T_{\min} = 0.964$, $T_{\max} = 0.988$ 9621 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.039$ $wR(F^2) = 0.113$ S = 1.04 1980 reflections 188 parameters H atoms treated by a mixture of independent and constrained	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0446P)^{2} + 0.1866P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.18 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.15 \text{ e} \text{ Å}^{-3}$ Extinction correction: none
refinement	

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Sel	ected	geometric	parameters	(A,	°)	١.
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N1-H1	0.90(1)	O2-C1	1.239 (3)
O1-C1	1.222 (3)	O3-H3	0.86 (1)
C8-O3-H3	112 (2)	C2-C3-C4	103.2 (2)
O1-C1-O2	127.6 (3)	N1-C5-C4	103.3 (2)
O1-C1-C2	117.8 (2)	C7-C6-N1	111.8 (2)
O2-C1-C2	114.5 (2)		
C5-N1-C2-C3	0.2 (3)	O2-C1-C2-N1	174.2 (3)
C6-N1-C2-C3	126.6 (2)	01-C1-C2-C3	106.4 (3)
O1-C1-C2-N1	-9.0 (4)	O2-C1-C2-C3	-70.5 (3)



Figure 2

The crystal structure of (I), viewed along the b axis. Dashed lines indicate hydrogen-bond interactions.

Table 2Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
N1−H1···O1	0.90(1)	1.99 (2)	2.599 (3)	123 (2)
$N1 - H1 \cdots O3$	0.90(1)	2.28 (2)	2.892 (3)	124 (2)
$O3 - H3 \cdots O2^i$	0.86(1)	1.73 (1)	2.586 (3)	175 (4)

Symmetry code: (i) -x + 1, $y + \frac{1}{2}$, $-z + \frac{3}{2}$.

All N- and O-bound H atoms were initially located in a difference Fourier map and were included in the refinement with restrained N– H and O–H distances of 0.85 and 0.90 Å. All other H atoms were positioned geometrically and refined using a riding model, with C–H = 0.93–0.98 Å and $U_{iso}(H) = 1.2U_{eq}(C)$. In the absence of significant anomalous dispersion effects, Friedel pairs were merged prior to refinement. The large anisotropic displacement parameters of some atoms in the naphthalene fused-ring system indicate either high thermal motion or possible unresolved disorder.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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