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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.003 \text{ Å}$ R factor = 0.043 wR factor = 0.112 Data-to-parameter ratio = 13.8

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

3-(1,3-Benzothiazol-2-yl)-2-naphthol

The molecule of the title compound, $C_{17}H_{11}NOS$, is planar, the dihedral angle between the benzothiazole and 2-naphthol systems being 3.44 (6)°. The face-to-face separation of 3.48 (2) Å indicates π - π stacking between parallel benzothiazole systems.

Comment

The title compound, (I), is an intermediate of photochromic compounds (Deligeorgiev *et al.*, 2002). We report here the molecular structure of (I).



The molecular structure of (I) is shown in Fig. 1. The molecule is planar, the dihedral angle between the benzothiazole and 2-naphthol mean planes being $3.44 (6)^{\circ}$. Intramolecular hydrogen bonding occurs between the benzothiazole and naphthol ring systems (Table 1).

The parallel benzothiazole systems partially overlap each other (Fig. 2). The face-to-face separation of 3.48 (2) Å indicates the existence of π - π stacking.

Experimental

The title compound was prepared according to the procedure described by Deligeorgiev *et al.* (2002). It was synthesized by reaction of 2-hydroxy-3-naphthalenecarboxylic acid and 2-aminothiophenol in boiling toluene. Single crystals of (I) were obtained by recrystallization from toluene solution.



Figure 1

The molecular structure of (I), with 40% probability displacement ellipsoids (arbitrary spheres for H atoms). Dashed lines indicate hydrogen bonding.

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Figure 2 The π - π stacking in (I) [symmetry code: (i) 1 - x, -y, 2 - z].

Crystal data

 $C_{17}H_{11}NOS$ $M_r = 277.33$ Monoclinic, $P2_1/n$ a = 6.3634 (9) Å b = 8.5293 (12) Å c = 23.588 (3) Å $\beta = 95.120 (3)^{\circ}$ $V = 1275.1 (3) Å^{3}$

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2002) $T_{\min} = 0.931, T_{\max} = 0.945$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.043$ $wR(F^2) = 0.112$ S = 0.862497 reflections 181 parameters Z = 4 D_x = 1.445 Mg m⁻³ Mo K α radiation μ = 0.25 mm⁻¹ T = 295 (2) K Prism, pale yellow 0.34 × 0.27 × 0.22 mm

6746 measured reflections 2497 independent reflections 1755 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.051$ $\theta_{\text{max}} = 26.0^{\circ}$

H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0699P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} = 0.002$ $\Delta\rho_{max} = 0.30 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{min} = -0.15 \text{ e } \text{\AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$O1-H1\cdots N1$	0.82	1.88	2.612 (2)	149

H atoms bonded to C atoms were positioned geometrically and treated as riding, with C–H = 0.93 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$. The hydroxy H atom was located in a difference Fourier map and refined as riding in its as-found relative position (O–H = 0.82 Å), with $U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}({\rm O})$.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; 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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