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#### **Key indicators**

Single-crystal X-ray study T = 295 KMean  $\sigma(\text{C}-\text{C}) = 0.007 \text{ Å}$  R factor = 0.076 wR factor = 0.215 Data-to-parameter ratio = 14.7

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# Naphthalene-2,7-diol-1,2,4-triazole (2/1)

In the crystal structure of the title compound,  $2C_{10}H_8O_{2}$ .  $C_2H_3N_3$ , intermolecular N-H···O, O-H···N and O-H···O hydrogen bonds connect naphthalene-2,7-diol molecules and 1,2,4-tridazole molecules into a linear ribbon motif. Received 7 August 2006 Accepted 4 September 2006

## Comment

Aromatic alcohols are generally not sufficiently acidic to protonate bases; however, naphthalene-2,7-diol represents an exception as it affords some cocrystals with bases, as noted from the reported crystal structures of 10-methylisoalloxazinium bromide hydrate (Langhoff & Fritchie, 1970), cefadroxil heptahydrate (Kemperman *et al.*, 2000) and diaza-18crown-6 (Watson *et al.*, 1989). In the present study, a 2:1 cocrystal, (I), was obtained when naphthalene-2,7-diol was treated with an equimolar quantity of 1,2,4-triazole. We find it interesting that the title structure is not a 1:1 cocrystal as the reactants were mixed in an equimolar ratio, as was the case for two previously reported crystal structures (Wang & Tang, 2006; Wang, Tang & Wan, 2006).



In the crystal structure, a ribbon motif is formed *via* intermolecular  $N-H\cdots O$ ,  $O-H\cdots O$  and  $O-H\cdots N$  hydrogen bonds. In detail, the repeat unit of this extended ribbon consists of four  $O-H\cdots O$  hydrogen-bonded naphthalene-2,7diol molecules which are, in turn,  $O-H\cdots N$  and  $N-H\cdots O$ hydrogen bonded to two 1,2,4-triazole molecules. There are no hydrogen bonds between triazole molecules (Fig. 2 and Table 2).



**Figure 1** View of the asymmetric unit of (I), shown with displacement ellipsoids at the 50% probability level.

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## Experimental

Naphthalene-2,7-diol (80 mg, 0.5 mmol) dissolved in methanol (5 ml) was treated with 1,2,4-triazole (35 mg, 0.5 mmol) dissolved in methanol (5 ml). After a few days, colorless bar-shaped crystals separated from the solution. Elemental analysis found: C 62.71, H 4.86, N 18.36%; calculated: C 62.87, H 4.84, N 18.33%.

Z = 4

 $D_x = 1.328 \text{ Mg m}^{-3}$ 

 $0.4 \times 0.3 \times 0.3$  mm

Mo  $K\alpha$  radiation  $\mu = 0.09 \text{ mm}^{-1}$ T = 295 (2) K

Bar, colorless

### Crystal data

$2C_{10}H_8O_2 \cdot C_2H_3N_3$
$M_r = 389.40$
Monoclinic, $P2_1/c$
$a = 20.754 (4) \text{\AA}$
b = 5.890(1) Å
c = 16.273 (4)  Å
$\beta = 101.682 \ (6)^{\circ}$
$V = 1948.0(7) \text{ Å}^3$

#### Data collection

Bruker SMART 1K area-detector	3918 independent reflections
diffractometer	1489 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\rm int} = 0.090$
Absorption correction: none	$\theta_{\rm max} = 26.3^{\circ}$
8911 measured reflections	

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0838P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.076$	where $P = (F_0^2 + 2F_c^2)/3$
$wR(F^2) = 0.215$	$(\Delta/\sigma)_{\rm max} = 0.001$
S = 0.97	$\Delta \rho_{\rm max} = 0.22 \ {\rm e} \ {\rm \AA}^{-3}$
3918 reflections	$\Delta \rho_{\rm min} = -0.19 \text{ e } \text{\AA}^{-3}$
267 parameters	Extinction correction: SHELXL97
H-atom parameters constrained	Extinction coefficient: 0.008 (2)

## Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
O1−H1 <i>o</i> ···N1	0.85	1.93	2.760 (6)	167
$O2-H2o\cdots O4^{i}$	0.85	2.02	2.806 (5)	153
O3-H30···N2	0.85	2.12	2.920 (5)	157
O4-H4o···O2 <sup>ii</sup>	0.85	1.90	2.743 (4)	172
$N3-H3n\cdots O3^{i}$	0.85	1.90	2,750 (5)	176

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



Figure 2 Part of the crystal structure of (I). Hydrogen bonds are shown as dashed lines.

The crystal did not diffract strongly and there are fewer reflections than normal which have  $I > 2\sigma(I)$ ; thismay lower the precision of the structure. H atoms were positioned geometrically (C-H = 0.93 Å, N-H = 0.85 Å and O-H = 0.85 Å), and were included in the refinement in the riding-model approximation, with  $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C,N,O})$ .

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *SHELXL97*.

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