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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.013 \text{ Å}$ R factor = 0.064 wR factor = 0.132 Data-to-parameter ratio = 12.3

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

Tetragonal polymorph of the 1:1 adduct of sulfathiazole with pyridine

The title tetragonal polymorph is one of the two sulfathiazole– pyridine adducts, $C_9H_9N_3O_2S_2\cdot C_5H_5N$, that can be formed either by crystallization from an *n*-propanol–pyridine solution, or by exposure of solid sulfathiazole to pyridine vapour. The asymmetric unit consists of a hydrogen-bonded sulfathiazole– pyridine pair. Hydrogen bonds of the form $N_{aniline}$ – $H \cdots O_{sulfonyl}$ form a three-dimensional network. Pyridine molecules linked to sulfathiazole molecules by N_{amino} – $H \cdots N_{pyridine}$ hydrogen bonds are located in the channels of the sulfathiazole framework which extend along the 4₁ axis. The angle between neighbouring pyridine rings in the channels is 55.2 (3)°. The adduct is stable in pyridine vapour, but decomposes in air under ambient conditions, giving the metastable polymorph I of sulfathiazole [Kruger & Gafner (1972). Acta Cryst. B**28**, 272–283].

Comment

Sulfathiazole gives over 100 solvates, for only a few of which crystal structures have been reported (Bingham *et al.*, 2001). Two polymorphs of the adduct of sulfathiazole with pyridine (1:1) could be crystallized from propanol–pyridine solution (Mikhailenko *et al.*, 2005). The structure of the pyramidal crystals of the tetragonal polymorph, (I), is described in the present contribution.



The pyramidal crystals of (I) have tetragonal symmetry (space group $P4_1$). The space group $P4_1$ is very rare for molecular crystals (Cambridge Structural Database; Version V5.27; Allen, 2002). Only one of the previously known solvates of sulfathiazole (Bingham *et al.*, 2001) can be

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Figure 1

The molecular structure of the 1:1 sulfathiazole adduct with pyridine in the tetragonal polymorph. Displacement ellipsoids are plotted at the 50% probability level. The dashed line indicates a hydrogen bond.



Figure 2

Parts of the structure of the tetragonal 1:1 sulfathiazole adduct with pyridine, shown in two projections. Dashed lines indicate hydrogen bonds.

supposed to be tetragonal, judging from the cell parameters.

The sulfathiazole molecules of (I) exist as imide tautomers (Fig. 1), similar to those found in five polymorphs of pure sulfathiazole (Kruger & Gafner, 1971, 1972; Babilev et al., 1987; Hughes et al., 1999), and in the monoclinic polymorph of the title adduct (Drebushchak et al., 2006). The sulfathiazolepyridine pairs are linked via an Namino-H···Npyridine hydrogen bond in both tetragonal and monoclinic polymorphs. In the tetragonal polymorph, the hydrogen-bonded sulfathiazole molecules (with thiazole and benzene rings being almost normal to each other) form a three-dimensional framework, in which pyridine guests fill the channels, which extend along the 4_1 axis. The angle between neighbouring pyridine rings in the channels is $55.2 (3)^{\circ}$. The parameters characterizing the hydrogen bonds are summarized in Table 1. The components of the crystal structure in two projections are shown in Fig. 2. The adduct can be classified as an appendage inclusion compound (Herbstein, 2005), with the host structure stabilized by the presence of the guest.

The tetragonal polymorph of the adduct is stable in pyridine vapour, but decomposes in air under ambient conditions, giving the metastable polymorph I of sulfathiazole [Kruger & Gafner, 1972; m.p. 474.8 K (Mikhailenko et al., 2005)]. It is interesting that the decomposition of all other sulfathiazole adducts has always been reported to give polymorphs II (Kruger & Gafner, 1971), III (Kruger & Gafner, 1972) and, particularly often, IV (Babilev et al., 1987), but never polymorph I (Bingham et al., 2001) of the sulfathiazole.

Experimental

The tetragonal polymorph, (I), of the 1:1 adduct of sulfathiazole with pyridine was obtained by slow evaporation of the solvent from a solution containing sulfathiazole (1.5 g), propanol (40 ml) and pyridine (20 ml). The synthesis of the polymorphs has been described in more detail by Mikhailenko et al. (2005).

Crystal data

$C_9H_9N_3O_2S_2 \cdot C_5H_5N$	$D_x = 1.412 \text{ Mg m}^{-3}$
$M_r = 334.43$	Mo $K\alpha$ radiation
Tetragonal, P4 ₁	$\mu = 0.35 \text{ mm}^{-1}$
a = 8.670 (2) Å	T = 295 (2) K
c = 20.927 (10) Å	Pyramid, colourless
V = 1573.2 (9) Å ³	$0.46 \times 0.41 \times 0.38 \text{ mm}$
Z = 4	

Data collection

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Stoe STADI-4 four-circle
  diffractometer
\omega/\theta scans
Absorption correction: \psi scan
   (X-RED; Stoe & Cie, 1997)
   T_{\min} = 0.771, T_{\max} = 0.874
2518 measured reflections
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Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.03P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.064$	+ 1.3187P]
$wR(F^2) = 0.132$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.17	$(\Delta/\sigma)_{\rm max} < 0.001$
2518 reflections	$\Delta \rho_{\rm max} = 0.23 \text{ e} \text{ Å}^{-3}$
205 parameters	$\Delta \rho_{\rm min} = -0.28 \text{ e} \text{ Å}^{-3}$
H atoms treated by a mixture of	Absolute structure: Flack (1983),
independent and constrained	with 1006 Friedel pairs
refinement	Flack parameter: 0.31 (17)

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
N3-H7···N4	0.86	1.93	2.773 (9)	165
$N1 - H1 \cdots O1^{i}$	0.86	2.12	2.976 (9)	176
$N1 - H2 \cdots O2^{ii}$	0.86	2.49	3.040 (8)	123

Symmetry codes: (i) x + 1, y, z; (ii) y + 1, -x + 2, $z - \frac{1}{4}$.

2518 independent reflections

3 standard reflections

frequency: 180 min intensity decay: 38.0%

 $R_{\rm int} = 0.021$

 $\theta_{\rm max} = 25.5^{\circ}$

1726 reflections with $I > 2\sigma(I)$

The tetragonal polymorph is not stable under ambient conditions and loses pyridine easily. Due to this instability, problems were encountered during data collection. One unique set of data was collected completely up to $\theta = 25.5^{\circ}$, but for the second equivalent set we were forced to stop data collection at $\theta = 22.5^{\circ}$, because of a strong decay in intensities due to crystal decomposition. For atom N4, U_{eq} is low compared with its neighbours, and this can be interpreted by the fact that N4 is involved in hydrogen-bond formation, in contrast with its neighbours, and therefore a slight disorder or high mobility of pyridine does not affect atom N4 very much - the molecule of pyridine is 'pinned' by atom N4 to the NH group of sulfathiazole. The amino atoms H1 and H2 were located in difference maps. The remaining H atoms were positioned geometrically at C-H = 0.93 and N-H = 0.86 Å. Atom H10 was refined freely but all other H atoms were refined using a riding model, with $U_{iso}(H) = 1.2U_{eq}$ of the carrier atom.

Data collection: *STADI4* (Stoe & Cie, 1997); cell refinement: *STADI4*; data reduction: *X-RED* (Stoe & Cie, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003) and *PowderCell* (Kraus & Nolze, 1999); software used to prepare material for publication: *SHELXL97*.

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