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Key indicators

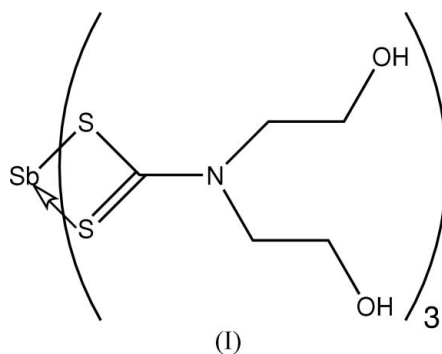
Single-crystal X-ray study
 $T = 93$ K
Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.020
 wR factor = 0.051
Data-to-parameter ratio = 19.1For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.A low-temperature refinement of tris[*N,N'*-bis(2-hydroxyethyl)dithiocarbamato- $\kappa^2\text{S,S}'$]-antimony(III)The title compound, $[\text{Sb}(\text{C}_5\text{H}_{10}\text{NO}_2\text{S}_2)_3]$, has been redetermined at 93 K, allowing a detailed analysis of the crystal packing, which is dominated by $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonding. Each terminal hydroxyl group participates in a 12-membered $[\cdots\text{O}-\text{H}]_6$ ring, leading to a three-dimensional array.

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Comment

The binary antimony(III) dithiocarbamates, $\text{Sb}(\text{S}_2\text{CNR}_2)_3$, usually associate into dimers *via* $\text{Sb}\cdots\text{S}$ interactions (Liu & Tiekink, 2005). Exceptions to this behaviour are found only when the steric bulk of the *R* groups precludes close association, such as in the example where $\text{NR}_2 = \text{N}(\text{CH}_2)_6$ (Liu & Tiekink, 2005). Indeed, steric bulk in remote substituents has been found to be an important factor in supramolecular aggregation in these and related systems (*e.g.* Lai *et al.*, 2002, 2004; Tiekink, 2003, 2006*a,b*). In the $\text{Sb}(\text{S}_2\text{CNR}_2)_3$ structures for which hydrogen-bonding functionality exists in one of the *R* groups, the same principle of steric control holds (Liu & Tiekink, 2005). However, in the case of $\text{Sb}[\text{S}_2\text{CN}(\text{CH}_2\text{CH}_2\text{OH})_2]_3$, (I), no $\text{Sb}\cdots\text{S}$ interactions were evident, despite being present in the *n*-propyl analogue $\text{Sb}[\text{S}_2\text{CN}(\text{CH}_2\text{CH}_2\text{OH})_2]_3$ (Kavounis *et al.*, 1982). In the original crystallographic analysis of (I) (Venkatachalam *et al.*, 1997), there were ambiguities in the positions of the $-\text{OH}$ groups. Here, a new analysis of (I) is reported from data measured at 93 K.



The molecular structure of (I) (Fig. 1 and Table 1) shows the expected features for compounds of this type (Liu & Tiekink, 2005). The molecule has non-crystallographic symmetry, at least in terms of the S_2CNC_2 residues, so that the S_1S_2 -dithiocarbamate ligand defines an approximate mirror plane. This ligand forms the most asymmetric chelate of the three dithiocarbamate ligands and this is reflected in the associated $\text{C}-\text{S}$ distances (Table 1). The coordination geometry is based on a pentagonal bipyramid, with atoms $\text{S}_2/\text{S}_3-\text{S}_6$ defining a distorted pentagon and with atom S_1 and the lone pair of electrons occupying axial positions. This conformation allows

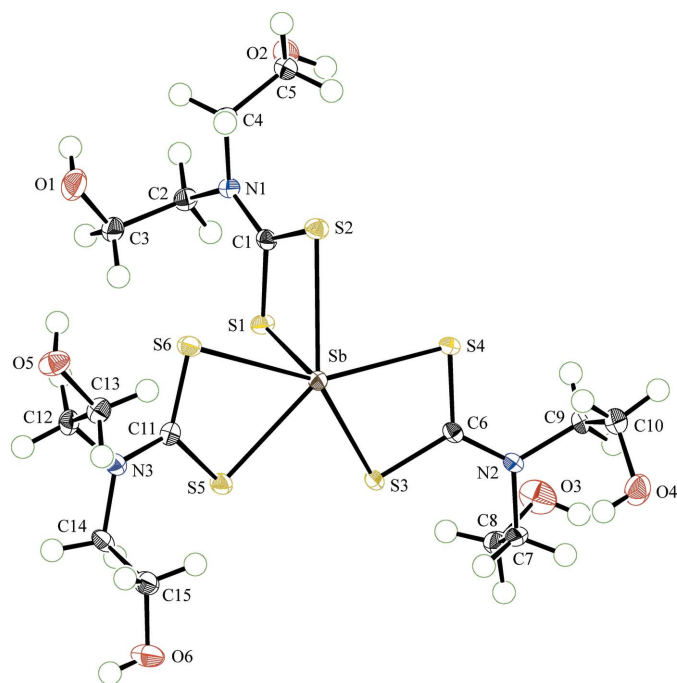


Figure 1
The molecular structure of (I), showing the crystallographic numbering scheme. Displacement ellipsoids are shown at the 50% probability level.

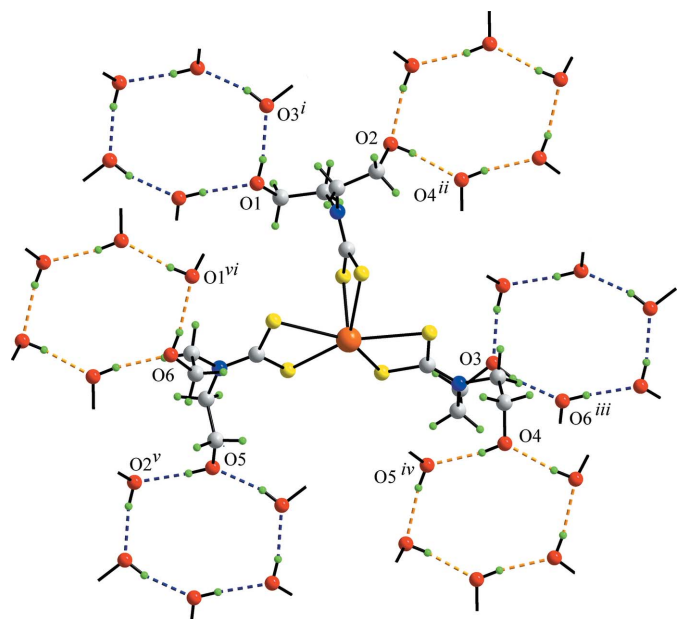


Figure 2
Hydrogen-bonding interactions in (I) mediated by O—H...O interactions, shown as dashed lines; see Table 2 for symmetry operations. Colour code: Sb orange, S yellow, O red, N blue, C grey and H green.

for the formation of intermolecular Sb...S interactions in many related systems, but no such interaction is present in (I).

Details of the hydrogen-bonding parameters are summarized in Table 2, from which it can be seen that each OH group functions as a hydrogen-bonding donor and acceptor. As shown in Fig. 2, each hydroxyl group is associated with an essentially planar 12-membered [\cdots O—H]₆ ring. Those rings, highlighted with blue dashed lines in Fig. 2, involve atoms O1,

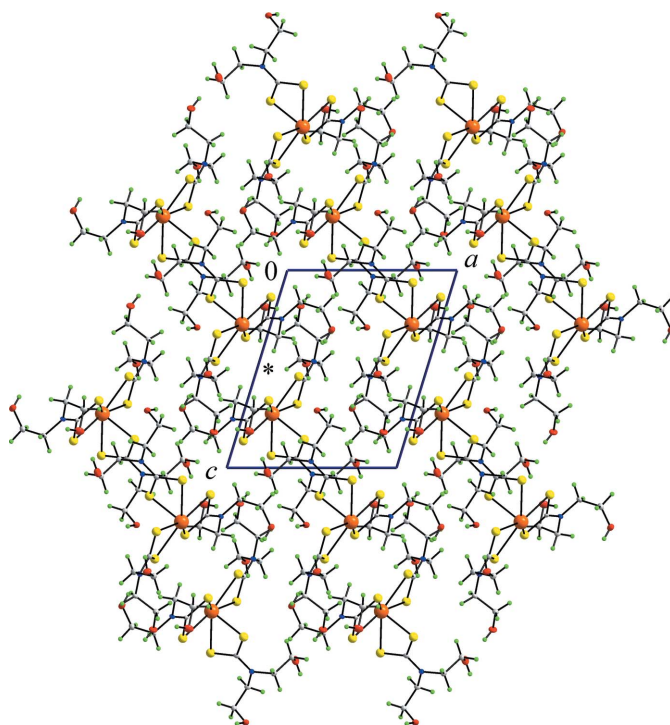


Figure 3
The crystal packing in (I); the colour code is as for Fig. 2.

O3 and O6. The net result of these interactions is the formation of a three-dimensional array, as shown in Fig. 3. Also shown in this figure is the relative disposition of a pair of centrosymmetric molecules which are aligned so as to form, potentially, the commonly observed dimeric motif. In Fig. 3, a potential point of aggregation is indicated with an asterisk, but the intermolecular Sb...S separation is greater than 5.7 Å. Association *via* Sb...S interactions leading to dimers is precluded in (I) due to the intervention of the hydrogen-bonded residues.

Experimental

Compound (I) was prepared in a manner that differed from the literature procedure (Venkatachalam *et al.*, 1997). A suspension of ammonium bis(hydroxyethyl)dithiocarbamate (5.94 g, 3.0 mmol), prepared after the procedure of Ramos *et al.* (2005), in methanol (200 ml) was added to a solution of SbCl₃ (Acros Organics, 2.29 g, 1.0 mmol) in methanol (30 ml) with stirring. The ligand dissolved and the solution turned iridescent yellow. The product was filtered off and recrystallized from an ethanol solution to provide 3.78 g (65.1%) of (I) [m.p. 413.5–414.5 K; literature 403–405 K (Venkatachalam *et al.*, 1997)].

Crystal data

[Sb(C₅H₁₀NO₂S₂)₃]
M_r = 662.53
 Triclinic, *P* $\bar{1}$
a = 9.942 (2) Å
b = 11.011 (2) Å
c = 12.458 (3) Å
 α = 104.422 (4)°
 β = 105.294 (4)°
 γ = 94.445 (3)°

V = 1258.9 (5) Å³
Z = 2
D_x = 1.748 Mg m⁻³
 Mo *K*α radiation
 μ = 1.63 mm⁻¹
T = 93 (2) K
 Block, yellow
 0.20 × 0.20 × 0.20 mm

Data collection

Rigaku AFC12K/SATURN724 31926 measured reflections
 diffractometer 5694 independent reflections
 ω scans 5555 reflections with $I > 2\sigma(I)$
 Absorption correction: multi-scan $R_{\text{int}} = 0.033$
 (ABSCOR; Higashi, 1995) $\theta_{\text{max}} = 27.5^\circ$
 $T_{\text{min}} = 0.736$, $T_{\text{max}} = 1.000$
 (expected range = 0.531–0.722)

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0275P)^2 + 0.8849P]$
 $R[F^2 > 2\sigma(F^2)] = 0.020$ where $P = (F_o^2 + 2F_c^2)/3$
 $wR(F^2) = 0.051$ $(\Delta/\sigma)_{\text{max}} = 0.002$
 $S = 1.08$ $\Delta\rho_{\text{max}} = 0.69 \text{ e } \text{\AA}^{-3}$
 5694 reflections $\Delta\rho_{\text{min}} = -0.52 \text{ e } \text{\AA}^{-3}$
 298 parameters
 H-atom parameters constrained

Table 1
 Selected geometric parameters (\AA , $^\circ$).

Sb—S1	2.4628 (6)	S3—C6	1.7340 (15)
Sb—S2	2.8733 (7)	S4—C6	1.7173 (16)
Sb—S3	2.6384 (6)	S5—C11	1.7322 (16)
Sb—S4	2.7548 (5)	S6—C11	1.7053 (17)
Sb—S5	2.6714 (6)	N1—C1	1.329 (2)
Sb—S6	2.8451 (6)	N2—C6	1.331 (2)
S1—C1	1.7664 (16)	N3—C11	1.348 (2)
S2—C1	1.6962 (16)		
S1—Sb—S2	67.359 (15)	S3—Sb—S6	141.007 (14)
S1—Sb—S3	91.326 (16)	S4—Sb—S5	142.877 (13)
S1—Sb—S4	89.770 (19)	S4—Sb—S6	152.124 (12)
S1—Sb—S5	92.367 (19)	S5—Sb—S6	64.852 (12)
S1—Sb—S6	85.095 (18)	Sb—S1—C1	92.07 (6)
S2—Sb—S3	138.619 (12)	Sb—S2—C1	80.36 (5)
S2—Sb—S4	78.059 (17)	Sb—S3—C6	88.54 (5)
S2—Sb—S5	136.185 (15)	Sb—S4—C6	85.13 (5)
S2—Sb—S6	74.674 (15)	Sb—S5—C11	89.64 (5)
S3—Sb—S4	66.339 (17)	Sb—S6—C11	84.54 (5)
S3—Sb—S5	76.559 (14)		

Table 2
 Hydrogen-bond geometry (\AA , $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1—H1o \cdots O3 ⁱ	0.84	1.84	2.671 (2)	170
O2—H2o \cdots O4 ⁱⁱ	0.84	1.91	2.718 (2)	162
O3—H3o \cdots O6 ⁱⁱⁱ	0.84	1.88	2.701 (2)	166
O4—H4o \cdots O5 ^{iv}	0.84	1.92	2.751 (2)	171
O5—H5o \cdots O2 ^v	0.84	1.99	2.8188 (19)	172
O6—H6o \cdots O1 ^{vi}	0.84	1.87	2.710 (2)	176

Symmetry codes: (i) $-x + 1, -y + 1, -z$; (ii) $-x + 2, -y + 1, -z$; (iii) $-x + 2, -y + 1, -z + 1$; (iv) $-x + 2, -y + 2, -z + 1$; (v) $-x + 1, -y + 2, -z$; (vi) $-x + 1, -y + 2, -z + 1$.

The methylene H atoms were included in a riding-model approximation, with C—H = 0.99 \AA and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The hydroxyl H atoms were located in a difference map and treated as riding, with O—H = 0.84 \AA and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$.

Data collection: *CrystalClear* (Rigaku/MS, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976) and *DIAMOND* (Crystal Impact, 2006); software used to prepare material for publication: *SHELXL97*.

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