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#### **Key indicators**

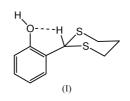
Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.004 Å R factor = 0.063 wR factor = 0.137 Data-to-parameter ratio = 15.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title compound,  $C_{10}H_{12}OS_2$ , the dithiane ring adopts a chair conformation. An intramolecular  $C-H\cdots O$  interaction forms an S(5) graph-set ring. Intermolecular  $O-H\cdots S$  and  $C-H\cdots O$  hydrogen bonds link the molecules into sheet-like assemblies parallel to the *ab* plane.

2-(2-Hydroxyphenyl)-1,3-dithiane

### Comment

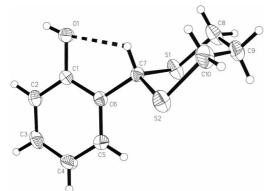
Protection of carbonyl compounds as cyclic S-acetals, such as 1,3-dithianes and dithiolanes, and regeneration of the parent carbonyl compounds from these derivatives are of considerable contemporary interest to organic chemists, in view of the central importance of carbonyl compounds in organic synthesis (Greene & Wuts, 1991). Derivatives of 1,3-dithiane are particularly important as versatile intermediates in organic synthesis because they are utilized as acyl anion equivalents via temporary inversion of electrophilic carbonyl carbon. This occurs via metallation, thereby allowing carbon-carbon bond formation via electrophiles (Grobel & Seebeck, 1977). These derivatives are easy to form and display remarkable stability towards acid as well as acidic hydrolysis. The presence of electron-donor and electron-withdrawing groups on the aromatic ring is a crucial factor which determines the ease of cleavage of C-S bonds, using soft electrophiles as reagents (Corey & Erickson, 1971). However, the presence of the dithiane moiety together with another functional group capable of forming hydrogen bonds in the solid state is of special interest, to obtain an insight into the orientation and hydrogen bonding of the dithiane ring (Jeffrey & Saenger, 1991; Desiraju, 1996). In view of this important factor, the crystal structure determination of the title compound, (I), was undertaken, and the results are reported here.



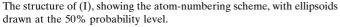
The bond lengths observed in (I) (Fig. 1) have normal values (Allen *et al.*, 1987) and the bond lengths and angles are comparable to those in the related structure 2-(2,6-diffuorophenyl)-1,3-dithiane, (II) (Adamson *et al.*, 1995). As observed in (II), the dithiane ring adopts a chair conformation. The conformation about the C6–C7 bond joining the dithiane and phenyl rings is best described by the torsion angles C1–C6–C7–S1 [–97.1 (2)°] and C5–C6–C7–S1 [80.2 (3)°]. An intramolecular C7–H7···O1 interaction forms an *S*(5) graphset ring (Etter *et al.*, 1990). In the solid state, intermolecular

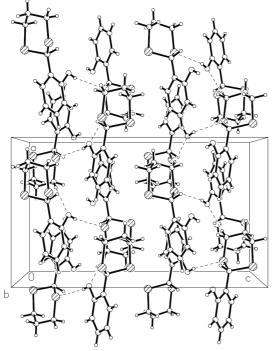
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#### Figure 1





#### Figure 2

Packing diagram of (I), showing molecular sheets parallel to the *ab* plane.

hydrogen bonds involving the hydroxy group and the dithiane ring,  $O1-H11\cdots S2^{i}$  and  $C10-H10A\cdots O1^{ii}$ , interconnect the molecules into sheet-like assemblies parallel to the *ab* plane (Fig. 2). The symmetry codes are as in Table 2.

## **Experimental**

To a solution of salicylaldehyde (18.7 mmol) in 1,3-propanedithiol (39.9 mmol), boron trifluoride etherate (2 ml) was added with stirring at 273 K for 15 min. After completion of the reaction, the mixture was poured into a saturated aqueous solution of sodium hydrogen carbonate (50 ml) and was extracted three times with ethyl acetate. The organic layer was washed repeatedly with sodium hydrogen carbonate and water. Finally, the organic layer was dried over anhydrous sodium sulfate. The solvent was removed under vacuum and the title compound was isolated by column chromatography, with petroleum ether as eluant. Single crystals suitable for X-ray analysis were obtained from a solution in petroleum ether.

### Crystal data

$C_{10}H_{12}OS_2$	Mo $K\alpha$ radiation
$M_r = 212.32$	Cell parameters from 4761
Orthorhombic, Pbca	reflections
a = 10.4039 (7)  Å	$\theta = 2.3 - 28.3^{\circ}$
b = 11.0435 (8) Å	$\mu = 0.48 \text{ mm}^{-1}$
c = 17.7471 (12)  Å	T = 293 (2)  K
$V = 2039.1 (2) \text{ Å}^3$	Block, colorless
Z = 8	$0.46 \times 0.40 \times 0.24 \text{ mm}$
$D_x = 1.383 \text{ Mg m}^{-3}$	

#### Data collection

Siemens SMART CCD areadetector diffractometer (i) scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  $T_{\min} = 0.810, \ T_{\max} = 0.894$ 10961 measured reflections

### Refinement

Refinement on  $F^2$  $w = 1/[\sigma^2(F_o^2) + (0.0449P)^2]$  $R[F^2 > 2\sigma(F^2)] = 0.063$  $wR(F^2) = 0.137$  $(\Delta/\sigma)_{\rm max} < 0.001$ S = 1.24 $\Delta \rho_{\rm max} = 0.55 \text{ e } \text{\AA}^{-3}$ 2524 reflections  $\Delta \rho_{\rm min} = -0.26 \text{ e } \text{\AA}^{-3}$ 166 parameters All H-atom parameters refined

### Table 1

Selected geometric parameters (Å, °).

S1-C8	1.814 (3)	S2-C7	1.813 (3)
S1-C7	1.814 (3)	O1-C1	1.364 (3)
S2-C10	1.812 (3)	C6-C7	1.502 (4)
C8-S1-C7	99.4 (2)	C6-C7-S1	109.1 (2)
C10-S2-C7	98.2 (2)	S2-C7-S1	111.8 (1)
C6-C7-S2	111.3 (2)		
C1-C6-C7-S2	139.0 (2)	C1-C6-C7-S1	-97.1 (2)
C5-C6-C7-S2	-43.6 (3)	C5-C6-C7-S1	80.2 (3)

2524 independent reflections

 $R_{\rm int}=0.034$ 

 $\theta_{\rm max} = 28.3^{\circ}$ 

 $h = -13 \rightarrow 11$ 

 $k = -14 \rightarrow 14$ 

 $l = -23 \rightarrow 19$ 

+ 2.164*P*]

2080 reflections with  $I > 2\sigma(I)$ 

where  $P = (F_o^2 + 2F_c^2)/3$ 

#### Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$\begin{array}{c} \hline C7-H7\cdots O1\\ O1-H11\cdots S2^{i}\\ C10-H10A\cdots O1^{ii} \end{array}$	0.96 (3) 0.73 (3) 0.96 (4)	2.35 (3) 2.56 (3) 2.54 (4)	2.769 (3) 3.254 (2) 3.387 (4)	105 (2) 159 (3) 147 (3)
	1.1 (**	<u>\ 1 1 .</u>		

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

All H atoms were located in a difference Fourier map and were refined isotropically. The C-H distances ranged from 0.89 (4) to 1.00 (4) Å and the O–H distance is 0.73 (3) Å. The highest peak was 0.89 Å from S1.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

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