## $C_{17}H_{13}NO_4S_2.C_2H_6OS$

#### Data collection

Enraf-Nonius CAD-4	$\theta_{\rm max} = 24.98^{\circ}$
diffractometer	$h = -38 \rightarrow 38$
$2\theta/\omega$ scans	$k = 0 \rightarrow 6$
Absorption correction: none	$l = 0 \rightarrow 26$
3603 measured reflections	3 standard reflections
3603 independent reflections	every 100 reflections
3285 reflections with	frequency: 120 min
$I > 2\sigma(I)$	intensity decay: 4%

## Refinement

Refinement on  $F^2$  $(\Delta/\sigma)_{\rm max} = 0.053$  $\Delta \rho_{\rm max} = 0.311 \ {\rm e} \ {\rm \AA}^{-3}$  $R[F^2 > 2\sigma(F^2)] = 0.054$  $wR(F^2) = 0.121$  $\Delta \rho_{\rm min} = -0.291 \ {\rm e} \ {\rm \AA}^{-3}$ S = 1.128Extinction correction: none 3285 reflections Scattering factors from 259 parameters International Tables for H atoms riding Crystallography (Vol. C)  $w = 1/[\sigma^2(F_o^2) + (0.0808P)^2]$ where  $P = (F_o^2 + 2F_c^2)/3$ 

Table 1. Selected geometric parameters (Å, °)

S1-C1	1.738 (2)	O2—C5	1.203 (2)
S1—C3	1.736 (2)	O3—C5	1.293 (2)
S2-C1	1.617 (2)	C2—C3	1.482 (2)
N—C1	1.385 (2)	C3—C6	1.352 (2)
N—C2	1.397 (2)	C4—C5	1.513 (2)
N—C4	1.431 (2)	C6C7	1.443 (2)
01—C2	1.197 (2)		
C1-S1-C3	92.96 (8)	N—C2—C3	109.46 (12)
C1—N—C2	116.78 (13)	C6—C3—C2	119.36 (13)
C1-N-C4	123.57 (13)	C6C3S1	130.21 (13)
C2—N—C4	118.61 (12)	C2-C3-S1	110.42 (11)
N—C1—S2	126.24 (13)	N-C4-C5	110.93 (14)
N-C1-S1	110.34(11)	O2—C5—O3	125.0 (2)
S2—C1—S1	123.42 (10)	02—C5—C4	123.35 (15)
01-C2-N	122.4 (2)	O3-C5-C4	111.69(14)
01—C2—C3	128.12 (15)	C3-C6C7	131.89 (14)

Most H atoms were geometrically placed, except for H3 which was unambiguously located by difference Fourier methods, and were constrained with a riding model.

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: CAD-4 Software. Program(s) used to solve structure: SHELXS86 (Sheldrick, 1990). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: Xtal\_GX (Hall & du Boulay, 1995) and PLUTON92 (Spek, 1992).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: CF1236). Services for accessing these data are described at the back of the journal.

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# 2,2'-Biphenol

Accepted.

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### Abstract

The structure of anhydrous 2,2'-biphenol,  $C_{12}H_{10}O_2$ , is described, together with the hydrogen-bonding scheme. The first hydrogen bond is an intramolecular one, while the second one connects the biphenol molecules to build infinite chains running in the **b** direction. A comparison with the structure of 2,2'-biphenol monohydrate is made.

## Comment

In relation to the synthesis of Lewis acid derivatives of biphenol (Schaverien *et al.*, 1992), we are interested in the organization of pure phenols in the solid state. Recently, biphenol has been used as an activator of coordination catalysts (Matsukawa & Mikami, 1996; Mikami & Matsukawa, 1997), but little is known about the structures of these complexes. The structure of 2,2'-biphenol monohydrate, (II), was recently described (Chen *et al.*, 1996), and a comparison can be made of the structures of the monohydrate, (II), and the anhydrous form, (I).



All bond lengths and angles found in (I) (Table 1) agree with those in the literature (Allen *et al.*, 1987). The C6—C7 single bond is significantly longer than the C—C bonds in the two aromatic rings. The dihedral angles between the aromatic rings, namely  $48.4(1)^{\circ}$  in (I) and  $67.6(1)^{\circ}$  in (II), are significantly different.

It is also worth noting that the two hydroxyl groups of (I) are cis with respect to one another, while in the monohydrate, this is not true. This cis arrangement allows the formation of an intramolecular hydrogen bond, in which the  $O \cdots O$  distance is 2.606(1)Å (Table 2). There is also a second hydrogen bond, linking the biphenyl molecules into a supramolecular chain running along the **b** direction (Fig. 1). This contrasts with the case of the monohydrate, where the trans arrangement of the O atoms inhibits an intramolecular bond, and hydrogen bonding via the lattice water molecules connects the biphenyl molecules into a twodimensional network.



Fig. 1. An ORTEPII (Johnson, 1976) projection of the crystal structure of (I) along the a direction, shown with 35% probability displacement ellipsoids. The hydrogen bonds are represented by dashed lines.

### **Experimental**

Crystals of the title compound were obtained by slow evaporation of a solution of amorphous 2,2'-biphenol (1 mmol) in dry toluene (10 ml).

Crystal data

$C_{12}H_{10}O_2$	Mo $K\alpha$ radiation
$M_r = 186.21$	$\lambda = 0.7107 \text{ Å}$

Monoclinic	
$P2_{1}/c$	
a = 7.392(2) Å	
b = 8.965 (6)  Å	
c = 14.925(4) Å	
$\beta = 102.90(2)^{\circ}$	
V = 964.1 (6) Å <sup>3</sup>	
Z = 4	_
D 1 000 14	-3

 $D_x = 1.283 \text{ Mg m}^{-3}$  $D_m$  not measured

#### Data collection

Enraf-Nonius CAD-4 diffractometer  $\omega$  scans Absorption correction: none 3079 measured reflections 2982 independent reflections 1695 reflections with  $I > 1.8\sigma(I)$ 

# Refinement

Refinement on F R = 0.042wR = 0.053S = 1.7681695 reflections 168 parameters All H atoms refined  $w = 1/[\sigma^2(F_o)]$  $+ 0.00042 |F_o|^2$  $(\Delta/\sigma)_{\rm max} = 0.050$ 

Cell parameters from 24 reflections  $\theta = 9.9 - 11.5^{\circ}$  $\mu = 0.087 \text{ mm}^{-1}$ T = 293 KStout monoclinic prism  $0.50 \times 0.48 \times 0.42$  mm Translucent pale yellow

- $R_{\rm int} = 0.019$  $\theta_{\rm max} = 29.96^{\circ}$  $h = -10 \rightarrow 10$  $k = 0 \rightarrow 12$  $l = 0 \rightarrow 20$ 2 standard reflections frequency: 120 min intensity decay: none
- $\Delta \rho_{\rm max} = 0.16 \ {\rm e} \ {\rm \AA}^{-3}$  $\Delta \rho_{\rm min} = -0.24 \ {\rm e} \ {\rm \AA}^{-3}$ Extinction correction: Zachariasen (1967, 1968) Extinction coefficient:  $3.20(4) \times 10^{-5}$ Scattering factors from International Tables for Crystallography (Vol. C)

# Table 1. Selected geometric parameters (Å, °)

01-C1	1.374 (1)	C6—C7	1.486(2)
O2-C12	1.381 (2)	C7—C8	1.396(2)
C1—C2	1.383 (2)	C7—C12	1.390(2)
C1—C6	1.389 (2)	C8—C9	1.375(2)
C2—C3	1.372 (2)	C9—C10	1.367 (2)
C3—C4	1.370(2)	C10-C11	1.376 (2)
C4C5	1.377 (2)	C11—C12	1.378(2)
C5—C6	1.395 (2)		
01—C1—C2	121.3(1)	C6—C7—C8	119.1 (1)
01—C1—C6	117.07 (9)	C6—C7—C12	124.2(1)
C2-C1-C6	121.6(1)	C8—C7—C12	116.7(1)
C1-C2-C3	119.6(1)	C7—C8—C9	121.8(1)
C2—C3—C4	120.3 (1)	C8—C9—C10	119.9(1)
C3-C4-C5	119.8(1)	C9-C10-C11	119.9(1)
C4C5C6	121.6(1)	C10C11C12	120.0(1)
C1-C6-C5	116.9(1)	O2—C12—C7	121.2(1)
C1-C6-C7	123.23 (9)	02—C12—C11	117.2(1)
C5—C6—C7	119.8(1)	C7—C12—C11	121.5(1)

# Table 2. Hydrogen-bonding geometry (Å, °)

D—-H···A	D—H	H···A	$D \cdot \cdot \cdot A$	D—H···A
O1—H1′···O2'	0.86(2)	1.79(2)	2.645(1)	168(1)
O2—H2′· · ·O1	0.86(2)	1.80(2)	2.606 (1)	156(1)
Symmetry code: (i)	$-x, y - \frac{1}{2}, \frac{3}{2}$	— z.		

The H atoms were found in a final difference Fourier map and were refined isotropically.

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: TEXSAN (Molecular Structure Corporation, 1995), Program(s) used to solve structure: SIR88 (Burla et al., 1989). Program(s) used to refine structure: *TEXSAN*. Software used to prepare material for publication: *TEXSAN*.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: OS1005). Services for accessing these data are described at the back of the journal.

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# Comment

Two types of transformation of the pentacyclic holostane skeleton in triterpene glycosides of holothurians resulting from the action of protonic acids are well known (Elyakov & Stonik, 1986). One type is related to a double-bond migration in aglycones with a labile  $9\beta$ -H-7(8)-ene fragment. The stereochemical details of this migration have been studied carefully (Ilyin *et al.*, 1991). The second transformation is typical for aglycones with a labile  $12\alpha$ -hydroxy-9(11)-ene fragment and forms holosta-7,9-dienes under appropriate conditions (Chanley & Rossi, 1969). We report herein the structure of a product (Fig. 1) of this type of transformation, namely the artificial genin of the holothurian Bohadschia argus glycosides (Antonov & Stonik, 1986), as its  $\frac{1}{3}$ -acetone solvate  $\frac{1}{3}$ -hydrate, (I). This holothurinogenin. seychellogenin, has been isolated from the sea cucumber Bohadschia koellikeri (Roller et al., 1969).



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# (20S)-3β-Hydroxyholosta-7,9-diene

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## Abstract

The crystal and molecular structures of the title triterpenoid have been determined. There are three molecules of this triterpenoid ( $C_{30}H_{46}O_3$ ), one molecule of acetone ( $C_3H_6O$ ) and a molecule of water ( $H_2O$ ) in the asymmetric unit, *i.e.* formula  $C_{30}H_{46}O_3$ . $\frac{1}{3}C_3H_6O$ . $\frac{1}{3}H_2O$ . The molecules of the triterpenoid have different conformations of both the side chains and the  $\gamma$ -lactone rings.

© 1998 International Union of Crystallography Printed in Great Britain – all rights reserved The conformation of the triterpenoid molecule is flexible (Fig. 2). Three different conformations are observed in the crystal as a consequence of this conformational flexibility. There are three molecules of seychellogenin in the asymmetric unit. While four rings possess stable conformations in these molecules (ring A chair, ring B distorted C5 $\alpha$ -sofa, ring C distorted C13 $\beta$ ,C14 $\alpha$ -halfchair and ring D C14 $\alpha$ -envelope), the fifth ring (E) differs between the three molecules, varying from a C17 $\beta$ envelope in molecule A to a C13 $\alpha$ ,C17 $\beta$ -half-chair in molecule C. The side-chain conformations differ from each other by rotation around the bonds C23—C24 and C24—C25.



Fig. 1. The molecular structure of molecule *A* of the title compound with 30% probability non-H-atom displacement spheres (H atoms as spheres of arbitrary radii).