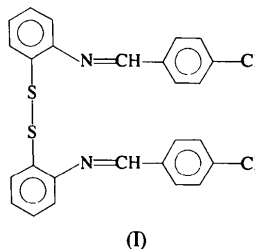


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their metal complexes have previously been used as models for biological systems (Sugiura & Hirayama, 1976; Casella, 1984). The direct approach in synthesizing a free Schiff's base from the condensation of 2-mercaptoaniline with a carbonyl compound often results in an undesirable side reaction involving ring closure with the formation of a heterocyclic compound (Jadamus, Fernando & Freiser, 1964; Charles & Freiser, 1953; Sugiura, Hirayama, Tanaka & Ishizu, 1975; İde, Öztaş, Ancın & Tüzün, 1996). The title compound, (I), was obtained by the condensation of 4-chlorobenzaldehyde with 2,2'-diaminodiphenyl disulfide. This preformed Schiff's base can then be used to obtain Ni^{II}, Zn^{II}, Cu^{II} and Cd^{II} complexes by an electrochemical reductive cleavage of the disulfide bond (Castro *et al.*, 1990). Disulfides including a C—S—S—C chain have been used also as antiwear and extreme-pressure additives in lubricating oils.



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N,N'-Bis(4-chlorobenzylidene)-2,2'-diaminodiphenyl Disulfide

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Abstract

In the title compound, *N,N'*-bis(4-chlorobenzylidene)-disulfanediybis(2-aniline), C₂₆H₁₈Cl₂N₂S₂, two 2-(4-chlorobenzylideneamino)phenyl sulfide moieties, which can act as tridentate ligands, are joined through their S atoms. The molecule has a *trans* configuration about the S—S bond. The bond lengths and angles are nearly identical between the two halves of the molecule.

Comment

Schiff's bases bearing imine N and anionic S atoms constitute an important class of polydentate ligands and

The present investigation is part of a research program involving the structural studies of disulfides. Structural properties of such compounds have been discussed previously by Hordvik (1966), Lee (1972), Meyer (1976), Muller & Diemann (1987), Marsden & Smith (1988) and Krasinski & Fackler (1994). The observed S—S and S—C bond lengths (Table 1) are in agreement with the values reported in the literature (Allen *et al.*, 1987; Orpen *et al.*, 1989). The torsion angles C7—C1—S1—S1' of 19.8 (4)° and C7'—C1'—S1'—S1 of 0.4 (4)° are similar to the corresponding angles in diphenyl disulfide (17.7 and 0.2°), but different from the values in 2,2'-diaminodiphenyl disulfide (83.6 and 83.4°; Lee & Bryant, 1970), which indicates a *pπ*–*dπ* interaction between the fully occupied *p_z* orbital on the C atom with an empty *d* orbital on the S atom in the latter compound. Such a *π* overlap does not seem as favourable in the title compound. In a disulfide group, the regular dihedral angle of about 90° corresponds to S—S bond lengths of about 2.03 Å (Hordvik, 1966). In the title compound, the C1—S1—S1'—C1' torsion angle of –89.8 (2)° and the S—S bond length of 2.032 (2) Å agree with this observation.

Mean plane A (formed by atoms C1–C3 and C5–C7) makes an angle of 86.3 (1)° with mean plane A' (formed by atoms C1'–C7'). The S1 atom deviates by 0.081 (1) Å from plane A, while there is no deviation of the S1' atom from plane A'. The N atoms

are coplanar with their respective rings. The torsion angles C2—N1—C8—C9 and C2'—N1'—C8'—C9' are 178.3(4) and $-178.3(4)^\circ$, respectively. The other bond distances and angles are close to those reported for *N,N'*-bis(*p*-chlorobenzylidene)-1,3-propanediamine (Tinant & Declercq, 1986) and *N*-(4-chlorobenzylidene)-2-(4-imidazolyl)ethylamine (Bocelli & Cantoni, 1992). There are weak intermolecular contacts. In the crystal structure, the molecules thus appear to be held together by van der Waals forces.

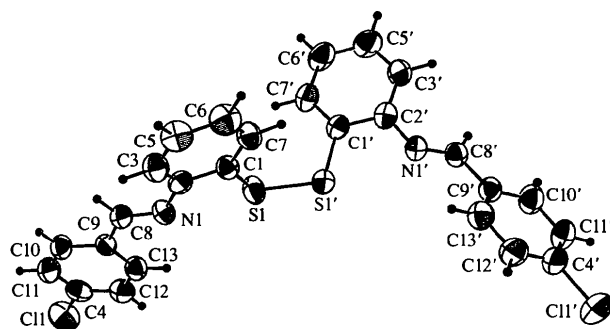


Fig. 1. The molecular structure of the title compound with the atomic labelling. The displacement ellipsoids are drawn at the 50% probability level. Unlabelled H atoms possess the same numerical label as their parent C atoms.

Experimental

To obtain the title compound, 4-chlorobenzaldehyde (0.283 g, 2 mmol) in alcohol (10 ml) was added to a solution of 2,2'-diaminodiphenyl disulfide (0.248 g, 1 mmol) in alcohol (20 ml). The solution was heated to 323 K for 5–6 h. During this time, a yellow crystalline product was deposited. It was recrystallized from chloroform (m.p. 411 K). IR (KBr) ν_{\max} : 1574 (s), 1500 (s, C=N), 680 (m, C—S), 476 (m, S—S), 1290 (m, C—N). $^1\text{H NMR}$ (δ_{H} , 400 MHz, CDCl_2): 8.46 (2H, s, 2CH=N), 7.91 (4H, d, Ar-12,12',11,11-H), 7.70 (2H, d, Ar-7,7'-H), 7.48 (4H, d, Ar-13,13'-10,10'-H), 7.19 (4H, m, Ar-3,3',6,6'-H), 7.06 (2H, dd, Ar-5,5'-H). UV (acetone) λ_{\max} : 327.7 nm ($\epsilon = 25240 \text{ dm}^3 \text{ mol}^{-1} \text{ cm}^{-1}$).

Crystal data

$\text{C}_{26}\text{H}_{18}\text{Cl}_2\text{N}_2\text{S}_2$
 $M_r = 493.48$
 Monoclinic
 $P2_1/c$
 $a = 11.7998(12) \text{ \AA}$
 $b = 10.1564(11) \text{ \AA}$
 $c = 19.5352(18) \text{ \AA}$
 $\beta = 93.410(9)^\circ$
 $V = 2337(5) \text{ \AA}^3$
 $Z = 4$
 $D_x = 2.805 \text{ Mg m}^{-3}$
 D_m not measured

Data collection

Enraf–Nonius CAD-4
 diffractometer

Mo $K\alpha$ radiation
 $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 25
 reflections
 $\theta = 10.31\text{--}18.01^\circ$
 $\mu = 0.948 \text{ mm}^{-1}$
 $T = 293(1) \text{ K}$
 Prismatic
 $0.56 \times 0.32 \times 0.10 \text{ mm}$
 Yellow

2418 reflections with
 $I > 2\sigma(I)$

$\omega/2\theta$ scans

Absorption correction:
 empirical *via* ψ scan
 (MolEN; Fair, 1990)
 $T_{\min} = 0.876$, $T_{\max} = 0.910$
 4755 measured reflections
 4477 independent reflections

$R_{\text{int}} = 0.037$

$\theta_{\max} = 26.32^\circ$
 $h = -14 \rightarrow 14$
 $k = 0 \rightarrow 12$
 $l = -23 \rightarrow 0$
 3 standard reflections
 frequency: 120 min
 intensity decay: 5.17%

Refinement

Refinement on F

$R = 0.046$
 $wR = 0.045$
 $S = 0.71$
 2418 reflections
 289 parameters
 H atoms: see below
 Weighting scheme: see
 below

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.285 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.114 \text{ e \AA}^{-3}$
 Extinction correction: none
 Scattering factors from
*International Tables for X-ray
 Crystallography* (Vol. IV)

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

N1—C2	1.406 (5)	N1'—C2'	1.415 (5)
N1—C8	1.258 (5)	N1'—C8'	1.259 (5)
C6—C7	1.364 (7)	C6'—C7'	1.375 (7)
C8—C9	1.457 (6)	C8'—C9'	1.472 (6)
C11—C4	1.735 (5)	C11'—C4'	1.740 (5)
S1—C1	1.770 (4)	S1'—C1'	1.781 (4)
S1—S1'	2.032 (2)		
S1—C1—C7	125.4 (3)	S1'—C1'—C7'	125.1 (3)
S1—C1—C2	115.1 (3)	S1'—C1'—C2'	114.5 (3)
S1'—S1—C1	104.6 (1)	S1—S1'—C1'	106.4 (1)
N1—C2—C1	114.9 (3)	N1'—C2'—C1'	116.3 (4)
N1—C2—C3	126.6 (4)	N1'—C2'—C3'	125.0 (4)
C2—N1—C8	122.0 (4)	C2'—N1'—C8'	120.2 (3)
N1—C8—C9	121.8 (4)	N1'—C8'—C9'	121.9 (4)
C8—C9—C10	120.2 (4)	C8'—C9'—C10'	120.3 (4)
C8—C9—C13	121.1 (4)	C8'—C9'—C13'	120.4 (4)
C11—C4—C11	120.1 (4)	C11'—C4'—C11'	119.7 (4)
C11—C4—C12	119.0 (3)	C11'—C4'—C12'	118.2 (4)

The weighting scheme applied was $w = (197.30/F)^2$ if $F \geq 197.30$ and $w = 1.0$ if $197.3 > F$ and $F^2 < 2\sigma F^2$, otherwise $w = 0$. H atoms were placed geometrically at a distance of 0.95 \AA from their corresponding C atoms, while the H atoms of the C8 and C8' atoms were taken from a difference map. For all H atoms, a riding model was used with $U_{\text{iso}}(\text{H}) = 1.3U_{\text{eq}}(\text{C})$.

Data collection: CAD-4 Express (Enraf–Nonius, 1993). Cell refinement: CAD-4 VAX/PC Fortran System (Enraf–Nonius, 1988). Data reduction: MolEN (Fair, 1990). Program(s) used to solve structure: SIR (Burla *et al.*, 1989) in MolEN. Program(s) used to refine structure: LSMF in MolEN. Molecular graphics: ORTEPII (Johnson, 1976). Software used to prepare material for publication: MolEN.

The authors wish to acknowledge the purchase of the CAD-4 diffractometer under Grant DPT/TBAG1 of the Scientific and Technical Research Council of Turkey.

Lists of atomic coordinates, displacement parameters, structure factors and complete geometry have been deposited with the IUCr (Reference: AB1401). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Acta Cryst. (1997). **C53**, 378–379

N-Acetyl-2,3,4,6-tetra-*O*-acetyl-*N*-benzyl-β-D-glucopyranosylamine

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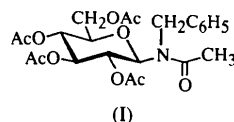
(Received 16 May 1996; accepted 21 October 1996)

Abstract

The solid-state conformation of the title compound, C₂₃H₂₉NO₁₀, has been unequivocally established. The configuration at C1 is β-D-glucopyranose and the pyranose ring is essentially a perfect chair. Only the *E* rotamer along the N—C bond is observed.

Comment

Many natural products and their derivatives have an *N*-acyl group joined to a sugar moiety and as the chemical and biological properties of these compounds depend on their structure and conformation, the correct determination is important. In some *N*-alkyl-*N*-glycosylamides, the presence of two conformational isomers is observed in solution due to hindered rotation about the amide C—N bond. In the present case, (I), no preferred conformation was established in solution so we undertook a crystal structure determination.



Crystals of (I) were obtained (Avalos, Babiano, Durán, Jiménez & Palacios, 1992) from the reaction of acetic anhydride with *N*-benzylglucopyranosylamine in pyridine. The same authors studied several sugar amides and thioamides, including the title compound, by NMR spectroscopy. Our structural analysis shows that only one anomer, *i.e.* β-D-glucopyranose, and one rotamer along the N—C bond (*E*) are present in the crystal.

A *PLATON* view (Spek, 1994) of the title molecule viewed along the *z* axis, together with the atomic numbering, is shown in Fig. 1. Bond lengths and angles agree with those of analogous compounds (Banic, Kojic-Prodic, Kroon-Beterburg & Keglevic, 1994; Vega, López-Castro & Márquez, 1986), although the pyranose endocyclic bond lengths [O1—C2 1.435 (3) and O—C5 1.427 (3) Å] do not show the anomeric effect characteristic of this ring. The phenyl, acetoxy and acetamide groups are essentially planar.

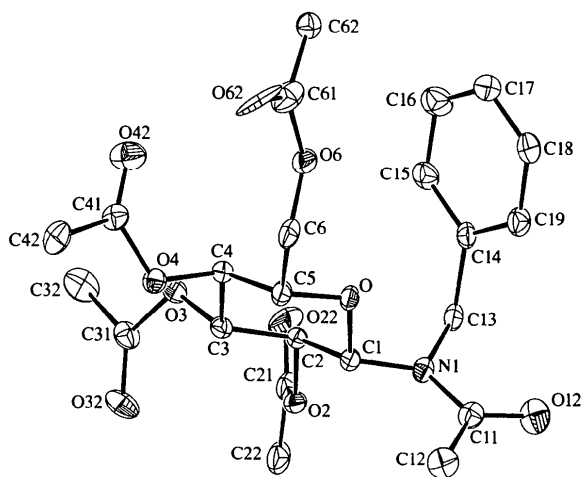


Fig. 1. A *PLATON* (Spek, 1994) plot of the title compound showing 30% probability displacement ellipsoids and the atom-numbering scheme. H atoms have been omitted for clarity.