

between TCNE molecules along this axis is 10.23 (1) Å. Electrical conduction measurements along the charge-transfer axis did not lead to an observable value. A resistivity higher than 1.5 GΩ cm⁻¹ was found.

In (II), each of the two TCNE molecules related by a centre of symmetry is associated with one of the two non-symmetrically equivalent benzene rings of the paracyclophane molecule (*B* in Fig. 3). This leads to a one-dimensional sandwich-type stacking along a face diagonal, **a** + **b** (Fig. 3). The shortest TCNE–TCNE distance along this axis is 14.15 Å. This sandwich arrangement is identical to that observed in [3.3]-paracyclophane–TCNE (Bernstein & Trueblood, 1971). Contrary to this previous study, no disorder of the TCNE molecules was found in the present compounds. Both of them show normal intermolecular distances involving van der Waals interactions. Other structure studies are in progress.

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Structure of Bis(*o*-chlorophenylaminocarbonylmethyl) Disulphide,* $\text{C}_{16}\text{H}_{14}\text{Cl}_2\text{N}_2\text{O}_2\text{S}_2$

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Abstract. $M_r = 401.33$, monoclinic, $P2_1/c$, $a = 9.394(5)$, $b = 17.84(1)$, $c = 10.576(5)$ Å, $\beta = 91.9(1)^\circ$, $V = 1771(1)$ Å³, $Z = 4$, $D_m = 1.5$, $D_x =$

1.495 g cm⁻³, $\lambda(\text{Mo } \text{K}\alpha) = 0.71073$ Å, $\mu = 6.0$ cm⁻¹, $F(000) = 824$, $T = 293$ K. Final $R = 0.031$ for 1403 observed reflections. The molecules adopt a folded conformation with non-parallel phenyl rings and are arranged in columns along the screw axis as a result of intermolecular hydrogen bonds [N–H…O distances of

* *N,N'*-Bis(*o*-chlorophenyl)-2,2'-dithiodiacetamide.

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2.85 (2) and 2.88 (2) Å]. A comparison is made with analogous disulphide compounds. The S–S distance of 2.022 (2) Å indicates partial double-bond character.

Introduction. Recently there has been considerable activity in the field of disulphide chemistry. Certain organic disulphides exhibit anti-seizing and anti-friction properties (Meyer, 1977), and disulphide compounds have been intensively studied in order to assess their ligand potentials. In addition to their chemical interest the disulphides are of biological importance insofar as they determine the structure, and therefore the biological activity, of molecules ranging from insulin to ribonuclease (Dickerson & Geis, 1969). Ribonuclease contains four intrachain disulphide linkages, the reduction of which results in total loss of enzymatic activity. The two peptide chains of insulin are held together by disulphide bridges (Adams *et al.*, 1969).

In view of the importance of the disulphide ligand we describe here the crystal and molecular structure of bis(*o*-chlorophenylaminocarbonylmethyl) disulphide.

Experimental. Title compound prepared by mild oxidation of *N*-(*o*-chlorophenyl)-2-mercaptoacetamide with 30% (v/v) H₂O₂ in 70% methanol–water mixture (Maire, Kapoor, Bachlas & Vincent, 1984). Product recrystallized from hot methanol solution to give needle-shaped white crystals, m.p. 413 K. Single crystals grown by slow evaporation of the methanolic solution of the above ligand at room temperature.

Crystal (0.1 × 0.2 × 0.4 mm) mounted on a CAD-4 Enraf–Nonius diffractometer (graphite monochromator for Mo Kα). Unit-cell parameters refined by least squares on 2sinθ/λ values for 25 reflections (15 < θ < 20°). Density obtained by flotation. Intensities measured using an ω–2θ scan of 0.91 to 10.06° min⁻¹, range Δω = (0.6 + 0.35 tanθ)°, θ_{max} = 25°. Intensities of three standard reflections measured every 5000 s throughout data collection, remained constant within 2%. A total of 1852 independent reflections obtained from which 1403 with *I* > 3σ(*I*) used in structure determination: *h*_{max} = 10, *k*_{max} = 18, *l*_{max} = ±11.

All computations performed on a PDP 11/44 computer (*SDP* software, Frenz, 1978). Structure solved by direct methods (*MULTAN77*, Main, Lessinger, Woolfson, Germain & Declercq, 1977) and completed by difference Fourier methods. Positions of H atoms defined on stereochemical grounds and introduced in the calculation before last refinement cycles. Full-matrix least-squares refinement included anisotropic thermal parameters for non-H atoms and minimized function $\sum w(|F_o| - |F_c|)^2$, $w = 1/\sigma^2(F_o)$. Final refinements converged with *R* = 0.031, *R*_w = 0.037, *S* = 1.22; $(\Delta/\sigma)_{\text{max}} = 0.21$. A final difference Fourier synthesis did not reveal any peak of density > 0.35 e Å⁻³. Atomic scattering factors from *International Tables for X-ray Crystallography* (1974).

Discussion. Final atomic parameters are given in Table 1,* bond distances and angles in Table 2.

* Lists of structure factors, anisotropic thermal parameters and H-atom parameters, distances and angles have been deposited with the British Library Lending Division as Supplementary Publication No. SUP 39638 (10 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

Table 1. Positional parameters and their e.s.d.'s

Anisotropically refined atoms are given in the form of the equivalent isotropic thermal parameter defined as: $\frac{1}{3}[a^2B(1,1) + b^2B(2,2) + c^2B(3,3) + ab(\cos\gamma)B(1,2) + ac(\cos\beta)B(1,3) + bc(\cos\alpha)B(2,3)]$.

	<i>x</i>	<i>y</i>	<i>z</i>	<i>B</i> _{eq} (Å ²)
Cl(1)	0.4723 (1)	0.2645 (8)	0.7832 (1)	4.64 (3)
Cl(2)	0.0466 (2)	0.7288 (8)	0.6741 (1)	5.16 (3)
S(1)	0.1270 (1)	0.4516 (8)	0.3294 (1)	4.29 (3)
S(2)	0.2433 (2)	0.5219 (8)	0.2243 (1)	4.15 (3)
C(1)	0.2612 (5)	0.4061 (3)	0.4306 (5)	4.1 (1)
C(2)	0.2009 (5)	0.3893 (3)	0.5571 (4)	3.2 (1)
N(1)	0.2977 (3)	0.3841 (2)	0.6533 (3)	3.19 (8)
C(3)	0.2649 (4)	0.3699 (3)	0.7808 (4)	3.0 (1)
C(4)	0.3393 (5)	0.3169 (3)	0.8518 (4)	3.3 (1)
C(5)	0.3077 (6)	0.3031 (3)	0.9760 (5)	4.5 (1)
C(6)	0.1993 (7)	0.3412 (4)	1.0286 (5)	6.0 (2)
C(7)	0.1245 (6)	0.3941 (4)	0.9614 (5)	5.2 (1)
C(8)	0.1564 (5)	0.4092 (3)	0.8375 (5)	4.2 (1)
C(9)	0.2311 (5)	0.6095 (3)	0.3086 (4)	3.4 (1)
C(10)	0.3008 (5)	0.6078 (3)	0.4391 (4)	3.3 (1)
N(2)	0.2122 (4)	0.6171 (2)	0.5339 (3)	3.32 (8)
C(11)	0.2508 (5)	0.6221 (3)	0.6634 (4)	3.3 (1)
C(12)	0.1778 (5)	0.6716 (3)	0.7400 (5)	3.8 (1)
C(13)	0.2113 (6)	0.6773 (4)	0.8665 (5)	5.5 (1)
C(14)	0.3191 (6)	0.6340 (4)	0.9192 (5)	6.4 (2)
C(15)	0.3923 (6)	0.5854 (4)	0.8461 (5)	5.2 (1)
C(16)	0.3594 (5)	0.5786 (3)	0.7179 (5)	4.1 (1)
O(1)	0.0743 (3)	0.3794 (2)	0.5712 (3)	4.77 (9)
O(2)	0.4289 (3)	0.5989 (2)	0.4542 (3)	5.20 (9)

Table 2. Bond lengths (Å) and angles (°) and e.s.d.'s

S(1)–S(2)	2.022 (2)		
S(1)–C(1)	1.817 (5)	S(2)–C(9)	1.804 (5)
C(1)–C(2)	1.500 (6)	C(9)–C(10)	1.508 (6)
C(2)–O(1)	1.216 (5)	C(10)–O(2)	1.219 (5)
C(2)–N(1)	1.345 (5)	C(10)–N(2)	1.335 (5)
N(1)–C(3)	1.416 (5)	N(2)–C(11)	1.408 (5)
C(3)–C(4)	1.383 (6)	C(11)–C(12)	1.394 (6)
C(4)–C(5)	1.378 (6)	C(12)–C(13)	1.368 (7)
C(5)–C(6)	1.359 (6)	C(13)–C(14)	1.377 (9)
C(6)–C(7)	1.363 (9)	C(14)–C(15)	1.363 (9)
C(7)–C(8)	1.380 (7)	C(15)–C(16)	1.386 (7)
C(8)–C(3)	1.389 (6)	C(16)–C(11)	1.390 (6)
C(1)–C(4)	1.739 (5)	C(1)–C(12)	1.729 (5)
S(2)–S(1)–C(1)	103.0 (2)	S(1)–S(2)–C(9)	102.8 (2)
S(1)–C(1)–C(2)	109.8 (3)	S(2)–C(9)–C(10)	113.7 (4)
C(1)–C(2)–O(1)	122.6 (4)	C(9)–C(10)–O(2)	121.3 (4)
O(1)–C(2)–N(1)	122.4 (4)	N(2)–C(10)–O(2)	123.7 (4)
C(1)–C(2)–N(1)	115.0 (4)	C(9)–C(10)–N(2)	115.0 (4)
C(2)–N(1)–C(3)	124.7 (4)	C(10)–N(2)–C(11)	126.3 (4)
N(1)–C(3)–C(4)	121.2 (4)	N(2)–C(11)–C(12)	119.3 (4)
N(1)–C(3)–C(8)	120.5 (5)	N(2)–C(11)–C(16)	122.0 (4)
C(3)–C(4)–C(5)	121.2 (5)	C(11)–C(12)–C(13)	120.9 (5)
C(4)–C(5)–C(6)	119.3 (5)	C(12)–C(13)–C(14)	119.8 (6)
C(5)–C(6)–C(7)	120.9 (5)	C(13)–C(14)–C(15)	120.3 (6)
C(6)–C(7)–C(8)	120.4 (5)	C(14)–C(15)–C(16)	120.7 (6)
C(7)–C(8)–C(3)	119.8 (5)	C(11)–C(16)–C(15)	119.6 (5)
C(8)–C(3)–C(4)	118.4 (5)	C(16)–C(11)–C(12)	118.8 (4)
C(3)–C(4)–Cl(1)	119.9 (3)	C(11)–C(12)–Cl(2)	119.9 (4)
C(1)–C(4)–C(5)	118.8 (4)	Cl(2)–C(12)–C(13)	119.2 (5)

The most striking feature of the molecular structure is the folded conformation, shown in Fig. 1, which brings the two phenyl rings close to each other. Owing to the bend on both sides of the S–S bond the molecule appears to be asymmetric. In the limits of the experimental errors there is no significant discrepancy between the two halves of the molecule which are almost alike.

Some bond distances and angles found in analogous disulphide compounds are cited in Table 3. The S–S distance of 2.022 (2) Å indicates some partial double-bond character. The low value of the C–S–S angle of the title compound in comparison with $(C_6H_5)_2S_2$ may be due to the attachment of S atoms to sp^3 C atoms. The two N(1)–C(3) [1.416 (5) Å] and N(2)–C(11) [1.408 (5) Å] bond lengths are different from the C–N value calculated elsewhere (1.51 Å, Kagan, 1979). Both values correspond to a partial double-bond character in the C–N linkage with conjugation between the carbonyl group and the N atom. In both halves of the molecule the dihedral angles between the amide group and the corresponding phenyl ring are slightly different: 48.1 (8)° for the C(1)C(2)O(1)N(1)/C(3)–C(8) angle and 35.7 (8)° for the C(9)C(10)O(2)–N(2)/C(11)–C(16) angle.

We observe that the shorter N–C(O) bond, N(2)–C(10), is related to the more coplanar amide group/phenyl ring. The large values of the valency angles of the N atoms [C(2)–N(1)–C(3) 124.7 (4) and C(10)–N(2)–C(11) 126.3 (4)°] indicate deviation from trigonal N and could be due to the interplay of the electronic repulsion between the phenyl rings and the carbonyl O atoms.

The two phenyl rings are (almost) planar with a maximum deviation from the mean least-squares plane

of 0.009 (6) Å. The N and Cl atoms are nearly coplanar with the phenyl rings: the distances to the plane of their respective phenyl rings are: 0.014 (5) Å for N(1) and N(2) and 0.062 (4) and 0.065 (4) Å for Cl(1) and Cl(2). The Cl–C distances fall within the normal range for chlorophenyl compounds and the Cl–C–C angles do not indicate any steric hindrance.

Finally the two conjugated systems are not at all parallel. The observed dihedral angle between the two phenyl rings is 151 (1)°. Moreover, as can be seen on Fig. 1, the C(6)–C(3)–N(1) and C(14)–C(11)–N(2) directions are oblique with an angle of 35 (1)° between them.

Molecular packing

Two intermolecular hydrogen bonds are observed between the amide N and the carbonyl O (Fig. 2). The

Table 3. Bond distances and angles of analogous disulphides

Compounds	(1)	(2)	(3)	(4)
Bond type (Å)				
S–S	2.030 (5)	2.019 (5)	2.045 (3)	2.022 (2)
S–C	1.79 (1)	1.767 (10)	1.784 (8)	1.804 (5)
	1.81 (1)		1.808 (8)	1.817 (5)
N–C		1.488 (13)	1.476 (13)	1.408 (5)
C–C	1.39 (3)	1.38 (3)	1.39 (3)	1.380 (12)
(phenyl rings)				
C–S–S	106.5 (4)	106.0 (6)	104.3 (6)	103.0 (2)
angle (°)	105.8 (4)		104.6 (6)	102.8 (2)

Compounds:

- (1) Diphenyl disulphide (Lee & Bryant, 1969).
- (2) Bis(*p*-nitrophenyl) disulphide (Ricci & Bernal, 1969).
- (3) Bis(*o*-nitrophenyl) disulphide (Ricci & Bernal, 1970).
- (4) Bis(*o*-chlorophenylaminocarbonylmethyl) disulphide (this work).

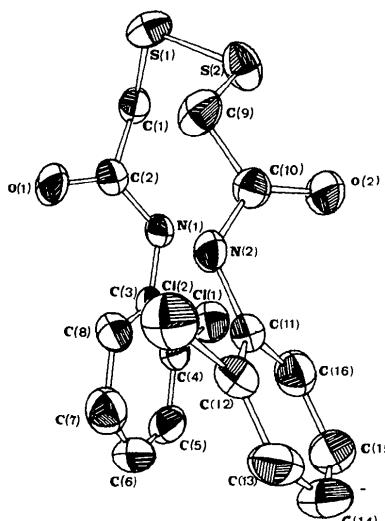


Fig. 1. ORTEP (Johnson, 1976) view of the molecule and labelling system used during the crystallographic study. The thermal parameters are shown as 50% probability envelopes.

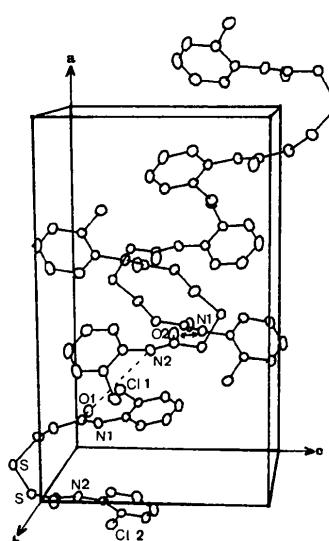


Fig. 2. View of the unit-cell contents down the *b* axis. The hydrogen bonds are indicated by broken lines.

N(1)—O(2) distance is 2.85 (2) Å, whilst the N(2)—O(1) distance is 2.88 (2) Å. In both cases the hydrogen-bond distance is less than 3.0 Å, hence these bonds play an important role in the packing of the molecules in a unit cell. The unit cell consists of four molecules with Cl(1) and Cl(2) as terminal atoms. The molecules are arranged in two pairs held by van der Waals forces of attraction and form columns along the screw axis as a result of the intermolecular hydrogen-bond system.

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Structure of the Twofold Addition Product Tetramethyl 6-Diethylamino-3,7a,10,11b-tetrahydro-7*H*-dipyridazino[4,5-*b*:4',5'-*d*]azepine-1,4,8,11-tetracarboxylate, C₂₂H₂₈N₆O₈

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Abstract. *M_r* = 504.51, triclinic, *P*̄1, *a* = 9.565 (2), *b* = 13.699 (3), *c* = 9.463 (2) Å, *α* = 96.95 (2), *β* = 97.12 (2), *γ* = 97.05 (2)°, *V* = 1209.34 Å³, *Z* = 2, *D_x* = 1.385 Mg m⁻³, *λ*(Cu *Kα*) = 1.5418 Å, *μ*(Cu *Kα*)

= 0.863 mm⁻¹, *F*(000) = 532, *T* = 290 (1) K, *R* = 0.072 for 2469 observed reflections. The molecule contains a seven-membered azepine ring which adopts a distorted boat conformation such that the dihedral angle between the best planes through the two dihydropyridazino rings is 88.5° and thus reduces the crowding of the ester groups on these rings.

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