

Table 1. *Fractional coordinates and equivalent isotropic thermal parameters with e.s.d.'s in parentheses*

$$B_{\text{eq}} = \frac{1}{3} \sum_i \sum_j \beta_{ij} \mathbf{a}_i \cdot \mathbf{a}_j$$

The IUPAC numbering scheme is given where appropriate.

	<i>x</i>	<i>y</i>	<i>z</i>	B_{eq} (Å ²)
Molecule (I)				
O(1) (6)	0.1963 (2)	0.8014 (2)	0.0044 (1)	4.3 (1)
C(2) (5a)	0.3233 (3)	0.8387 (2)	-0.0163 (2)	4.0 (2)
C(3) (5)	0.4168 (4)	0.8252 (2)	0.0424 (2)	4.1 (2)
C(4) (4)	0.5433 (4)	0.8652 (3)	0.0149 (3)	4.5 (2)
C(5) (3)	0.5819 (4)	0.9175 (2)	-0.0687 (2)	4.5 (2)
C(6) (2a)	0.4849 (4)	0.9310 (2)	-0.1258 (2)	4.4 (3)
C(7) (2)	0.4889 (4)	0.9780 (3)	-0.2188 (2)	5.3 (3)
C(8) (1)	0.3539 (5)	0.9610 (3)	-0.2475 (3)	5.7 (3)
C(9) (11b)	0.2635 (4)	0.9130 (3)	-0.1640 (2)	4.6 (2)
C(10) (11c)	0.3586 (4)	0.8947 (2)	-0.0975 (2)	4.1 (2)
C(11) (11a)	0.2015 (4)	0.8099 (3)	-0.1592 (2)	4.4 (2)
C(12) (11)	0.2997 (4)	0.7361 (3)	-0.1949 (2)	4.6 (2)
C(13) (10a)	0.3041 (3)	0.6444 (3)	-0.1290 (2)	4.0 (2)
C(14) (10)	0.3765 (3)	0.5571 (3)	-0.1348 (2)	4.5 (2)
C(15) (9)	0.3608 (4)	0.4797 (3)	-0.0629 (2)	4.8 (2)
C(16) (8)	0.2749 (4)	0.4878 (3)	0.0108 (2)	4.6 (2)
C(17) (7)	0.2029 (3)	0.5757 (3)	0.0158 (2)	4.4 (1)
C(18) (6b)	0.2189 (3)	0.6539 (2)	-0.0552 (2)	4.0 (2)
C(19) (1a)	0.1525 (3)	0.7557 (3)	-0.0635 (2)	4.4 (2)
O(20)	0.3751 (2)	0.7730 (2)	0.1234 (1)	5.3 (1)
C(21)	0.4671 (5)	0.7528 (4)	0.1858 (3)	6.4 (3)
C(22)	0.7221 (4)	0.9553 (3)	-0.0956 (3)	5.8 (2)
O(23)	0.5810 (3)	1.0223 (2)	-0.2665 (2)	6.8 (2)
O(24)	0.3586 (3)	0.7515 (2)	-0.2685 (2)	6.0 (2)
C(25)	0.4675 (5)	0.5455 (3)	-0.2146 (3)	6.0 (2)
O(26)	0.2531 (3)	0.4139 (2)	0.0827 (2)	5.6 (2)
C(27)	0.3203 (5)	0.3205 (3)	0.0811 (3)	6.0 (3)
O(28)	0.1122 (2)	0.5825 (2)	0.0873 (1)	5.0 (1)
C(29)	0.1710 (5)	0.5840 (4)	0.1642 (3)	6.2 (3)
C(30)	0.1507 (5)	0.9869 (3)	-0.1427 (3)	6.8 (3)
C(31)	0.0011 (4)	0.7520 (4)	-0.0447 (3)	5.6 (2)
Molecule (II)				
O(1A)	0.7417 (2)	0.7955 (2)	0.3997 (1)	4.2 (2)
C(2A)	0.8449 (3)	0.8183 (2)	0.4406 (2)	4.1 (2)
C(3A)	0.9710 (4)	0.8443 (2)	0.3978 (2)	4.7 (2)
C(4A)	1.0692 (5)	0.8710 (3)	0.4433 (3)	5.3 (2)
C(5A)	1.0466 (4)	0.8694 (3)	0.5317 (3)	5.2 (2)
C(6A)	0.9193 (4)	0.8421 (2)	0.5738 (2)	4.7 (2)
C(7A)	0.8601 (4)	0.8270 (3)	0.6658 (3)	5.3 (2)
C(8A)	0.7220 (5)	0.7862 (4)	0.6723 (3)	5.6 (3)
C(9A)	0.6887 (3)	0.7957 (3)	0.5786 (2)	4.5 (2)
C(10A)	0.8218 (3)	0.8208 (2)	0.5264 (2)	4.1 (2)
C(11A)	0.6443 (3)	0.7017 (3)	0.5509 (2)	4.3 (2)
C(12A)	0.7230 (3)	0.6059 (3)	0.5827 (2)	4.5 (2)
C(13A)	0.7601 (3)	0.5568 (3)	0.5097 (2)	4.3 (2)
C(14A)	0.8244 (4)	0.4655 (3)	0.5102 (2)	4.8 (2)
C(15A)	0.8556 (4)	0.4375 (3)	0.4310 (3)	5.1 (2)
C(16A)	0.8235 (4)	0.4986 (3)	0.3545 (2)	4.9 (2)
C(17A)	0.7568 (3)	0.5894 (3)	0.3553 (2)	4.5 (2)
C(18A)	0.7252 (3)	0.6179 (2)	0.4338 (2)	4.0 (2)
C(19A)	0.6563 (3)	0.7133 (3)	0.4501 (2)	4.3 (2)
O(20A)	0.9889 (2)	0.8405 (2)	0.3121 (2)	5.7 (1)
C(21A)	1.1172 (5)	0.8662 (4)	0.2645 (3)	7.8 (3)
C(22A)	1.1563 (5)	0.8954 (4)	0.5785 (4)	7.2 (3)
O(23A)	0.9121 (3)	0.8408 (2)	0.7274 (2)	7.3 (2)
O(24A)	0.7470 (3)	0.5762 (2)	0.6568 (1)	6.0 (2)
C(25A)	0.8591 (6)	0.3971 (4)	0.5913 (3)	6.7 (4)
O(26A)	0.8538 (3)	0.4760 (2)	0.2750 (2)	5.7 (2)
C(27A)	0.9362 (5)	0.3896 (4)	0.2675 (3)	6.6 (3)
O(28A)	0.7165 (2)	0.6473 (2)	0.2805 (1)	5.1 (1)
C(29A)	0.8222 (5)	0.6918 (4)	0.2176 (3)	6.8 (3)
C(30A)	0.5878 (4)	0.8821 (3)	0.5562 (3)	5.6 (2)
C(31A)	0.5214 (4)	0.7308 (4)	0.4154 (3)	5.8 (2)

Only the positional parameters for the H atoms were included in the final cycles of refinement. They were assigned *B* values equal to the final isotropic *B* values for the atoms to which they were bonded. Not all parameters could be varied in any one cycle of refinement. Consequently, only one of the molecules in the asymmetric unit was varied per refinement cycle.

Table 2. *Fractional coordinates for H atoms with e.s.d.'s in parentheses*

	<i>x</i>	<i>y</i>	<i>z</i>
Molecule (I)			
H(4)	0.606 (3)	0.857 (2)	0.055 (2)
H(8)	0.372 (3)	0.916 (3)	-0.293 (2)
H(8)	0.313 (4)	1.026 (3)	-0.279 (3)
H(11)	0.125 (4)	0.823 (3)	-0.197 (2)
H(15)	0.416 (4)	0.413 (3)	-0.064 (2)
H(21)	0.418 (4)	0.703 (3)	0.236 (3)
H(21)	0.502 (4)	0.810 (3)	0.196 (3)
H(21)	0.556 (4)	0.716 (3)	0.165 (3)
H(22)	0.777 (4)	0.924 (3)	-0.145 (3)
H(22)	0.721 (4)	1.021 (3)	-0.125 (3)
H(22)	0.777 (4)	0.942 (3)	-0.047 (3)
H(25)	0.495 (4)	0.487 (3)	-0.215 (3)
H(25)	0.539 (4)	0.587 (3)	-0.224 (3)
H(25)	0.411 (4)	0.554 (3)	-0.270 (3)
H(27)	0.422 (4)	0.330 (3)	0.071 (3)
H(27)	0.285 (4)	0.276 (3)	0.135 (3)
H(27)	0.303 (4)	0.295 (3)	0.026 (3)
H(29)	0.217 (4)	0.527 (3)	0.184 (3)
H(29)	0.100 (4)	0.592 (3)	0.209 (3)
H(29)	0.217 (4)	0.647 (3)	0.160 (3)
H(30)	0.094 (4)	0.959 (3)	-0.080 (3)
H(30)	0.185 (5)	1.047 (4)	-0.145 (3)
H(30)	0.074 (4)	0.991 (3)	-0.186 (3)
H(31)	-0.040 (4)	0.719 (3)	-0.089 (3)
H(31)	-0.026 (4)	0.717 (3)	0.022 (3)
H(31)	-0.032 (4)	0.822 (3)	-0.053 (3)
Molecule (II)			
H(4A)	1.156 (4)	0.886 (3)	0.413 (2)
H(8A)	0.657 (4)	0.824 (3)	0.705 (2)
H(8A)	0.716 (4)	0.711 (3)	0.707 (3)
H(11A)	0.529 (4)	0.686 (3)	0.572 (2)
H(15A)	0.898 (4)	0.374 (3)	0.427 (2)
H(21A)	0.102 (5)	0.852 (4)	0.209 (3)
H(21A)	0.133 (6)	0.938 (4)	0.264 (3)
H(21A)	1.191 (5)	0.828 (4)	0.295 (3)
H(22A)	1.210 (5)	0.947 (4)	0.547 (3)
H(22A)	1.214 (5)	0.844 (4)	0.594 (3)
H(22A)	1.119 (4)	0.938 (3)	0.628 (3)
H(25A)	0.791 (5)	0.377 (4)	0.626 (3)
H(25A)	0.904 (4)	0.429 (3)	0.628 (3)
H(25A)	0.905 (4)	0.337 (3)	0.583 (3)
H(27A)	1.016 (4)	0.390 (3)	0.297 (3)
H(27A)	0.884 (4)	0.324 (3)	0.302 (3)
H(27A)	0.950 (4)	0.384 (3)	0.200 (3)
H(29A)	0.796 (4)	0.747 (3)	0.178 (3)
H(29A)	0.845 (5)	0.655 (4)	0.173 (3)
H(29A)	0.904 (5)	0.708 (4)	0.237 (3)
H(30A)	0.584 (4)	0.901 (3)	0.496 (3)
H(30A)	0.610 (4)	0.949 (3)	0.583 (3)
H(30A)	0.489 (4)	0.858 (3)	0.580 (3)
H(31A)	0.545 (4)	0.735 (3)	0.350 (3)
H(31A)	0.482 (4)	0.796 (3)	0.426 (3)
H(31A)	0.452 (4)	0.670 (3)	0.448 (3)

The function minimized by the least squares was $\sum w(|F_o| - |F_c|)^2$, where the weights (w) were derived from e.s.d.'s of observed intensities (Gilardi, 1973). All reflections were included in the refinement and the final R factors were $R = 0.089$ and $R_w = 0.062$. At the end of the refinement the standard deviation for an observation of unit weight was 1.4. Table 1 lists the final positional parameters and B_{eq} values for the non-H atoms. Table 2 lists the hydrogen coordinates.*

Discussion

The results of the X-ray study are illustrated in Fig. 1. Only one molecule has been drawn since both molecules in the asymmetric unit were found to have the same conformation. The molecule consists of three essentially planar segments: the two indenone moieties [excluding methyls C(29), C(30) and C(31)], and a four-atom plane made up of C(9), C(11), C(19) and O(1). The four-atom plane bridges the two indenone systems forming a dihedral angle of $59.8(5)^\circ$ with the AB ring system and an angle of $48.2(5)^\circ$ with the $A'B'$ ring system. For molecule (II) these angles are $56.5(5)$ and $43.4(5)^\circ$ respectively. Methoxy groups bonded to aromatic rings tend to be coplanar with the plane of the ring and except for C(29) the results of the present study follow this trend. C(29) is prevented from being coplanar with ring A by the H atoms on methyl C(31). The stereochemistry at the three chiral centers, C(9), C(11) and C(19) was established by this study and showed that the H atom at C(11) is *cis* with respect to C(31) and *gauche* with respect to C(30). Pertinent torsion angles are H(11)–C(11)–C(19)–C(31) at $-10.6(5)^\circ$ and H(11)–C(11)–C(19)–C(30) at $51.4(5)^\circ$. For molecule (II) these values are $-15.7(6)$ and $45.8(6)^\circ$. Bond lengths and angles for the two molecules in the asymmetric unit, illustrated in Fig. 2, have expected values. There is some strain in the fused ring system as evidenced by the long bonds to C(11). The C(11)–C(9) bond length is $1.563(5)$ Å in (I) and $1.557(5)$ Å in (II) and the C(11)–C(19) bond length is $1.574(5)$ Å in (I) and $1.567(5)$ Å in (II). The large external angles at C(6), C(13) and C(18) are common to indole moieties. The average C–H distance is $1.00(5)$ Å for both molecules. The molecules form pairs around centers of symmetry as can be seen in the packing diagram in Fig. 3. There are no intermolecular contacts between molecule (II) and its symmetry-equivalent molecules that are less than van der Waals distances. The only intermolecular contacts in the cell

less than 3.5 Å are either (I)⋯(I) or (I)⋯(II) approaches, e.g. C(4)⋯C(27) at $3.41(5)$, C(5)⋯C(27) at $3.43(5)$, C(4)⋯C(5) at $3.44(5)$ and C(12)⋯C(27A) at $3.48(5)$ Å.

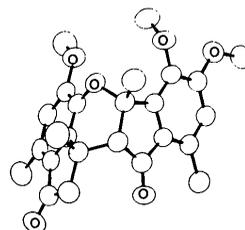


Fig. 1. ORTEP (Johnson, 1965) drawing of the methoxydiindenopyradione. Atoms are shown at their final refined positions.

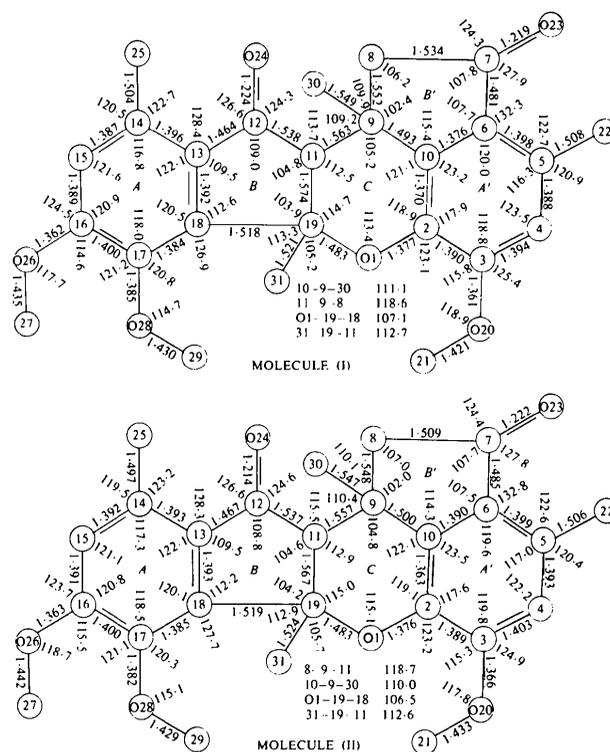


Fig. 2. Bond lengths (Å) and angles ($^\circ$). Standard deviations for both molecules are on the order of 0.005 Å for bond lengths and 0.3° for bond angles.



Fig. 3. Contents of one unit cell. Bonds have been blackened for both symmetry-equivalent positions for one molecule. The view is drawn with c horizontal, b approximately vertical and a coming up out of the paper.

* Lists of structure factors and anisotropic thermal parameters have been deposited with the British Library Lending Division as Supplementary Publication No. SUP 36324 (27 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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The Structure of *S,S'*-Methylenebis(L-cysteine) Monohydrochloride

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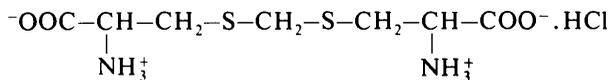
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Abstract

The crystal structure of the title compound has been determined from diffractometer data by Patterson and Fourier methods and refined to $R = 0.039$ for 1147 counter reflections: $C_7H_{15}N_2O_4S_2 \cdot Cl^-$, $M_r = 290.779$, $a = 10.405$ (6), $b = 5.074$ (3), $c = 11.972$ (6) Å, $\beta = 104.9$ (1)°, $V = 610.81$ (65) Å³, space group $P2_1$, $Z = 2$, $D_m = 1.60$, $D_c = 1.58$ Mg m⁻³, $F(000) = 304$, $\lambda(Cu K\alpha) = 1.54178$ Å, $\mu = 6.0203$ mm⁻¹. The monovalent cation arranges the polar groups in such a way as to enclose the chloride ion. Packing is determined by a three-dimensional network of hydrogen bonds involving the protonated amino and carboxyl groups. In the very strong $O_i'' \cdots O_k''$ [2.444 (8) Å] hydrogen bond, the H atom is disordered. Other contacts involve H atoms bonded to C atoms: $C^\alpha-H \cdots O = 3.165$ (8), 3.267 (8) Å and $C(\text{methylene})-H \cdots O = 3.377$ (9) Å. The two chemically equivalent parts of the molecule are not equivalent crystallographically and the most relevant difference between them is observed for the conformation around the $C^\beta-S^p$ bond, χ^2 being -169.3 (5) and -130.6 (5)°.

Introduction

As a continuation of a research programme on the conformational aspects of the sulphur-containing amino acid structures, the crystal structure analysis of *S,S'*-methylenebis(L-cysteine) monohydrochloride (djenkolic acid monohydrochloride),



has been carried out. The present study is connected with previous work on crystals of DL-homocystine itself and of its monohydrogen oxalate (Bigoli, Lanfranchi, Leporati, Nardelli & Pellinghelli, 1981).

Experimental

Attempts to obtain single crystals of L-djenkolic hydrochloride from aqueous solutions were unsuccessful, while clear, colourless needle-like single crystals of the title compound were obtained when attempting to prepare a copper(I) complex, *i.e.* by adding commercial L-djenkolic acid to a saturated solution of CuCl in 37% hydrochloric acid. From these

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