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The Crystal Structure of 3-Thiadodecanoic Acid

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3-Thiadodecanoic acid is triclinic with

a = 4.69, b = 5.10, c = 29.3 Å, $\alpha = 95^{\circ}8', \beta = 92^{\circ}36'$ and $\gamma = 114^{\circ}48'$.

The space group is $P\overline{1}$. The molecules are bent at the sulphur atom and linked together to dimers, which form a layer structure. The carbon chain tails of the molecules form a new type of chain packing with a nearly monoclinic subcell (M||).

Introduction

The structure determination of long-chain compounds is simplified if a heavy atom is introduced in the molecule. As our main interest lies in the naturally occurring unsubstituted compounds, different synthetic heavy-atom-containing compounds have been studied in order to find out if any of them are isomorphous with or structurally similar to the corresponding unsubstituted ones. The structure analysis of 3-thiadodecanoic acid was performed as part of this investigation, though the acid did not meet these requirements.

Experimental

The sulphur-containing acid was prepared by Smith & Hernestam (1954) by addition of thioglycollic acid to nonene-1. Crystals for the X-ray work were grown from glacial acetic acid. They had a melting point of 51.8 °C.

Rotation and Weissenberg photographs about the a- and b-axes were taken in a calibrated camera using Cu $K\alpha$ radiation.

Molecular formula: $C_{11}H_{22}O_2S$. Molecular weight: 218.35. Unit cell: triclinic $a = 4.69 \pm 0.05, \ b = 5.10 \pm 0.05, \ c = 29.3 \pm 0.03 \text{ Å};$ $\alpha = 95^{\circ}8' \pm 30', \ \beta = 92^{\circ}36', \ \gamma = 114^{\circ}48' \pm 30'.$ $d(001) = 29.04 \pm 0.15 \text{ Å}.$ Z = 2. $\varrho_c = 1.134 \pm 0.011 \text{ g.cm}^{-3},$ $\rho_{2} = 1.128 \text{ g.cm}^{-3}.$

The space group was considered to be $P\overline{1}$ which was later proved to be correct by the refinement.

The crystals decomposed slowly in the X-ray beam and different crystals had to be used for each zone of reflexions. The shape of the reflexions on non-zero layers was very unsatisfactory for intensity estimation. As, moreover, an attempt to scale the threedimensional data showed that different layers had apparently different temperature factors, it hardly seemed worth while using the full three-dimensional intensity data considering the rather limited accuracy reached in the analysis of phenoxymethylpenicillin (Abrahamsson *et al.*, 1962) where similar experimental difficulties were encountered.

*h*0*l* and 0*kl* intensities were estimated visually using the multiple-film technique. The Lorentz and polarization factors were applied but no absorption correction was made.



Fig. 1. Sharpened Patterson projection along the *b*-axis. Contours at arbitrary but equal levels.

Structure determination

From Patterson projections along the two short axes, with coefficients modified to correspond roughly to those from point atoms at rest (Abrahamsson & Maslen, 1962), the S–S vectors were easily recognized (Fig. 1). Electron-density projections from signs given by the sulphur position showed the whole molecule



Fig. 2. Sulphur phased electron density projection along the *b*-axis. Contours at arbitrary but equal levels. The correct structure is outlined.

without many spurious features (Fig. 2). A few rounds of difference syntheses reduced R to about 0.25 for both projections. The two sets of data were refined independently by anisotropic least-squares treatment to R about 0.15. An inspection of the vibrational parameters showed that those common for the two projections were similar enough to allow the two sets of data to be refined together. At this stage the hydrogen atoms on the methylene groups and the end methyl group were included in the structure-factor calculations with coordinates calculated with a C-H distance of 1.10 Å and a C-C-H angle of 110°. Isotropic temperature factors only were used for the hydrogen atoms (Table 3). The hydrogen parameters were not refined. Another few rounds of least-squares reduced R to 0.11 after which no further improvement occurred. The carbon shifts were still relatively large being up to half the standard deviations.

A Ferranti Mercury computer was used for the calculations with programs described by Mills &



Fig. 3. Final electron density projection along the *b*-axis. Contours given at intervals of 1 e.Å⁻² starting at 1 e.Å⁻² except for the sulphur atom which has heavy lines at intervals of 5 e.Å⁻².



Fig. 4. Final electron density projection along the *a*-axis. Contours given at intervals of $1 \text{ e.} \text{Å}^{-2}$ starting at $1 \text{ e.} \text{Å}^{-2}$ except for the sulphur atom which has heavy lines at intervals of $5 \text{ e.} \text{Å}^{-2}$.

Rollett (1961). For sulphur the scattering curve of Viervoll & Øgrim (1949) was used, for carbon and oxygen those of Berghuis *et al.* (1955) and for hydrogen that of McWeeny (1952).

Results and discussion

The final coordinates for the carbon and oxygen atoms are listed in Table 1, the vibrational parameters in Table 2. The electron-density projections along the short axes are given in Figs. 3 and 4. Observed and calculated structure factors are given in Table 5.

The bond distances and angles in the molecule are shown in Table 4. The average C–C distance is 1.53 ± 0.06 Å and the average C–C–C angle is $115 \pm 3^{\circ}$. This spread of the values is reasonable in comparison



Fig. 5. Molecular arrangement of 3-thiadodecanoic acid as seen along the *b*-axis.

Table 1. Coordinates of the heavier atoms

8 1.0602 1.5290 0.1	$352 \\ 422$
	422
O ₁ 0.7169 1.0425 0.0	
O_{0}^{-} 1.1962 1.3348 0.0	227
C, 0.9322 1.2762 0.0	454
C_{0}^{1} 0.8952 1.5134 0.0	802
C_{4}^{2} 0.7652 1.2091 0.1	577
C_{5}^{*} 0.8429 1.2230 0.2	082
C. 0.6012 0.9578 0.2	299
C, 0.6859 0.9797 0.2	798
C. 0.4570 0.7180 0.3	038
C. 0.5314 0.7363 0.3	538
$C_{10} = 0.3025 = 0.4880 = 0.3$	780
C_{11} 0.3743 0.5025 0.4	279
C_{12}^{11} 0.1509 0.2530 0.4	539

with the average standard deviations as estimated from the least-squares treatment (Ahmed & Cruickshank, 1953; Darlow, 1960). They are approximately independent of direction: $\sigma(S) = 0.006$ Å, $\sigma(O) = 0.01$ Å and $\sigma(C) = 0.03$ Å.

The molecular arrangement is shown in Fig. 5. The molecules are as usual for long-chain fatty acids linked together to dimers which form a layer structure.

Table 2. Vibrational parameters for the heavier atoms $(2^{-(h^2b_{11}+k^2b_{22}+l^2b_{33}+klb_{23}+hlb_{13})})$

	b_{11}	b_{33}	b_{22}	b_{23}	b_{13}
\mathbf{S}	0.12401	0.00175	0.06553	0.00728	-0.00869
C ₁	0.11829	0.00187	0.10548	0.02033	-0.00863
C_2	0.18265	0.00231	0.06165	0.00911	-0.01363
C_4	0.12958	0.00343	0.09262	0.01655	-0.01209
C ₅	0.13573	0.00271	0.14163	0.00937	-0.00706
C ₆	0.12972	0.00286	0.08282	-0.00068	-0.00165
C ₇	0.17026	0.00240	0.07390	0.01658	-0.00840
C ₈	0.17081	0.00229	0.10728	0.01452	-0.00894
C ₉	0.22527	0.00283	0.10427	0.01351	-0.01758
C ₁₀	0.18500	0.00269	0.09948	0.01134	-0.01290
C11	0.28786	0.00250	0.10809	0.00686	-0.01846
C ₁₂	0.24915	0.00287	0.15894	0.02520	-0.00200
01	0.11576	0.00283	0.09604	0.00082	-0.00497
O_2	0.12807	0.00228	0.14525	-0.00092	0.00001

Table 3. Parameters for the hydrogen atoms

	x/a	y/b	z/c	B
н,	0.9567	0.7233	0.0720	$5\cdot 8$ Å ²
H_2	0.6504	0.4724	0.0861	5.8
H_{3}	0.7618	1.0128	0.1398	5.8
H₄	0.5337	1.2029	0.1514	5.8
H_5	1.0773	1.2364	0.2146	5.8
H_6	0.8551	$1 \cdot 4231$	0.2258	5.8
н,	0.5858	0.7575	0.2119	5.9
H_8^{\cdot}	0.3678	0.9486	0.2246	5.9
H_9	0.9219	0.9862	0.2853	5.9
H_{10}	0.7125	1.1753	0.2976	5.9
H_{11}	0.4383	0.5137	0.2869	$5 \cdot 9$
$H_{12}^{}$	0.2194	0.6953	0.2969	$5 \cdot 9$
$H_{13}^{}$	0.7683	0.7569	0.3602	5.9
H_{14}	0.5540	0.9421	0.3706	$5 \cdot 9$
H_{15}	0.2774	0.2821	0.3608	6.0
H_{16}	0.0655	0.4667	0.3709	6.0
H ₁₇	0.6144	0.5330	0.4351	6.0
H_{18}	0.4029	0.7097	0.4449	6.0
H ₁₉	0.1222	0.0458	0.4369	6·0
H_{20}	-0.0893	0.2226	0.4467	6·0
н	0.9036	0.9697	0.4006	6.0

Table 4. Bond distances and angles in 3-thiadodecanoic acid

Bond		Angle	
C1-01	1·20 Å	O1-C1-O2	123°
$C_{1} - O_{2}$	1.37	$O_1 - C_1 - C_2$	114
$C_1 - C_2$	1.58	$O_{2} - C_{1} - C_{2}$	122
$C_2 - S$	1.74	$C_1 - C_2 - S$	112
S-C ₄	1.85	$\tilde{C_2}-\tilde{S-C_4}$	103
$C_4 - \tilde{C}_5$	1.50	$S - C_4 - C_5$	110
$C_5 - C_6$	1.58	$C_4 - \hat{C}_5 - \check{C}_6$	113
$C_6 - C_7$	1.48	$C_{5} - C_{6} - C_{7}$	112
$C_7 - C_8$	1.55	$C_6 - C_7 - C_8$	115
$C_8 - C_9$	1.47	$C_7 - C_8 - C_9$	115
C ₀ -C ₁₀	1.53	$C_{0}-C_{0}-C_{10}$	116
C10-C11	1.47	$C_{9} - C_{10} - C_{11}$	116
$C_{11} - C_{12}$	1.57	$\dot{C_{10}} - \dot{C_{11}} - \dot{C_{12}}$	118

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Table 5. Final structure factors for 3-thiadodecanoic acid

An asterisk marks changes of h and k. Each non-asterisk line contains l, $100F_o$ and $100F_c$

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-14 -13	309 - 6 37	500 - 490	- 22 - 21	319 - 110	401 95	11 13 14	- 594 318 558	- 629 305 652	15 16 16	103 87 - 28	115 36 - 122

- 122

There is a bend at the sulphur atom similar to that found in branched fatty acids (Abrahamsson, 1959), *cis*-cyclopropane fatty acids (Craven & Jeffrey, 1959) *cis*-unsaturated acids (Abrahamsson & Ryderstedt-Nahringbauer, 1962) and in ethyl stearate (Aleby, 1962). Bent long-chain molecules are thus rather common and found in structures where the earbon chain packing is disturbed by the presence of other than methylene groups along the molecule.

The chain part C_4-C_{12} has the usual planar zig-zag conformation. The maximum deviation from the best least-squares plane is 0.04 Å. The sulphur atom also lies in this plane whereas the bond $S-C_2$ is rotated 12° out of the plane.



Fig. 6. Idealized subcell $M \parallel$ found in 3-thiadodecanoic acid.

The regular chain parts pack side by side with all chain planes parallel. The dimensions of the subcell are calculated to be (average values):

$$a_s = 4.69, \ b_s = 4.56, \ c_s = 2.56 \text{ Å};$$

 $\alpha_s = 90^\circ, \ \beta_s = 84^\circ, \ \gamma_s = 122^\circ.$

The subcell is not similar to those previously known (Abrahamsson *et al.*, 1962; Abrahamsson & Ryderstedt-Nahringbauer, 1962) but can be regarded as a distortion of the theoretical monoclinic subcell deduced on close-packing considerations by Kitaigorodskii (1957). 3-Thiadodecanoic acid appears to be the first long-chain compound reported to show this packing. The idealized subcell, $M \parallel$, ($\beta_s = 90$) is shown in Fig. 6. The space group is $P2_1/m$ with 2 CH₂ groups per cell. The volume per CH₂ group (23.2 Å³) is similar to that in the common orthorhombic (O \perp) and triclinic ($T \parallel$) chain packing.

The average values for the hydrogen contacts in the subcell region were calculated from the positions in Table 3. The H_A-H_B distance is 2.68 Å whereas two alternate distances grouped around 2.52 Å and 2.93 Å are found for the H_A-H_C and H_B-H_D distances due to the actual deviations from the idealized subcell. The spread from these mean values is large (~ 0.1 Å) as is to be expected as the hydrogen positions were calculated from carbon positions with appreciable standard deviations.



Fig. 7. Hydrogen contacts in the methyl group contact planes. The directions of the a- and c-axes are indicated.

The angle of tilt of the chain axes to the (001) planes is 57° which is comparable to that in other simple long-chain compounds (Abrahamsson *et al.*, 1962). The methyl end group packing is illustrated in Fig. 7.

The carboxylic groups are as usual around centres of symmetry. The O-H···O distance is 2.74 Å. The atoms C₂, C₁, O₁ and O₂ lie in a plane (max. dev. 0.01 Å) which is separated 0.16 Å from that of the hydrogen bonded carboxylic group. A corresponding separation is found in α -pimelic acid (~0.3 Å, Kay & Katz, 1958) in oleic acid (0.35 Å, Abrahamsson & Ryderstedt-Nahringbauer, 1962) and in the *D*-form of 11-bromoundecanoic acid (1.17 Å, Larsson, 1962).

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A Direct Determination of the Molecular Structure of Prostaglandin F_{2-1}

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The tri-*p*-bromobenzoate of the methyl ester of prostaglandin $F_{2-1}(C_{42}H_{47}O_8Br_3)$ is orthorhombic with $a = 26 \cdot 14$, $b = 33 \cdot 93$ and $c = 4 \cdot 76$ Å. The space group is $P2_12_12_1$ with Z = 4. The structure was solved by heavy-atom methods and refined by three-dimensional anisotropic least-squares treatment to give the molecular structure and configuration of prostaglandin F_{2-1} , which belongs to a new class of hormones having smooth-muscle-stimulating and blood-pressure-depressing activity.

Introduction

A smooth-muscle-stimulating and blood-pressuredepressing activity was discovered in human semen by Goldblatt (1933) and by von Euler (1934, 1935) who also found a similar activity in sheep sperm and named the factor 'prostaglandin'.

In 1960 Bergström & Sjövall (1960) isolated two active crystalline compounds $PGE_1(C_{20}H_{34}O_5)$ and $PGF_{1-1}(C_{20}H_{36}O_5)$ from sheep prostate glands. Reduction of PGE₁ with borohydride (Bergström *et al.*, 1962) yielded PGF_{1-1} together with the isomeric compound PGF_{2-1}.

The tri-*p*-bromo- and tri-*p*-iodobenzoates of the methyl ester of PGF_{2-1} were prepared by Prof. S. Bergström who kindly provided them for an X-ray analysis in order to determine *ab initio* the molecular structures.

Experimental

The two benzoates crystallize as beautiful needles from ethanol. All crystals studied gave excellent X-ray photographs, which indicated that the two derivatives are isomorphous. The unit cell is orthorhombic with space group $P2_12_12_1$. The cell dimensions are:

	a	ь	с
Br-derivative	26·14 Å	33∙93 Å	4·76 Å
I-derivative	26·49	34∙39	4·76

Full three-dimensional data (Cu $K\alpha$ radiation) were collected for the bromo compound. Multiple-film Weissenberg photographs were recorded for the 0-2 layers about c (needle direction) and the 0-5 layers about a. Only hk0 intensities were estimated for the iodo derivative. The intensities were corrected for the Lorentz and polarization factors but not for absorption.

Structure determination

The initial work was done on the (001) projection. From a sharpened Patterson projection of the Iderivative, the relative positions of the heavy atoms of the four molecules per cell were obtained by simple vector addition and subtraction. The origin was then determined from the known symmetry of the projection. As expected the Patterson synthesis was completely dominated by I-I vectors (Fig. 1).

An electron-density projection showed most of the atoms of the molecule (Fig. 2) but also contained some spurious detail. This, however, was still present,