

The positional parameters are in good agreement with those given by Geller and Wood,⁸ who obtained $x_{\text{Rh}} = 0.144 \pm 0.003$ and $x_{\text{Si}} = 0.840 \pm 0.007$. However, the accuracy of the parameters is higher than in the earlier investigation.

In Table 1 the interatomic distances in RhSi are listed. A comparison of the distances with those given in the earlier investigation shows no significant differences.

Table 1. Interatomic distances in RhSi (FeSi-type).

Rh—6 Rh:	2.87 ₆
—7 Si:	2.47, 2.45 (3), 2.57 (3)
Si—7 Rh:	2.47, 2.45 (3), 2.57 (3)
—6 Si:	2.90

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The Crystal Structure of Tetramethylthiuram Disulphide

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Tetramethylthiuram disulphide, or bis(dimethylthiocarbamyl) disulphide, crystallizes in the space group $C2/c$ (No. 15) with four $[(\text{CH}_3)_2\text{NC}(\text{S})]_2\text{S}_2$ molecules per unit cell, of dimensions, $a = 9.66 \text{ \AA}$, $b = 9.95 \text{ \AA}$, $c = 11.85 \text{ \AA}$, $\beta = 99\frac{1}{2}^\circ$. With re-orientation of axes, these agree well with reported data.¹ The molecule possesses, by space group requirements, a twofold axis of symmetry.

The crystal structure has been solved, and refined by least squares analysis of 239 observed $h0l$, hhl , and $hk0$ reflections, estimated visually from zero-layer Weissenberg photographs taken with $\text{CuK}\alpha$ radiation. The reliability index, R , is 0.10 at the present stage. Anisotropic temperature factors were used for the sulphur atoms. The atomic coordinates, listed in Table 1, give 2.00 \AA for the length

Table 1. Atomic coordinates, in fractions of monoclinic cell edges, for tetramethylthiuram disulphide. Origin at a centre of symmetry.

	x	y	z
S ₁	-0.0283	0.3259	0.1651
S ₂	0.1955	0.1086	0.2033
C ₁	0.085	0.195	0.115
N	0.063	0.191	0.003
C ₂	-0.045	0.271	-0.077
C ₃	0.142	0.088	-0.056

of the S—S bond, with an e.s.d. of about 0.01 \AA . The bond angle of the sulphur atoms of the disulphide group is 104° , and the dihedral angle is 88° . The dimethyl-dithiocarbamate groups, excluding the hydrogen atoms, are planar within the experimental error.

Further refinement, based on more complete three-dimensional data, is in progress.

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