

métal du groupe IIa (Be, Mg ou Ca), et B, un métalloïde de la colonne V (N ou P). Ces composés possèdent une structure anti- α - Mn_2O_3 dont le paramètre varie avec la grosseur de chacun des types d'atome en présence, la distorsion de la structure étant principalement due à la différence d'électroaffinité entre A et B.

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The Crystal Structure of Monomethylammonium Perchlorate*

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The crystal structure of monomethylammonium perchlorate, $CH_3NH_3^+ClO_4^-$ has been determined by a three-dimensional X-ray analysis. The space group is $P2_1/n$ and the cell dimensions are: $a_0=10.59$, $b_0=7.67$, $c_0=12.87$ Å, $\beta=101^\circ 26'$; there are two molecules per asymmetric unit. The structure has been solved by the heavy-atom technique and refined by least-squares methods. The average Cl-O and N-C bond lengths are 1.43 and 1.52 Å respectively. Each of the nitrogen atoms has six oxygen neighbours with distances varying from 2.86 to 3.14 Å probably forming bifurcated hydrogen bonds. Each of the two independent $CH_3NH_3^+$ cations in the structure is surrounded by twelve oxygen atoms with distances less than 3.5 Å.

Introduction

Little information is available in the literature on the crystal structure of monomethylammonium salts. Only two of these compounds, namely the tetragonal monomethylammonium chloride (Hughes & Lipscomb, 1946) and the monomethylammonium aluminum sulphate alum in the room temperature cubic phase (Okaya, Ahmed, Pepinsky & Vand, 1957) and in the low temperature orthorhombic phase (Fletcher & Steeple, 1964), have been examined by X-ray methods during the last twenty years.

In the chloride, the C-N axis of the cations lies on the fourfold symmetry axis; since this is physically im-

possible with normal valences, the cations must exhibit either internal rotation or orientation disorder in the crystal. In cubic monomethylammonium alum, the $CH_3NH_3^+$ ions are statistically arranged around the cube body diagonal. Therefore in both structures, the exact coordination around the methylammonium ion is not considered, the only determination being that in orthorhombic 'alum'.

The present crystal structure determination on monomethylammonium perchlorate was undertaken with the purpose of contributing towards a better knowledge of the crystallochemistry of the $CH_3NH_3^+$ ion.

Experimental

Crystals of monomethylammonium perchlorate, prepared by neutralization of an alcoholic solution of methylamine with an aqueous solution of perchloric acid, were kindly supplied by Dr Marzocchi of the

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Laboratorio di Spettroscopia Molecolare of Florence University. The crystals, grown from alcoholic solution, are colourless needles elongated in the [010] direction, stable in air and in the X-ray beam.

The cell parameters determined from Weissenberg and rotation photographs were refined by a least-squares method, employing data from a powder diffractogram. The results are:

$$a_0 = 10.59 \pm 0.01, \quad b_0 = 7.67 \pm 0.01, \\ c_0 = 12.87 \pm 0.01 \text{ \AA}, \quad \beta = 101^\circ 26' \pm 15'.$$

From the systematically absent reflexions the space group was determined as $P2_1/n$ (C_{2h}^5). For eight unit $\text{CH}_3\text{NH}_3^+\text{ClO}_4^-$ in the cell the calculated density is $D_x = 1.71 \text{ g.cm}^{-3}$, in agreement with the observed value of 1.69 g.cm^{-3} measured by flotation in a mixture of methylene iodide and carbon tetrachloride. The linear absorption coefficient for the Cu $K\alpha$ wave-length is $\mu = 60.3 \text{ cm}^{-1}$.

For the collection of the intensities a crystal of prismatic shape with a cross section of $0.13 \times 0.06 \text{ mm}^2$ was used. The crystal was rotated about the b axis and equi-inclination Weissenberg photographs from $h0l$ to $h4l$ layers were taken with the use of Ni-filtered Cu $K\alpha$ radiation and the multiple film technique. In each layer, reflexions at high Bragg angles were too weak to be measured and were neglected. Of the 885 independent reflexions examined, 381 were in the observable range. Unidimensionally integrated intensities were measured with a microdensitometer and were roughly brought to a common relative scale on the basis of the exposure time of different layers. On account of the small cross section of the specimen, a sufficiently accurate correction was made for absorption by assuming the crystal to be a cylinder with a radius of 0.05 mm. Phillips's (1954) corrections for spot elongation and Lorentz-polarization factors to obtain the F^2 's were applied.

Determination and refinement of the structure

A three-dimensional Patterson synthesis was first calculated. The $(\frac{1}{2}, v, \frac{1}{2})$ Harker line showed only a relevant peak, and this allowed the assignment of the same y coordinate to the two independent chlorine atoms in the structure. The sets of x and z coordinates obtained from the $(u, \frac{1}{2}, w)$ Harker plane were then correctly correlated with the positions of the peaks due to the Cl(1)–Cl(2) vectors. The approximate coordinates of remaining non-hydrogen atoms were found from a three-dimensional 'heavy-atom' Fourier synthesis calculated without the coefficients whose signs had a low probability of being correct. The atomic coordinates were improved by a second electron density map and the discrepancy index $R = \sum ||F_o| - |F_c|| / \sum |F_o|$ for all the observed structure factors was at this stage 0.20.

A first refinement of the structure was performed with a least-squares program written by Albano, Bellon, Pompa & Scatturin (1963) for the IBM 1620 computer, using the block diagonal approximation and

individual isotropic thermal parameters. The weighting scheme suggested by Cruickshank (1961), $|w| = 1/(a + F_o + cF_o^2)$ with $a \approx 2F_{\min}$ and $c \approx 2/F_{\max}$, was employed for the structure factors and the unobserved reflexions were not included in the calculations. After three cycles the R index dropped to 0.14. At this stage the full-matrix least-squares refinement was undertaken with the program of Busing & Levy, adapted for the IBM 7090 computer by Stewart (1964). For the observed planes a weighting scheme like that specified by Hughes (1941) was used, with $|w| = 1$ for reflexions with $F_o \leq 4F_{\min}$; $|w| = 4F_{\min}/F_o$ for $F_o > 4F_{\min}$. To provide additional data in this new series of calculations the 'accidentally absent' reflexions were included with a value just below the lowest intensity observed in that part of the spectrum. These reflexions were weighted as follows: $|w| = 1$ for reflexions with $F_o > F_{\min}$, and $|w| = 0$ when $F_o \leq F_{\min}$. Ten intense low order reflexions suspected to be affected by secondary extinction were excluded from the last cycle of refinement; they are marked by an E in Table 1. After one isotropic and three anisotropic cycles, each followed by a proper rescaling of F_o 's, the R index for all observed structure factors reached the final value of 0.09.

The observed and calculated structure factors are listed in Table 1. For the calculations the scattering factor values from the self-consistent model as listed in *International Tables for X-ray Crystallography* (1962) for Cl, O, N and C were used.

Discussion of the structure

The atomic coordinates and anisotropic temperature factors, as from the last cycle of refinement, are given in Tables 2 and 3 with their standard deviations, estimated by the least-squares program using the formula:

$$\sigma_j = \sqrt{\frac{a^{jj} \sum w \Delta F^2}{m - n}},$$

where m is the number of reflexions, n is the number of parameters and a^{jj} is the jj element of the inverse matrix.

The high values reached by the temperature factors should indicate that in the structure the thermal motion is relevant and this fact may be responsible for the weakness of the diffraction effects at high Bragg angles.

The atomic arrangement projected along the b axis is shown in Fig. 1. The lists of bond lengths and bond angles with their standard deviations are included in Tables 4 and 5.

Both ClO_4 tetrahedra in the structure are slightly distorted. The O–Cl–O angles range from 105 to 112° . The average Cl–O distances are 1.44 and 1.42 \AA respectively for Cl(1) and Cl(2) tetrahedra. These values are in agreement with those reported in the literature for Cl–O distances in perchlorates (Truter, Cruickshank & Jeffrey, 1960; Truter, 1961; Coulter, Gantzel & McCullough, 1963).

Table 2. *Fractional atomic coordinates with their standard deviations*

| | x | 10 ⁴ σ(x) | y | 10 ⁴ σ(y) | z | 10 ⁴ σ(z) |
|-------|--------|----------------------|--------|----------------------|--------|----------------------|
| Cl(1) | 0.3818 | 4 | 0.6895 | 8 | 0.1503 | 3 |
| Cl(2) | 0.5469 | 5 | 0.3037 | 9 | 0.3911 | 4 |
| O(1) | 0.3866 | 13 | 0.8086 | 21 | 0.2403 | 9 |
| O(2) | 0.3539 | 12 | 0.7916 | 21 | 0.0533 | 10 |
| O(3) | 0.2847 | 15 | 0.5603 | 19 | 0.1493 | 14 |
| O(4) | 0.5023 | 13 | 0.6110 | 18 | 0.1556 | 12 |
| O(5) | 0.5764 | 13 | 0.1911 | 20 | 0.4844 | 11 |
| O(6) | 0.4278 | 15 | 0.3859 | 20 | 0.3910 | 13 |
| O(7) | 0.5383 | 14 | 0.1864 | 23 | 0.3014 | 12 |
| O(8) | 0.6416 | 13 | 0.4265 | 22 | 0.3906 | 12 |
| N(1) | 0.1883 | 15 | 0.6663 | 23 | 0.3615 | 11 |
| N(2) | 0.8715 | 12 | 0.3280 | 29 | 0.1068 | 12 |
| C(1) | 0.1016 | 20 | 0.7075 | 38 | 0.4410 | 16 |
| C(2) | 0.7744 | 22 | 0.3547 | 38 | 0.1766 | 18 |

The two independent N-C distances in the methylammonium ions are 1.54 and 1.51 Å. These values are somewhat larger than the expected normal single-bond lengths between carbon and quaternary nitrogen atoms: 1.49, 1.50 and 1.51 Å in spermine phosphate (Iitaka & Huse, 1965), 1.47 Å in tetramethylammonium perchlorate (McCullough, 1964), 1.47 Å in monomethylammonium chloride (Hughes & Lipscomb, 1946), 1.51 Å in monomethylammonium alum (Fletcher & Steeple, 1964); but the difference seems not to be significant considering their standard deviations.

The shortest distances among the nitrogen atoms of the methylammonium ions and oxygen atoms of ClO₄ tetrahedra are listed in Table 6. The directions of the N-H...O bonds cannot be deduced simply, but it

Table 3. *Anisotropic thermal parameters*

exp $[-(h^2\beta_{11} + k^2\beta_{22} + l^2\beta_{33} + 2hk\beta_{12} + 2hl\beta_{13} + 2kl\beta_{23})]$.
(σ) applies to the rightmost digit of the quantity in question.

| | β ₁₁ | β ₂₂ | β ₃₃ | β ₁₂ | β ₁₃ | β ₂₃ |
|-------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|
| Cl(1) | 0.0063 (6) | 0.0161 (16) | 0.0058 (4) | 0.0012 (8) | 0.0002 (3) | -0.0010 (7) |
| Cl(2) | 0.0060 (5) | 0.0217 (16) | 0.0050 (4) | 0.0005 (9) | -0.0004 (3) | 0.0006 (8) |
| O(1) | 0.0131 (17) | 0.0223 (45) | 0.0047 (10) | -0.0007 (21) | -0.0006 (10) | -0.0030 (17) |
| O(2) | 0.0130 (16) | 0.0370 (48) | 0.0028 (10) | -0.0004 (22) | 0.0001 (10) | 0.0053 (19) |
| O(3) | 0.0124 (19) | 0.0170 (44) | 0.0117 (15) | -0.0043 (25) | -0.0012 (14) | -0.0015 (20) |
| O(4) | 0.0098 (18) | 0.0199 (43) | 0.0103 (15) | 0.0019 (19) | 0.0009 (14) | -0.0039 (18) |
| O(5) | 0.0102 (16) | 0.0272 (40) | 0.0083 (13) | 0.0003 (21) | -0.0002 (11) | 0.0025 (22) |
| O(6) | 0.0093 (16) | 0.0272 (49) | 0.0154 (16) | 0.0058 (22) | 0.0011 (13) | 0.0017 (22) |
| O(7) | 0.0152 (19) | 0.0330 (49) | 0.0060 (14) | 0.0027 (26) | 0.0002 (13) | -0.0038 (25) |
| O(8) | 0.0072 (19) | 0.0467 (57) | 0.0085 (13) | -0.0035 (27) | 0.0000 (13) | -0.0001 (19) |
| N(1) | 0.0117 (19) | 0.0238 (50) | 0.0031 (10) | -0.0025 (31) | 0.0019 (10) | 0.0030 (22) |
| N(2) | 0.0058 (16) | 0.0502 (56) | 0.0055 (12) | -0.0052 (29) | -0.0001 (11) | -0.0008 (25) |
| C(1) | 0.0085 (22) | 0.0660 (88) | 0.0058 (16) | 0.0068 (36) | 0.0006 (14) | -0.0060 (32) |
| C(2) | 0.0106 (22) | 0.0415 (77) | 0.0110 (22) | 0.0032 (36) | 0.0065 (17) | 0.0009 (30) |

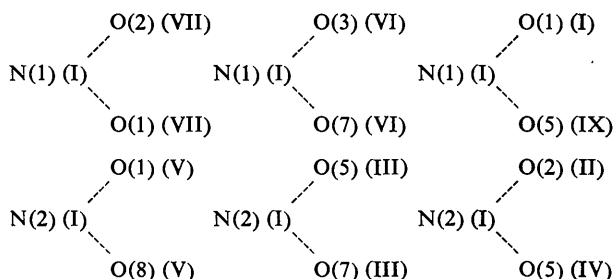
Table 4. *Bond lengths with standard deviations*

| | |
|---|--|
| (I) x, y, z | (VI) $\frac{1}{2}-x, \frac{1}{2}+y, \frac{1}{2}-z$ |
| (II) 1-x, 1-y, -z | (VII) $\frac{1}{2}-x, y-\frac{1}{2}, \frac{1}{2}-z$ |
| (III) $\frac{3}{2}-x, \frac{1}{2}+y, \frac{1}{2}-z$ | (VIII) $x-\frac{1}{2}, \frac{3}{2}-y, \frac{1}{2}+z$ |
| (IV) $\frac{1}{2}+x, \frac{1}{2}-y, -\frac{1}{2}+z$ | (IX) 1-x, 1-y, 1-z |
| (V) $\frac{3}{2}-x, -\frac{1}{2}+y, \frac{1}{2}-z$ | |
| Cl(1)(I) - O(1)(I) 1.47 ± 0.01 Å | Cl(2)(I) - O(5)(I) 1.46 ± 0.01 Å |
| - O(2)(I) 1.45 0.01 | - O(6)(I) 1.41 0.02 |
| - O(3)(I) 1.43 0.02 | - O(7)(I) 1.45 0.02 |
| - O(4)(I) 1.40 0.01 | - O(8)(I) 1.38 0.02 |
| N(1)(I) - C(1)(I) 1.54 ± 0.02 | N(2)(I) - C(2)(I) 1.51 ± 0.02 |

Table 5. *Bond angles in ClO₄ tetrahedra and their standard deviations*

| | |
|---------------------------------|--------------|
| O(1) (I) - Cl(1) (I) - O(2) (I) | 108.2 ± 0.9° |
| - O(3) (I) | 110.5 0.8 |
| - O(4) (I) | 109.9 0.8 |
| O(2) (I) - Cl(1) (I) - O(3) (I) | 110.1 0.8 |
| - O(4) (I) | 107.6 0.8 |
| O(3) (I) - Cl(1) (I) - O(4) (I) | 110.5 0.9 |
| O(5) (I) - Cl(2) (I) - O(6) (I) | 108.1 ± 0.8 |
| - O(7) (I) | 104.9 0.9 |
| - O(8) (I) | 111.8 0.8 |
| O(6) (I) - Cl(2) (I) - O(7) (I) | 111.2 0.9 |
| - O(8) (I) | 110.3 1.0 |
| O(7) (I) - Cl(2) (I) - O(8) (I) | 110.6 0.9 |

seems likely from stereochemical considerations that there are bifurcated hydrogen bonds in the following manner:



The resultant system of hydrogen bonding is rather complex.

Table 6. N—O distances less than 3.5 Å

| | | | |
|---------------------|--------|---------------------|--------|
| N(1) (I) — O(1) (I) | 3.06 Å | N(2) (I) — O(1) (V) | 2.91 Å |
| — O(1) (VII) | 3.08 | — O(2) (II) | 2.97 |
| — O(2) (VII) | 3.14 | — O(5) (IV) | 2.93 |
| — O(3) (VI) | 3.04 | — O(5) (III) | 3.11 |
| — O(5) (IX) | 3.06 | — O(7) (III) | 3.07 |
| — O(7) (VI) | 2.86 | — O(8) (V) | 3.08 |
| — O(3) (I) | 3.20 | — O(3) (II) | 3.49 |
| — O(6) (I) | 3.29 | — O(6) (IV) | 3.38 |
| — O(8) (IX) | 3.41 | | |

The interatomic distances less than 3.5 Å among methyl carbon atoms and the neighbours oxygen atoms are listed in Table 7. These distances appear to be normal, although C(1)(I) shows five contacts [with O(2)(VII), O(2)(VIII), O(8)(IX), O(7)(VI) and O(3)(VI)] and C(2)(I) two contacts [with O(2)(II) and O(7)(III)], which are slightly less than the sum of the van der Waals radii for a methyl group and oxygen, *i.e.* 3.4 Å. The environments of the two non-equivalent CH_3NH_3^+ ions are shown in Fig. 2. Each of them is surrounded by twelve oxygen atoms at distances less than 3.5 Å.

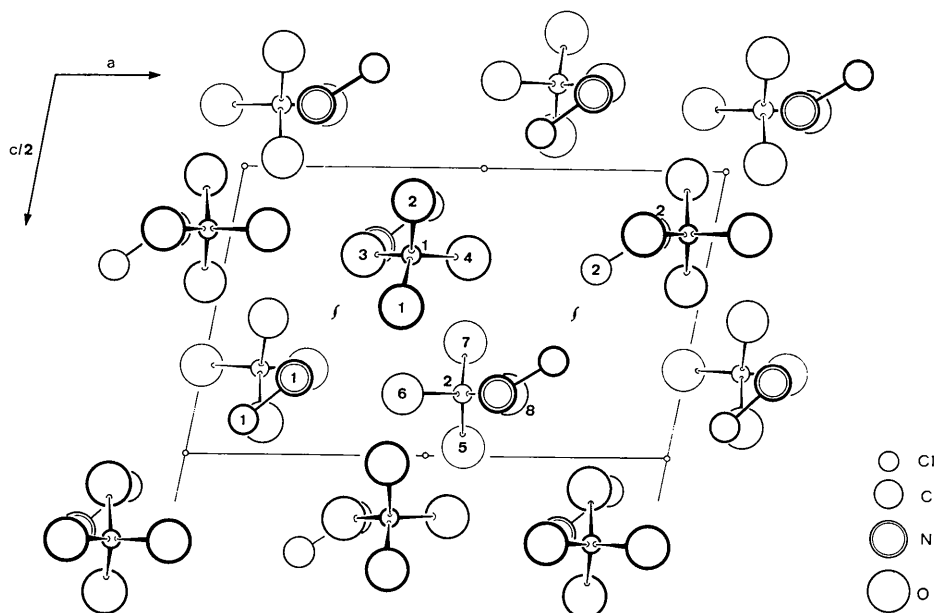


Fig. 1. Projection of the structure along the *b* axis.

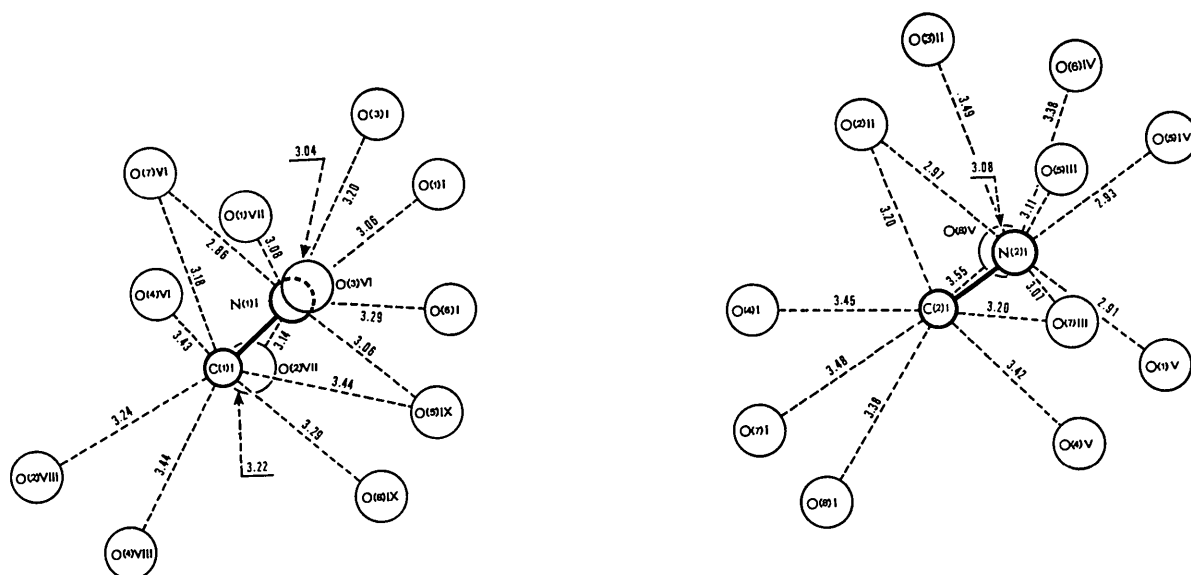


Fig. 2. The environments of the two non-equivalent $[\text{NH}_3\text{CH}_3]^+$ ions projected along the *b* axis. Distances in Å.

Table 7. Distances less than 3.5 Å between methyl carbon atoms and neighbouring atoms

| | | | |
|-----------------------|--------|----------------------|--------|
| C(1) (I) – O(2) (VII) | 3.22 Å | C(2) (I) – O(2) (II) | 3.20 Å |
| – O(2) (VIII) | 3.24 | – O(4) (V) | 3.42 |
| – O(3) (VI) | 3.27 | – O(4) (I) | 3.45 |
| – O(4) (VI) | 3.43 | – O(7) (III) | 3.20 |
| – O(4) (VIII) | 3.44 | – O(7) (I) | 3.48 |
| – O(5) (IX) | 3.44 | – O(8) (I) | 3.38 |
| – O(7) (VI) | 3.18 | | |
| – O(8) (IX) | 3.29 | | |

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A New Method of Locating Heavy Atoms Bound to Protein Crystals

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It is shown that heavy atoms bound isomorphously to protein crystals can be located by using a direct method (Sayre's equation) to phase X-ray diffraction reflections in centrosymmetric projections. This method has been tested successfully with three derivatives of carboxypeptidase A, one of which contains four heavy atoms per protein molecule. Quite possibly several heavy atoms can be located in this manner with greater ease and assurance than by difference Patterson methods alone and thus the method may be especially useful in the X-ray study of proteins larger than 125,000 molecular weight, where binding of more than one or two heavy atoms will probably be both common and necessary.

Multiple isomorphous replacement phasing of X-ray diffraction data is the method which has proven most useful for determining protein structures. Its application requires the preparation of several isomorphous heavy atom derivatives of the native protein and the determination of the positions of these heavy atoms in the derivative crystals. The difference Patterson synthesis (Green, Ingram & Perutz, 1954), which is used to locate heavy atoms when no estimate of the protein phases is available, is quite adequate if only one or two heavy atoms are bound per asymmetric unit. However, by this method the location of larger numbers of heavy atoms becomes increasingly difficult.

It is shown here that a direct phase determination using Sayre's equation may be applied to reflections in centrosymmetric zones to locate heavy atoms bound isomorphously to protein crystals. The method has

been successfully tested with three heavy atom derivatives of carboxypeptidase A_x (CPA_x), one of which contains eight mercury atoms per unit cell (four per molecule), and hence may prove to be a useful and powerful alternative to the difference Patterson syntheses for locating large numbers of heavy atoms.

The method

Sayre's (1952) equation expresses a relationship between the signs (phases) of certain structure factors for centrosymmetric structures of non-overlapping atoms:

$$s(F_{\mathbf{h}}) = s \sum_{\mathbf{k}} (F_{\mathbf{k}} F_{\mathbf{h}-\mathbf{k}}),$$

where $s()$ means 'sign of' and $F_{\mathbf{h}}$, $F_{\mathbf{k}}$ and $F_{\mathbf{h}-\mathbf{k}}$ are the structure factors of reflections $\mathbf{h}(=hkl)$, \mathbf{k} , and $\mathbf{h}-\mathbf{k}$. Using this equation, and assuming the signs of a few structure factors, the signs of other structure factors can be predicted and these in turn can be used to pre-

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