

The Determination of the Phases of the Structure Factors of Non-Centrosymmetric Crystals by the Method of Double Isomorphous Replacement*

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It is shown that the complete structure of a non-centrosymmetric crystal can be determined from the X-ray diffraction data obtained from the members of an isomorphous series in which two different parts of the structure can be varied independently. The structure so determined may be an enantiomorph of the true structure.

1. Introduction

Two crystals are said to be isomorphous if they have essentially the same structures, but are composed of chemically different atoms. (This includes the case of additional atoms, since these can be thought of as replacing vacancies, i.e. atoms of zero atomic number.) Examples of isomorphism are very common: KMnO_4 and BaSO_4 , Ag_3AsS_3 and Ag_3SbS_3 , benzene hexachloride and benzene hexabromide, tetraphenyl tin and tetraphenyl lead, etc. It often happens that the several hydrohalides of a complicated organic amine, or the various alkali metal salts of an organic acid, form series of isomorphous crystals. It has recently been discovered that certain proteins can crystallize in apparently identical fashion either with or without the substitution of mercury or silver atoms for the hydrogen of sulfhydryl groups, or with or without dye molecules bound to the protein molecules; these are also examples of isomorphism. The existence of such isomorphous pairs or series often greatly facilitates the determination of the structure of these crystals by X-ray diffraction methods.

A good account of how isomorphous replacement methods have been used for determining the structures of crystals is given in the new book by Lipson & Cochran (1953). These methods have been applied in the past with great success to centrosymmetric crystals.

It is often important to find the structure of a complicated organic molecule containing dozens, or even hundreds, of light atoms. This can be done when a crystal containing this molecule is capable of forming two different isomorphous series by the addition or substitution of heavy atoms in two different sets of positions in the unit cell. For example, suppose that a certain complicated sugar forms a set of isomorphous crystalline addition compounds with the alkali halides NaCl, NaBr and KCl. The intensities of the X-rays

diffracted by these three crystals can be used to define completely the structure of the sugar, as will be shown below, provided only that the arrangements of the alkali and halide atoms are sufficiently simple that their positions can be determined easily (see, however, the discussion in § 3). This latter problem, that of finding the positions of the heavy atoms, is usually soluble by the use of Patterson functions or related methods; in the discussion to follow, it is assumed that the arrangements of the heavy atoms have been found, and that the question of interest concerns the nature of the complicated arrangements of the light atoms in the crystal.

Bokhoven, Schoone & Bijvoet (1951; see especially the first full paragraph on p. 279) have outlined very briefly the method of double isomorphous replacement which is the subject of the present paper. They did not, however, mention the difficulties connected with the choice of origin and enantiomorphism, which are discussed at some length below in § 5. The resolution of these ambiguities is imperative for the solution of a non-centrosymmetric structure. The detailed development of the entire method has therefore been undertaken here.

2. Statement and solution of the problem

The structure factors, $F(hkl)$, of a crystal depend on the natures and positions of the atoms in one unit cell of the crystal according to the formula:

$$F(hkl) = \sum_{j=1}^N f_j(hkl) \exp 2\pi i(hx_j + ky_j + lz_j), \quad (1)$$

where the various symbols have their usual meanings. The sum on the right of (1) can be conveniently divided into terms corresponding to the atoms which do not change from one isomorphous crystal to another and other terms which correspond to atoms which do change. Call the constant part of the structure O and the variable part of the structure X . Then

$$F_{O+X} = F_O + F_X. \quad (2)$$

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The indices h , k and l have been omitted here, since no confusion is possible, and the subscripts indicate the parts of the structure to which each term corresponds. Inasmuch as $F(hkl)$ is, in general, a complex number, equation (2) is a vector equation in the complex plane, and can be represented by a diagram such as Fig. 1.

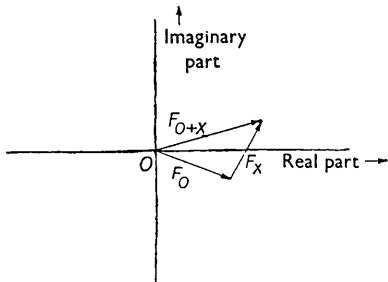


Fig. 1. Vector addition of the structure factors of the invariable and variable parts of a crystal structure.

It is assumed in what follows that the experimental quantities $|F(hkl)|$ are known for the various members of the isomorphous series. For definiteness, assume that, for each h , k , l , the magnitudes $|F_O|$, $|F_{O+X}|$ and $|F_{O+M}|$ are known, where X and M refer to two different isomorphous additions to the structure O . Assume also that the structures X and M have been found, and that the complex numbers F_X and F_M have been computed from them for each h , k , l . The problem is to find the complex numbers $F_O(hkl)$ from which the structure O can be computed directly by Fourier methods.

The solution for F_O is easily obtained graphically; the procedure is illustrated in Fig. 2. Vectors $-F_X$

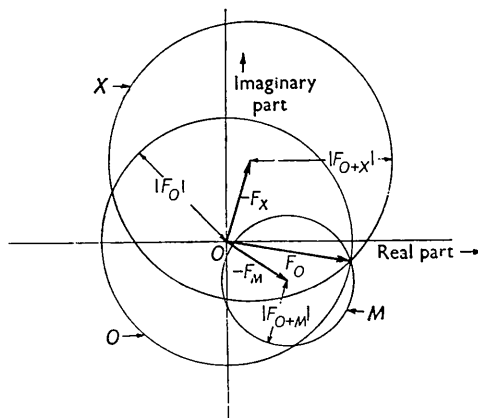


Fig. 2. Solution of the vector equations for F_O from a knowledge of $|F_O|$, $|F_{O+M}|$, $|F_{O+X}|$, F_M and F_X .

and $-F_M$ are drawn from the origin of the complex plane, and a circle (marked O) with radius equal to $|F_O|$ is described about the origin as center, a circle (marked X) of radius $|F_{O+X}|$ is centered at the end of the vector $-F_X$ and another circle (marked M) of radius $|F_{O+M}|$ is centered at the end of vector $-F_M$.

The three circles intersect at a point which is the end of vector F_O . It is to be noted that a knowledge of F_X , $|F_O|$ and $|F_{O+X}|$ alone gives two solutions for F_O , one for each of the two intersections of circles O and X ; similarly, there are two solutions for F_O obtainable from a knowledge of F_M , $|F_O|$ and $|F_{O+M}|$ alone, corresponding to the two intersections of circles O and M . The correct solution for F_O is the one common to these pairs of solutions, i.e. the one corresponding to the point where all three circles O , M and X intersect.

Analytically, each complex number F can be written $F = A + iB$. The known data are $|F_O|^2 = A_O^2 + B_O^2$, $|F_{O+X}|^2 = A_{O+X}^2 + B_{O+X}^2$, $|F_{O+M}|^2 = A_{O+M}^2 + B_{O+M}^2$, $F_X = A_X + iB_X$ and $F_M = A_M + iB_M$. We wish to find A_O and B_O . Now,

$$\begin{aligned} |F_{O+X}|^2 &= (A_O + A_X)^2 + (B_O + B_X)^2 \\ &= A_O^2 + B_O^2 + A_X^2 + B_X^2 + 2A_O A_X + 2B_O B_X \end{aligned}$$

or

$$|F_{O+X}|^2 = |F_O|^2 + |F_X|^2 + 2A_O A_X + 2B_O B_X.$$

This gives:

$$P_X = |F_{O+X}|^2 - |F_X|^2 - |F_O|^2 = 2A_X A_O + 2B_X B_O. \quad (3)$$

Similarly,

$$P_M = |F_{O+M}|^2 - |F_M|^2 - |F_O|^2 = 2A_M A_O + 2B_M B_O. \quad (4)$$

All the quantities in (3) and (4) are known, except A_O and B_O ; consequently these two equations can be solved simultaneously, as follows:

Let

$$\Delta = 4(A_X B_M - A_M B_X),$$

then

$$A_O = \frac{2}{\Delta} \begin{vmatrix} P_X & B_X \\ P_M & B_M \end{vmatrix} \quad \text{and} \quad B_O = \frac{2}{\Delta} \begin{vmatrix} A_X & P_X \\ A_M & P_M \end{vmatrix}. \quad (5)$$

From the values of A_O and B_O obtained by means of (5) it is possible to write immediately $F_O = A_O + iB_O$.

The analytical method just outlined corresponds to locating the end of the vector F_O from a knowledge of its projections on the directions of the two vectors F_M and F_X . The geometry is thus quite different from that of the most convenient graphical method, and the effect of experimental errors in the intensity measurements will not be the same. The analytical method always gives a unique determination of A_O and B_O (except when F_M and F_X are parallel), while the graphical method may not, since there may be no point at which the three circles O , M and X intersect. If the graphical method fails in this way, the best position for the end of the vector F_O would be in the center of the small curved triangle where the three circles nearly intersect; an averaging of errors is also required in the analytic method, because $A_O^2 + B_O^2$ may differ from the observed value of $|F_O|^2$, but the averaging is different.

3. Situations to be avoided

It is easy to see that, if F_M and F_X are parallel or anti-parallel (i.e. collinear in Fig. 2), there will be two points where the three circles O , M and X intersect, corresponding to two solutions for F_O . In this case the analytical method just described fails to give a unique solution for F_O , since $\Delta = 0$. This situation may occur occasionally in any one isomorphous series for special values of h , k , l , but such a sporadic ambiguity usually does not seriously hamper a structure determination. If, on the other hand, F_M and F_X are systematically collinear, no structure determination by this method is possible for a non-centrosymmetric crystal. For instance, this last is true for every h , k , l , if structures M and X are both centrosymmetric about the same point in structure O , or centrosymmetric about points which differ in coordinates by integral numbers of halves; and there are several other special situations which make F_M and F_O collinear for a hopelessly large proportion of the h , k , l , combinations. Such situations cannot occur if one of the structures M or X is non-centrosymmetric, or if they are both centrosymmetric, but about points which differ in coordinates by irrational numbers. Thus, in the case of the series of alkali halide-sugar addition compounds mentioned in § 1, if each unit cell contains only one alkali metal atom and one halogen atom, the coordinates of these must differ by irrational numbers, in order to make the structure determination possible, since single atoms are always centrosymmetric. (Of course, if the crystal is centrosymmetric—a case not under consideration here—the structure determination is usually possible, even though F_M and F_X are required by symmetry to be collinear.)

4. Solution of a hypothetical structure

These methods were applied to a hypothetical one-dimensional structure consisting of point atoms, of unit scattering power for X-rays, arranged along a line. All atomic coordinates are multiples of one-twelfth of the period, so that the F 's are periodic functions of the order of 'reflection', h , with a period of twelve, and the relationship $F(h) = F^*(12n-h)$ holds, where the asterisk indicates the complex conjugate and n is an integer. It is therefore only necessary to study the first six orders of reflection. The structure O was taken to consist of six atoms, while the two different structures M and X were each taken to have three atoms. None of these structures has a center of symmetry. The atomic coordinates appear in Table 1, and the various values of F , $|F|$ and $|F|^2$, together with the phase angle α , from $F = |F| \exp(i\alpha)$, are presented in Table 2.

A test of the graphical method is illustrated in Fig. 3. For each order of reflection h , two diagrams appear in the figure; one shows the determination of the two solutions for F_O found by the use of F_M ,

Table 1. *Coordinates of unit point atoms in hypothetical test structures*

Structure O :

$$\text{Atoms at } x = \frac{1}{12}, \frac{2}{12}, \frac{3}{12}, \frac{7}{12}, \frac{10}{12}, \frac{11}{12}$$

$$F_O(h) = 2 \cos 2\pi \frac{h}{12} + 2 \cos 2\pi \frac{h}{6} + (i)^h + (-1)^h \exp\left(2\pi i \frac{h}{12}\right)$$

Structure M :

$$\text{Atoms at } x = 0, \frac{5}{12}, \frac{8}{12}$$

$$F_M(h) = 1 + (-1)^h \exp\left(-2\pi i \frac{h}{12}\right) + \exp\left(-2\pi i \frac{h}{3}\right)$$

Structure X :

$$\text{Atoms at } x = \frac{4}{12}, \frac{6}{12}, \frac{9}{12}$$

$$F_X(h) = \exp\left(2\pi i \frac{h}{3}\right) + (-1)^h + (-i)^h$$

Structure $O+M$:

Atoms at positions of structures O and M

$$F_{O+M}(h) = F_O(h) + F_M(h)$$

Structure $O+X$:

Atoms at positions of structures O and X

$$F_{O+X}(h) = F_O(h) + F_X(h)$$

$|F_{O+M}|$ and $|F_O|$, the other shows the two solutions found by the use of X , instead of M . Comparing these two diagrams, it is always found that one solution for F_O is common to both and has the same phase angle α as appears in Table 2. It is easy to carry out the computations of the analytical method and to show that it, too, gives the values of F_O in Table 2.

5. Ambiguities and their removal

In an actual application of this method of double isomorphous replacement, some complications arise which have not yet been mentioned and which have to do with the indeterminacy of the origin to which a structure is referred and with the near impossibility of distinguishing between two enantiomorphs by X-ray diffraction. These difficulties will be discussed using the structures O , M and X of Table 1. Structure M , for instance, must be discovered by comparing the Patterson functions of structures O and $O+M$, since a crystal of structure M alone cannot exist. This process results in a knowledge of the interatomic vectors in M , and from this it is usually possible to find the relative positions of its atoms, except that either enantiomorph will have the same interatomic vectors and that any origin can be used. Thus, starting from the values of $|F_O|^2$ and $|F_{O+M}|^2$ of Table 2, it could be found that structure M consists of three atoms in a row, the two end ones being, respectively, $3/12$ and $4/12$ of the period from the middle one, but there would be no way to tell which end should be in the direction of increasing X . (This description of structure M corresponds to choosing the origin at $x = 8/12$ and using the atoms at $x = 5/12, 8/12$ and 1 in the descrip-

Table 2. Values of F , $|F|$, $|F|^2$ and α for the structures O , M , X , $O+M$ and $O+X$ of Table 1
$$(F = |F|e^{i\alpha} = A + iB)$$

h	0	1	2	3	4	5	6
$F_O(h)$	6	$\frac{1}{2}(2+\sqrt{3})+\frac{1}{2}i$	$-\frac{1}{2}+\frac{1}{2}\sqrt{3}i$	$-2-2i$	$-\frac{3}{2}+\frac{1}{2}\sqrt{3}$	$\frac{1}{2}(2-\sqrt{3})+\frac{1}{2}i$	-2
$\alpha_O(h)$ ($^\circ$)	0	15	120	225	150	75	180
$ F_O(h) $	6	$\frac{1}{2}\sqrt{2}(\sqrt{3}+1)$	1	$2\sqrt{2}$	$\sqrt{3}$	$\frac{1}{2}\sqrt{2}(\sqrt{3}-1)$	2
$ F_O(h) ^2$	36	$2+\sqrt{3}$	1	8	3	$2-\sqrt{3}$	4
$F_M(h)$	3	$-\frac{1}{2}(\sqrt{3}-1)-\frac{1}{2}(\sqrt{3}-1)i$	1	$2+i$	$-\sqrt{3}i$	$\frac{1}{2}(\sqrt{3}+1)+\frac{1}{2}(\sqrt{3}+1)i$	1
$\alpha_M(h)$ ($^\circ$)	0	225	0	26.56	270	45	0
$ F_M(h) $	3	$\frac{1}{2}\sqrt{2}(\sqrt{3}-1)$	1	$\sqrt{5}$	$\sqrt{3}$	$\frac{1}{2}\sqrt{2}(\sqrt{3}+1)$	1
$ F_M(h) ^2$	9	$2-\sqrt{3}$	1	5	3	$2+\sqrt{3}$	1
$F_X(h)$	3	$-\frac{3}{2}-\frac{1}{2}(2-\sqrt{3})i$	$-\frac{1}{2}-\frac{1}{2}\sqrt{3}i$	i	$\frac{3}{2}+\frac{1}{2}\sqrt{3}i$	$-\frac{3}{2}-\frac{1}{2}(2+\sqrt{3})i$	1
$\alpha_X(h)$ ($^\circ$)	0	185.11	240	90	30	231.21	0
$ F_X(h) $	3	$\sqrt{4-\sqrt{3}}$	1	1	$\sqrt{3}$	$\sqrt{4+\sqrt{3}}$	1
$ F_X(h) ^2$	9	$4-\sqrt{3}$	1	1	3	$4+\sqrt{3}$	1
$F_{O+M}(h)$	9	$\frac{3}{2}+\frac{1}{2}(2-\sqrt{3})i$	$\frac{1}{2}+\frac{1}{2}\sqrt{3}i$	$-i$	$-\frac{3}{2}-\frac{1}{2}\sqrt{3}i$	$\frac{3}{2}+\frac{1}{2}(2+\sqrt{3})i$	-1
$\alpha_{O+M}(h)$ ($^\circ$)	0	5.11	60	270	210	51.21	180
$ F_{O+M}(h) $	9	$\sqrt{4-\sqrt{3}}$	1	1	$\sqrt{3}$	$\sqrt{4+\sqrt{3}}$	1
$ F_{O+M}(h) ^2$	81	$4-\sqrt{3}$	1	1	3	$4+\sqrt{3}$	1
$F_{O+X}(h)$	9	$\frac{1}{2}(\sqrt{3}-1)+\frac{1}{2}(\sqrt{3}-1)i$	-1	$-2-i$	$\sqrt{3}i$	$-\frac{1}{2}(\sqrt{3}+1)-\frac{1}{2}(\sqrt{3}+1)i$	-1
$\alpha_{O+X}(h)$ ($^\circ$)	0	45	180	206.56	90	225	180
$ F_{O+X}(h) $	9	$\frac{1}{2}\sqrt{2}(\sqrt{3}-1)$	1	$\sqrt{5}$	$\sqrt{3}$	$\frac{1}{2}\sqrt{2}(\sqrt{3}+1)$	1
$ F_{O+X}(h) ^2$	81	$2-\sqrt{3}$	1	5	3	$2+\sqrt{3}$	1

Table 3. Values of F , α , F and F^2 for structures M' and X'

h	0	1	2	3	4	5	6
$F_{M'}(h)$	3	$\frac{1}{2}+\frac{1}{2}(2-\sqrt{3})i$	$-\frac{1}{2}+\frac{1}{2}\sqrt{3}i$	$2-i$	$\frac{3}{2}-\frac{1}{2}\sqrt{3}i$	$\frac{1}{2}+\frac{1}{2}(2+\sqrt{3})i$	1
$\alpha_{M'}(h)$ ($^\circ$)	0	15	120	333.44	330	75	0
$ F_{M'}(h) $	3	$\frac{1}{2}\sqrt{2}(\sqrt{3}-1)$	1	$\sqrt{5}$	$\sqrt{3}$	$\frac{1}{2}\sqrt{2}(\sqrt{3}+1)$	1
$ F_{M'}(h) ^2$	9	$2-\sqrt{3}$	1	5	3	$2+\sqrt{3}$	1
$F_{X'}(h)$	3	$\frac{3}{2}+\frac{1}{2}(2-\sqrt{3})i$	$-\frac{1}{2}-\frac{1}{2}\sqrt{3}i$	$-i$	$\frac{3}{2}+\frac{1}{2}\sqrt{3}i$	$\frac{3}{2}+\frac{1}{2}(2+\sqrt{3})i$	1
$\alpha_{X'}(h)$ ($^\circ$)	0	5.11	240	270	30	51.21	0
$ F_{X'}(h) $	3	$\sqrt{4-\sqrt{3}}$	1	1	$\sqrt{3}$	$\sqrt{4+\sqrt{3}}$	1
$ F_{X'}(h) ^2$	9	$4-\sqrt{3}$	1	1	3	$4+\sqrt{3}$	1

tion of Table 1, with x increasing either to the right or left.) It might well be decided to use the coordinates $x = 0, 3/12, 8/12$ for the atoms in M , and this would be a 'correct' structure determination, as this phrase is usually used; this structure will be called M' . In a similar way, structure X might be 'correctly' determined to have atoms at $x = 0, 3/12, 10/12$; this structure will be called X' . It will be noted that structure M' is the enantiomorph of M , while structure X' is congruent to X ; also that the origins of M' and X' are not properly chosen with respect to one another. The values of F , α , $|F|$ and $|F|^2$ for structures M' and X' appear in Table 3. ($|F|$ and $|F|^2$ are, of course, the same as in Table 2.)

Since structures M' and X' are referred to different origins and correspond to different enantiomorphs, the solution for F_O cannot be obtained directly from the values of $F_{M'}$ and $F_{X'}$ by the use of the methods previously described. However, as will be shown, it is possible to extend these methods so as to obtain a 'correct' set of values for F_O , in the sense that these values will correspond to structure O or its enantiomorph and that structure O may be referred to a different origin from that in the original description.

First, consider the effect on the values of $F(hkl)$ caused by changing the origin to which the crystal structure is referred. Suppose the new origin has coordinates x_0, y_0, z_0 in the original coordinate system, and let $F'(hkl)$ be the value of $F(hkl)$ when the new origin is used.

If the complex number $F(hkl)$ is written in the form $F = |F| \exp(i\alpha)$, then the value of $F'(hkl)$ is

$$F'(hkl) = |F(hkl)| \exp i[\alpha(hkl) - 2\pi(hx_0 + ky_0 + lz_0)]. \quad (6)$$

This is the same as saying that the magnitudes of the F 's are not changed by a change of origin, but the phase angles α (expressed in radians) are reduced by $2\pi(hx_0 + ky_0 + lz_0)$. For a one-dimensional structure, the reduction in α is $2\pi hx_0$.

Second, the effect of changing from a structure to its enantiomorph is most easily expressed by changing all coordinates x_j, y_j, z_j into negatives: $-x_j, -y_j, -z_j$. This clearly changes each F into F^* and therefore changes α into $-\alpha$.

Let us now attempt to find the 'correct' values of F_O for the hypothetical one-dimensional structure of Table 1, starting with the measured values of $|F_O|$,

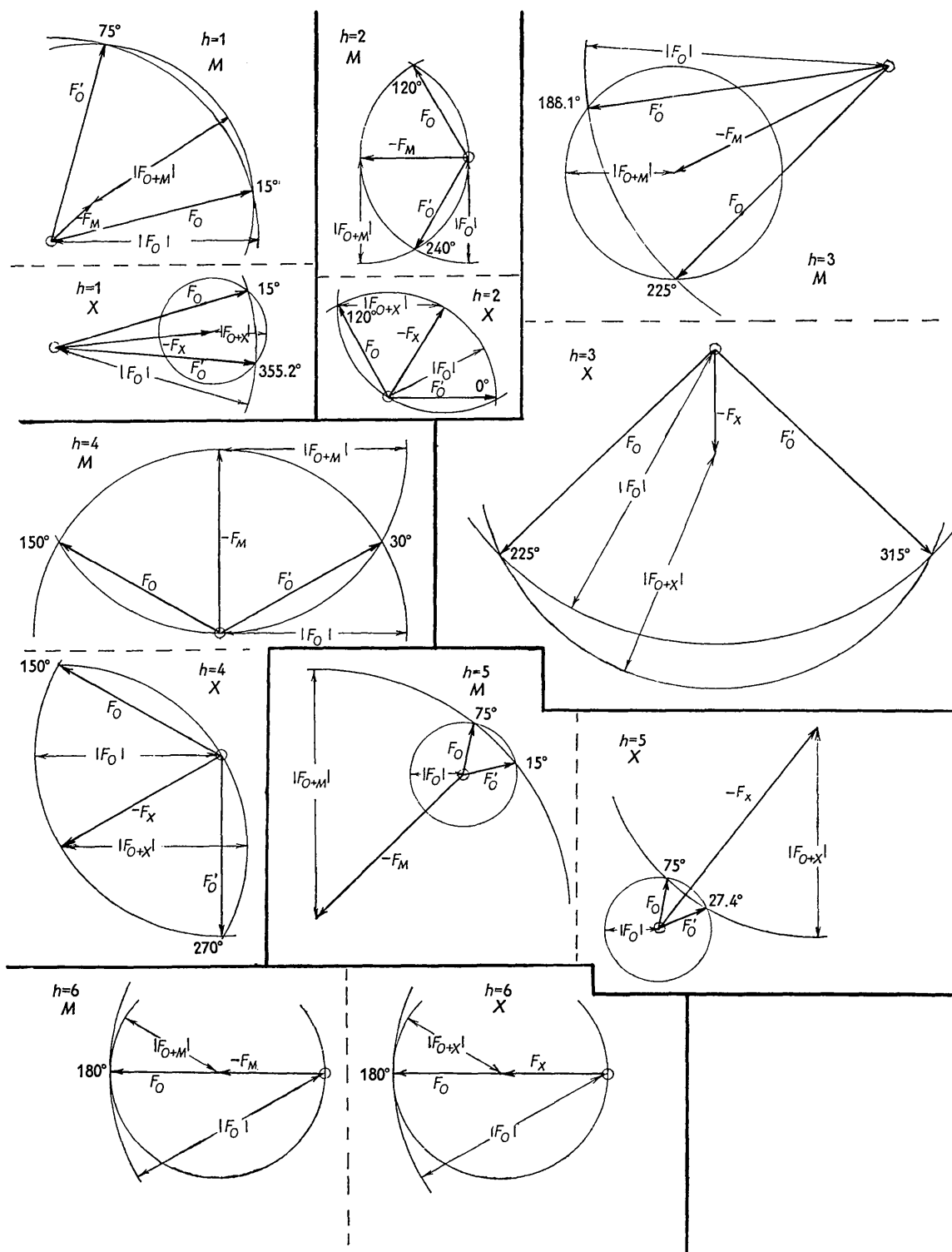


Fig. 3. Test of the double isomorphous replacement method of phase determination.

$|F_{O+M}|$, $|F_{O+X}|$ and the values of F_M and F_X found arbitrarily and listed in Table 3; we shall keep in mind, however, that structures M' and X' may be

referred to different origins, and may correspond to different enantiomorphs.

The first step is to find the two values of $\alpha_0(h)$

Table 4. Values of α_0 found from $|F_O|$, $|F_{O+M}|$ and $F_{M'}$ and from $|F_O|$, $|F_{O+X}|$ and $F_{X'}$.(All values of α are in degrees)

	(a)						
h	0	1	2	3	4	5	6
$ \alpha_O - \alpha_{M'} $	0	150	120	198.44	240	30	180
$\alpha_{M'}$	0	15	120	333.44	330	75	0
$\alpha_{M'} + \alpha_O - \alpha_{M'} = \alpha_0^i$	0	165	240	171.88	210	105	180
$\alpha_{M'} - \alpha_O - \alpha_{M'} = \alpha_0^{iv}$	0	225	0	135	90	45	180
	(b)						
$ \alpha_O - \alpha_{X'} $	0	189.89	240	135	120	203.79	180
$\alpha_{X'}$	0	5.11	240	270	30	51.21	0
$\alpha_{X'} + \alpha_O - \alpha_{X'} = \alpha_0^{ii}$	0	195	120	45	150	255	180
$\alpha_{X'} - \alpha_O - \alpha_{X'} = \alpha_0^{iv}$	0	175.22	0	135	270	207.42	180
	(c)						
$\alpha_0^i - \alpha_0^{ii}$	0	330	120	126.88	60	210	0
$\alpha_0^i - \alpha_0^{iv}$	0	349.78	240	36.88	300	257.58	0
$\alpha_0^{ii} - \alpha_0^{iv}$	0	30	240	90	300	150	0
$\alpha_0^i - \alpha_0^{iv}$	0	49.78	0	0	180	197.58	0
	(d)						
$\alpha_0^i + \alpha_0^{ii}$	0	0	0	216.88	0	0	0
$\alpha_0^i + \alpha_0^{iv}$	0	340.22	240	306.88	120	312.42	0
$\alpha_0^{ii} + \alpha_0^{iv}$	0	60	120	180	240	300	0
$\alpha_0^i + \alpha_0^{iv}$	0	40.22	0	270	0	252.42	0
	(e)						
$F_{X'''}(h)$	3	$\frac{1}{2}\sqrt{3} + \frac{1}{2}(2\sqrt{3}-1)i$	$-\frac{1}{2} - \frac{1}{2}\sqrt{3}i$	$-i$	$-\frac{3}{2} - \frac{1}{2}\sqrt{3}i$	$-\frac{1}{2}\sqrt{3} - \frac{1}{2}(2\sqrt{3}+1)i$	1
$\alpha_{X'''}$	0	54.89	240	270	210	248.79	0
$ \alpha_O - \alpha_{X'''} $	0	189.89	240	135	120	203.79	180
$\alpha_{X'''} + \alpha_O - \alpha_{X'''} = \alpha_0^{iv}$	0	244.78	120	45	330	42.58	180
$\alpha_{X'''} - \alpha_O - \alpha_{X'''} = \alpha_0^{iv}$	0	225	0	135	90	45	180

that correspond to each combination $|F_O|$, $|F_{O+M}|$, $F_{M'}$ and $|F_O|$, $|F_{O+X}|$, $F_{X'}$. This can be done either graphically, as in Fig. 3, or analytically. It is easily shown that:

$$\cos(\alpha_O - \alpha_M) = \frac{1}{2|F_O| |F_M|} [|F_{O+M}|^2 - |F_M|^2 - |F_O|^2], \quad (7)$$

where α_O and α_M are the phase angles of F_O and F_M , respectively. From (7), the magnitude of $\alpha_O - \alpha_M$ can be found, but not its sign; if α_M is known, this gives two values of α_O —the same two values to be found by the graphical method. Formula (7) applies equally well, of course, if M is everywhere replaced by X . Table 4 presents the results of calculations based on this method.

In Table 4 (a), the first row lists the values of $|\alpha_O - \alpha_{M'}|$ for each h from 0 to 6; the second row lists the values of $\alpha_{M'}$, from Table 3, and the third and fourth rows list the values of α_0^i and α_0^{iv} which are the two values of α_0 compatible with the values of $|F_{O+M}|^2$, $|F_O|^2$ and $F_{M'}$ used. Table 4 (b) presents the corresponding results obtained from X' ; the last two rows are called α_0^{ii} and α_0^{iv} . A comparison at each h , of the values of α_0^i and α_0^{ii} with those of α_0^{ii} and α_0^{iv} reveals no consistent correspondence. It is possible, however, that these values of α_0 are referred to different origins. (In fact, this is known to be true, from the way the data were obtained in this case.)

If this is so, the differences between the α_0 's obtained from the M' structure and those obtained from the X' structure should be of the form $h\beta$ (where $\beta = 2\pi x_0$ expressed in degrees and x_0 is the shift in origin between structures M' and X'). Table 4 (c) presents all these differences, the four rows being, respectively: $\alpha_0^i - \alpha_0^{ii}$, $\alpha_0^i - \alpha_0^{iv}$, $\alpha_0^{ii} - \alpha_0^{iv}$ and $\alpha_0^i - \alpha_0^{iv}$. It is seen that it is impossible to find a value of β which will make a value of the difference in each column equal to $h\beta$. It is still possible, however, that not only are M' and X' referred to different origins but also correspond to different enantiomorphs. If this is so (and we know it is), all the values of α_0 from either the M' or the X' data should be changed in sign, before taking the differences of Table 4 (c). In other words, instead of differences, sums should be taken; these sums appear in Table 4 (d), the four rows corresponding, respectively, to $\alpha_0^i + \alpha_0^{ii}$, $\alpha_0^i + \alpha_0^{iv}$, $\alpha_0^{ii} + \alpha_0^{iv}$ and $\alpha_0^i + \alpha_0^{iv}$. It is seen at once that choosing $\beta = 60^\circ$ expresses all the values in the third row in the form $h\beta$. (These numbers need not all have been in the third row, but might have varied from one row to another; that they are in one row is fortuitous.) This observation proves that structures M' and X' correspond to enantiomorphs of structure O , and that the origins were taken $60/360$ apart, i.e. $x_0 = 2/12$.

In order to refer structures M' and X' to the same origin and have them correspond to the same enantio-

morph of O , one of them must be changed to its mirror image and the origin of one of them moved by two-twelfths of the period. It is assumed that this treatment is to be given to X' . X' has atoms at $x = 0, 3/12, 10/12$; changing these to their negatives gives X'' , the enantiomorph of X' , with atoms at $x = 0, 2/12, 9/12$, X'' is still not referred to the same origin as M' ; to accomplish this, each atomic coordinate in X'' must be increased by $2/12$ to give structure X''' , with atoms at $x = 2/12, 4/12, 11/12$. The values of $F_{X'''}$, $\alpha_{X'''}$ and the corresponding two values of α_o (called α_o^b and α_o^c) are set forth in Table 4 (e).

It is seen at once that there is a perfect correspondence at each value of h between α_o'' , obtained (Table 4 (a)) from $|F_o|$, $|F_{O+M}|$ and $F_{M'}$, and α_o^b , (Table 4 (e)) obtained from $|F_o|$, $|F_{O+X}|$ and $F_{X'''}$. Thus, structures M' and X''' are referred to the same origin, and correspond to the same enantiomorph of O . The values of α_o'' (or, what is the same thing, α_o^b) are a self-consistent set of phase angles for the structure factors, F_o , of structure O . If angles α_o'' , from Table 4 (a), are compared with the original phase angles α_o , from Table 2, it is found that $\alpha_o''(h) = -\alpha_o(h) + 240^\circ h$, which means that structure O , as found 'correctly' from the data, is the enantiomorph of the true structure and is referred to an origin $8/12$ of the period from that in the original description. This, however, is all that can be expected from a structure determination using X-ray diffraction data.

6. Generalization to three dimensions

The procedure just described is easily generalized to three dimensions. In general, structures M' and X' , as found from the intensities of the X-rays diffracted by crystals with the structures O , $O+M$ and $O+X$, will be referred to different origins and are as likely as not to correspond to different enantiomorphs. The two different origins will be separated by a vector in the crystal which has components x_0 , y_0 and z_0 and the differences or sums corresponding to the entries in Tables 4 (c) or 4 (d) will be of the form $2\pi(hx_0 + ky_0 + lz_0)$ when structures M' and X' refer to the same enantiomorph of O . In consequence of this, a sequence of reflections such as $(0kl)$, $(1kl)$, $(2kl)$, ..., etc. can be used to find x_0 alone, since, for these, the entries will be of the form $h\beta + \Delta$, where β is $2\pi x_0$ and Δ is $2\pi(ky_0 + lz_0)$, both expressed in degrees. It is interesting—and advantageous—that this process can be carried through for each combination of k and l , thus providing a multitude of checks on the value of x_0 . Simultaneously, it will be discovered whether or not structures M' and X' correspond to enantiomorphs of O , and, if they do, one of them can be changed into its mirror image. In a similar way, the sequences $(h0l)$, $(h1l)$, $(h2l)$, ..., etc., and $(hk0)$, $(hk1)$, $(hk2)$, ..., etc., can be used to find y_0 and z_0 , respectively. Of course, the phases of the various F_o 's are found simultaneously. Once M' and X' are properly located and referred to

the same enantiomorph of O , all the phases of the F 's can be found. With this knowledge, it is then possible to compute directly the Fourier series for the electron density of structure O .

7. A more general treatment

The method for phase determination described in the preceding paragraphs has been based on the isomorphous replacement of vacancies in structure O by the atoms of structures M or X . This type of situation is rare in nature: in the more usual case, one kind of atom replaces another. Thus, one would usually be faced with a set of crystals with structures such as this: $O+M+X$, $O+M'+X$, $O+M+X'$ and, perhaps, $O+M'+X'$, where structures M and M' are the same, except that they are composed of different atoms, and similarly for X and X' . Consider the pair of structures $O+M+X$ and $O+M+X'$. If we suppose, as is usually true in practice, that all the atoms in X are of the same kind (for instance, chlorine atoms) and that the same holds for X' , but with a different kind of atoms (for instance, bromine atoms), then we have that

$$F_{X'} = \{f_{X'}/f_X\}F_X \quad (8)$$

and

$$F_{X'} - F_X = \{(f_{X'} - f_X)/f_X\}F_X \quad (9)$$

for each (hkl) . Thus we can write

$$F_{O+M+X'} = F_{O+M+X} + F_{X'-X}. \quad (10)$$

In most cases, the values of $F_{X'-X}$ can be found directly by comparing the Patterson functions of structures $O+M+X$ and $O+M+X'$, and, from these, working out the structure $X'-X$. This structure can be considered as substituted for a set of vacancies in structure $O+M+X$. Now, by using the experimentally determined values of $|F_{O+M+X}|^2$, $|F_{O+M+X'}|^2$ and $F_{X'-X}$, two solutions can be found for F_{O+M+X} . (This is done in the same way that two solutions are found for F_o from the measured values of $|F_o|^2$, $|F_{O+X}|^2$ and F_X , as described previously.)

By using a parallel procedure, starting with the data obtained from structures $O+M+X$ and $O+M'+X$, two independent solutions can be found for F_{O+M+X} , from $|F_{O+M'+X}|^2$, $|F_{O+M+X}|^2$ and $F_{M'-M}$. If a common solution occurs in each pair, $F_{M'-M}$ and $F_{X'-X}$ are referred to the same origin and correspond to the same enantiomorph, if not, then a 'correct' origin and enantiomorph can be found by the methods outlined in previous paragraphs, and the values of $F_{M'-M}$, $F_{X'-X}$ and F_{O+M+X} can all be referred to this origin and enantiomorph. Now, using

$$F_X = \{f_X/f_{X'-X}\}F_{X'-X} \quad \text{and} \\ F_M = \{f_M/f_{M'-M}\}F_{M'-M}, \quad (11)$$

we can find F_o from the relation

$$F_o = F_{O+M+X} - F_{M'-M} - F_X, \quad (12)$$

and, from this, structure O can be found by summing a Fourier series.

8. The effect of variations from exact isomorphism

It has been assumed in the foregoing discussion that the lattice constants and atomic parameters of the crystals in an isomorphous series are all exactly alike; this state of affairs is not likely to be found in nature, although it is sometimes closely approximated. The question then arises: how far can two crystals differ from isomorphism without seriously vitiating the method of determining phases just described?

In order to make this question definite, let us assume that structure O is a complicated organic molecule which maintains its dimensions and orientation with respect to the crystal axes, and that the effect of adding structures M or X is only to increase the lengths of the crystal axes. For convenience, we choose the origin at the centroid of scattering of structure O and assume that the greatest possible value of $|x|$, $|y|$ or $|z|$ for an atom of O is 0.5. Let us suppose that there are two equal atoms in O , with x coordinates $+0.5$ and -0.5 , respectively, when the lattice constants a_0 , b_0 and c_0 of the crystal are at a minimum. As the lattice constants increase, by fractional amounts α , β and γ , respectively, these x coordinates decrease by the fractional amount α (to a sufficient approximation). The contribution of these two atoms to the structure factors of the $h00$ reflections of the crystal is given by

$$2f \cos 2\pi h \cdot \frac{1}{2}(1-\alpha) = 2f(-1)^h \cos \pi h \alpha.$$

This contribution never changes by more than 15% of its maximum range as $h\alpha$ changes by 0.1. All other pairs of atoms in structure O contribute smaller changes to these structure factors. It seems reasonable to say that crystals will behave as isomorphous for purposes of phase determination if the change from one member of the series to another in the contribution of any atom to $|F_O|$ is less than 15% of its maximum contribution; consequently, $h\alpha$ (also $k\beta$ and $l\gamma$) should be held below 0.1. For instance, if the largest value of h under consideration is 25, then α must be less than 0.004 (0.4%), etc.

There are other kinds of deviations from isomorphism, besides those due to changes in axial lengths; for instance, structure O may be rotated or distorted by the addition of structures M and X , without changing appreciably the lattice parameters of the crystal. Such deviations from isomorphism are almost impossible to detect *a priori*, but their importance can be minimized by the use of chemical sense. In any event, the effects of such deviations will be too small to spoil this phase-determination method if the change in $hx+ky+lz$ for any atom in O and for any index triple hkl is less than 0.05 between different crystals

in the series. This criterion reduces to that of the last paragraph under the special assumptions stated there.

9. Absolute intensities

The whole of the argument presented here depends on having available the correct absolute intensities of X-ray reflection for all members of the isomorphous series of crystals under study. It is fairly easy to put all members on the same *relative* basis by dividing the measured intensities by the volumes of the crystals, since the unit cell volume does not change* throughout an isomorphous series. The factor converting to the *absolute* scale can then be determined by comparing the calculated and observed values of F_M , F_X , F_{M-M} or F_{X-X} , after the corresponding structures have been found. Wilson's method (Wilson, 1942) for placing on an absolute scale the measured intensities of the X-rays diffracted by a crystal frequently gives results in error by as much as 50%; it is, consequently, more useful as an approximate check on the absolute scale, than as a method of determining that scale exactly. On the other hand, Wilson's method should work very well for placing the various members of an isomorphous series on the same *relative* scale, since the general structure of the crystals is here always the same, and the errors in this method are due to peculiarities of the structure.

10. Application to protein crystals

It has been found recently (King, Magdoff, Adelman & Harker, to be published) that crystals of several proteins react with solutions containing large organic dye molecules in such a way that these attach themselves to the protein molecules in definite positions, but without distorting appreciably the unit cells of the protein crystals. These dye molecules can be made to contain various arrangements of heavy atoms and so can, by themselves, furnish both structures M and X . This is a particularly convenient way of forming these structures, since it can be known *a priori*, on stereo-chemical grounds, how M and X are related to one another in space and also, if they are asymmetrical, how corresponding enantiomorphs of each should be chosen.

It is probable that other ways can be found of relating structures M and X , on the basis of *a priori* knowledge, which would be applicable to the solution of the structures of other non-centrosymmetrical crystal structures.

It is a pleasure to thank all of the author's colleagues in the Protein Structure Project for their comments, criticisms and suggestions, all of which have served to improve this work. It is also a pleasing duty to thank

* If the lattice constants change by such amounts as are allowed by the considerations of the previous paragraph, the change in cell volume is still completely unimportant.

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References

- BOKHOVEN, C., SCHOONE, J. C. & BIJVOET, J. M. (1951). *Acta Cryst.* **4**, 275.
 LIPSON, H. & COCHRAN, W. (1953). *The Determination of Crystal Structures*. London: Bell.
 WILSON, A. J. C. (1942). *Nature, Lond.* **150**, 152.

Acta Cryst. (1956). **9**, 9

A New Aid to the Determination of the Point-Group Symmetry of Transparent Crystals

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The patterns formed by electrical breakdown paths provide a new way to investigate the point-group symmetry of transparent crystals because the paths lie along equivalent directions. A point-group is uniquely determined from the observation in a single-crystal slice of non-degenerate breakdown configurations. Such non-degenerate configurations are formed in most monoclinic and in all triclinic crystals, while most crystals of higher symmetry reveal degenerate configurations which are compatible with more than one point group. Degeneracy, however, is not an inherent property of the breakdown process and might be removed by suitable techniques. Since the paths are believed to be formed by electron avalanches, the symmetry shown by the breakdown pattern probably refers to the symmetry of the electrical fields within the crystal that influence the motion of electrons. The symmetry of the breakdown pattern is usually identical with the symmetry of the crystal determined by other methods. A new test for the lack of centrosymmetry in transparent crystals is described.

Introduction

This paper introduces the study of the patterns formed by electrical breakdown paths as a new aid to the determination of the point-group symmetry of transparent crystals. Each crystal shows an overall breakdown pattern that conforms with one or more point-group symmetries. Individual breakdown patterns, consisting of paths that lie along equivalent directions of the crystal, are called here breakdown configurations. Each configuration conforms with the symmetry of the overall pattern, but may differ from other configurations in orientation and in number of paths. Since the paths are believed to be formed by electron avalanches, the symmetry revealed by the pattern is probably the symmetry of the electrical fields within the crystal that influence the motion of electrons. Although, for crystals of high symmetry, the method suffers from the same limitations that are encountered in morphological studies, in that 'special forms' (here called degenerate configurations*) tend to develop, it is

sometimes possible to obtain less degenerate configurations, or non-degenerate configurations, by changing the conditions of breakdown, such as by changing the temperature of the crystal or by applying overvoltages. While the degeneracy is seldom completely removed, the partial information that is obtained is usually sufficient to show that certain point groups are incompatible with the pattern. The incompatible groups may have higher or lower symmetries than the true symmetry of the crystal for the breakdown process, depending upon the nature of the degeneracy. If one then gains knowledge by some other means of the existence of those symmetry elements which are not established by the degenerate configuration, or, if one obtains a non-degenerate breakdown configuration, a point-group symmetry is uniquely determined. Since the breakdown pattern lies in three dimensions, its full symmetry is revealed in a single-crystal section or slice. Thus euhedral crystals are not necessary for the symmetry determination.

The point-group symmetry is most simply determined by examining the breakdown configurations in

* A degenerate electrical breakdown path configuration is one that consists of paths that lie in planes or along axes that could be symmetry operators. When the paths are in such positions it is not possible to determine whether or not the planes or axes are symmetry operators for the breakdown process. Hence degenerate configurations are compatible with

two or more point-group symmetries. When all configurations are degenerate the overall pattern is degenerate and the point-group symmetry of the crystal for the breakdown process is not established.