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A Thesis

Entitled

**The Structure of Yohimbine
and other Organic Molecules
by X-Ray Crystal Analysis**

Submitted for the Degree of

Doctor of Philosophy

in the

University of Glasgow

by

Kathleen Ann Kerr

May 1966.

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SUMMARY

This thesis describes work carried out under the supervision of Professor J.M. Robertson since October 1962. It is divided into two sections.

Section I describes two structures solved by the heavy-atom method. In both cases, it is the actual structure of the molecule in the chemical sense and the stereochemistry that is of interest, and for this reason refinement was terminated at a fairly early stage.

Part I of this section reviews certain aspects of heavy-atom and Patterson methods.

Part II is concerned with the structure of the methiodide of the 7-acetoxy-7-*H*-yohimbine. The configuration of the acetoxy group is the point of major interest here. The results of this analysis have been used by N. Finch and W.I. Taylor of the G.I.B.A. Pharmaceutical Company to establish the stereochemistries of a number of related compounds. This work is also an independent proof of the structure of yohimbine.

Part III is an investigation of the stereochemistries of the monoximes of benzil through the crystal structure analysis of the *para*-bromobenzoate of the α -oxime. There is some evidence in the literature that suggests that Moscovice's original assignment should be reversed.

but the present work confirms his assignment.

Section III describes an accurate determination of bond lengths in 1,1'-binaphthyl and in perylene. Both molecules can be described as composed of two naphthalenic residues joined by long bond(s). The perylene molecule is planar but the binaphthyl molecule is not. Valence bond theory predicts that the aromatic character of the perylene molecule is localised in the naphthalenic residues and that the linking bonds are pure single bonds with no π -bond character. If this is so, the dimensions of the molecule should be very similar to those of 1,1'-binaphthyl where the dihedral angle between the planes of the naphthalenic residues is 68° - large enough to preclude any interaction between the two aromatic systems.

The refinements of the two structures are described, and the dimensions of the two molecules are compared with other molecules of similar bond type.

Appendix I is a short note concerning the choice of unit cell in a triclinic lattice.

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SECTION I

Two Heavy-Atom Structures

INTRODUCTION

This section of the thesis describes the structural analysis of 7-acetoxy-7H-yohimbine and the para-bromobenzoate of benzil monoxime. It is divided into three parts.

Part one discusses some aspects of the theory of heavy-atom and Patterson methods in the context of the problem to which they were applied. The structures in point illustrate different aspects of the cooperative use of the two methods. This is not intended to be a general discussion of the theory or application of either method.

Part two is concerned with the structure of the methiodide of the acetoxy-indolenine of yohimbine. This is an independent proof of the structure and stereochemistry of yohimbine and is in agreement with the previous assignment by chemical means. The feature of interest in this analysis is the stereochemistry at C(7), the junction between the indolenine moiety and the aliphatic ring system. This feature is discussed in terms of the work that has been done on oxidative transformations of indole alkaloids at CIBA in Summit, New Jersey.

Part three describes the analysis and refinement of the structure of the para-bromobenzoate of benzil monoxime.

Certain phenomena suggest that the accepted configuration of the α and β isomers should be reversed. The structure was solved by the use of a "minimum function" derived from a sharpened Patterson synthesis. The configuration of the molecule agrees with that currently accepted in the literature. Since the molecule crystallizes in the triclinic system, the choice of unit cell is not dictated by symmetry considerations. The cell chosen is the Bravais reduced cell. A short discussion of the problem of choosing the unique cell in the triclinic system is included in Appendix 1.

PART I1.1 The Phase Problem

It is well known that the ideal diffraction pattern can be calculated for a structure containing N atoms.

The structure factor expression gives the amplitude and phase of the scattered wave and has the form

$$(1) \quad \underline{F}(\underline{H}) = \sum_{j=1}^N f_j(\underline{H}) \exp(2\pi i \underline{H} \cdot \underline{r}_j) = |F(\underline{H})| \exp(i \phi(\underline{H}))$$

where $F(\underline{H})$ is the structure factor associated with the

plane of Miller index $\underline{H} = (h, k, l)$; $\underline{r}_j = (x_j, y_j, z_j)$ the fractional coordinates of the j^{th} atom; $\phi(\underline{H})$ is

the phase angle associated with $F(\underline{H})$ and $f_j(\underline{H})$ is the

scattering factor of the j^{th} atom. The intensities of the diffraction maxima in the ideal pattern are obtained from $|F(\underline{H})|^2$, the square of the scattering amplitude.

The positions of the maxima can be calculated from a knowledge of the size and geometry of the unit cell.

The problem facing the crystallographer is the reverse of the one stated above; given the diffraction pattern, calculate the distribution of the atoms in the unit cell.

If we regard the electron density function in the unit cell as the sum of N atomic densities, then

$$(2) \quad \rho(\underline{r}) = \sum_{j=1}^N \rho(\underline{r} - \underline{r}_j) = \sum_{j=1}^N \rho_j(\underline{r}_j)$$

where $\rho(\underline{x})$ is the density function; \underline{x} is the vector (x, y, z) ; and $\underline{x}_j' = (\underline{x} - \underline{x}_j)$. The function $\rho_j(\underline{x}_j')$ is the atomic density function of the j^{th} atom defined with the atomic centre $\underline{x}(j)$ as origin.

The scattering factor, $f_j(\underline{H})$, is the Fourier transform of the atomic density function of the j^{th} atom.

Let $(\underline{x} - \underline{x}_j) = \underline{x}_j'$. Then, by definition,

$$(3) \quad f_j(\underline{H}) = \int_{-\infty}^{\infty} \rho_j(\underline{x}_j') \exp(2\pi i \underline{H} \cdot \underline{x}_j') dV_{\underline{x}_j'}$$

Again, $\rho_j(\underline{x}_j')$ is the atomic density function referred to \underline{x}_j as origin. By substituting this expression into equation (1), it can be shown that the structure factor expression is the Fourier transform of the unit cell contents.

$$(4) \quad F(\underline{H}) = \int_{-\infty}^{\infty} \rho(\underline{x}) \exp(2\pi i \underline{H} \cdot \underline{x}) dV_{\underline{x}}$$

By the properties of Fourier series, $F(\underline{H})$ is exactly equal to the value of the appropriate Fourier coefficient of $\rho(\underline{x})$, so that

$$(5) \quad \rho(\underline{x}) = \frac{1}{V} \sum_{\underline{H}} F(\underline{H}) \exp(-2\pi i \underline{H} \cdot \underline{x})$$

The maxima in this function represent the positions of atoms in the unit cell. If the Fourier series could be calculated directly from the experimental data, solution of crystal structures would be trivial. However, the observed intensities yield only the structure amplitudes,

$|F(\underline{h})|$. The phases are not observed. This is the "phase problem" of crystallography.

Referring back to equation (1), the scattering factors and structure amplitudes can be regarded as known since the former are available in tables and the latter can be obtained from the experiment. This means that the structure factor expression can be formulated in terms of two types of unknowns, the atomic positions or the phases. If either is known, the other can be calculated. This suggests two possible approaches to the phase problem. "Direct" methods as developed by Sayre, Zachariasen, Karle and Hauptman and others, attempt to find the phases directly by finding relationships among the structure amplitudes. "Indirect" methods seek to solve the phase problem through the atomic positions.

1.2. METHODS INVOLVING HEAVY ATOMS

The heavy-atom method and isomorphous replacement are distinct techniques, but both methods have developed from Robertson's work on the phthalocyanines.

"Free" phthalocyanine and its nickel derivative are isomorphous; they have the same unit cell dimensions and the same distribution of light atoms in the crystal. In the metal derivative, the metal atom is situated at

the centre of the molecule. Since the phthalocyanines crystallize in space group $I2_1/a$ with two molecules in the unit cell, the molecular centres must lie on special positions. Thus the metal atom lies on a centre of symmetry and makes a positive contribution to all reflections. By comparing the structure amplitudes of "free" phthalocyanine and its nickel isomorph, Robertson (1935, 1936) was able to determine the phases of nearly all of the observed reflections from phthalocyanine. This is the earliest example of the solution of a crystal structure without recourse to chemical information.

The first direct application of the heavy-atom method was the solution of the structure of platinum phthalocyanine by Robertson and Woodward (1940). The platinum compound is not isomorphous with the two phthalocyanines described above, although it has the same space group. A Fourier synthesis using the observed structure amplitudes and the phase of the heavy atom alone revealed the positions of all of the light atoms other than hydrogen.

If the scattering power of one atom is very much larger than that of the other atoms in the structure and if the position of the heavy atom is known, the structure factor expression can be factored into light atom and heavy atom components;

$$(6) \quad F(\underline{H}) = f_H(\underline{H}) \exp(2\pi i \underline{H} \cdot \underline{R}_H) + \sum_{j=1}^{N-1} f_j(\underline{H}) \exp(2\pi i \underline{H} \cdot \underline{R}_j).$$

With a suitable choice of heavy atom, it will so dominate the scattering that for most planes the phase angle calculated from the first term in the series is a suitable approximation to the true phase. If a Fourier synthesis is calculated with the observed structure amplitudes as coefficients and with phases calculated from the heavy atom position, it will reveal more atomic positions. These can be used in subsequent phasing calculations to give a better approximation to the true phases. The process is continued until the positions of all atoms are known.

Lipson and Cochran (1957) suggest that the method is most effective when the scattering power of the heavy atom is about equal to that of the light atoms in the structure. If a very heavy atom is used, the contribution of the light atoms to the total scattering is so small that experimental errors and diffraction effects become important. Under these conditions it is difficult to obtain accurate parameters for the light atoms. If the heavy atom is too light, the phase of the heavy atom is not a good approximation to the true phase and the Fourier map may reveal no new atomic positions. Sim (1957, 1959) has calculated

the proportion of structure factors correctly phased by a heavy atom or a group of heavy atoms in space group $P\bar{1}$, and has developed formulae for the distribution of phase angles around the phase of the heavy atom in space group $P\bar{1}$. His tables provide a good guide in the selection of a derivative suitable for X-Ray analysis.

1.3. THE PATTERSON FUNCTION

In 1935 Patterson defined the function

$$(7) \quad P(\underline{x}') = V_0 \int \rho(\underline{x}) \rho(\underline{x} + \underline{x}') \, dV_{\underline{x}}.$$

Substituting equation 5 in equation 7 we get

$$(8) \quad P(\underline{x}') = \frac{1}{V_0} \int_{-\infty}^{+\infty} \sum_{\underline{H}} \sum_{\underline{H}'} F(\underline{H}) \exp(-2\pi i \underline{H} \cdot \underline{x}) \times F(\underline{H}') \exp(-2\pi i \underline{H}' \cdot \underline{x}) \\ \times \exp(-2\pi i \underline{H}' \cdot \underline{x}') \, dV_{\underline{x}}.$$

This function is zero except when $\underline{H} = -\underline{H}'$; when this condition applies

$$(9) \quad P(\underline{x}') = \frac{1}{V_0} \sum_{\underline{H}} F(\underline{H}) F(-\underline{H}) \exp(-2\pi i \underline{H}' \cdot \underline{x}') \\ = \frac{1}{V_0} \sum_{\underline{H}} |F(\underline{H})|^2 \exp(2\pi i \underline{H}' \cdot \underline{x}').$$

Thus a Patterson function is a Fourier series with the square of the structure amplitudes as coefficients. As defined in equation 7, it is the self-convolution of the electron density. The maxima of this function represent interatomic vectors. If there are N atoms in the unit cell, there will be N^2 peaks in the Patterson

map with N of these superimposed at the origin and one half of the remaining $N(N-1)$ related to the other half by a centre of symmetry at the origin. The weight of the j^{th} atom is its atomic number Z_j , and the weight of the ij^{th} Patterson peak is $Z_i Z_j$. Both the atomic functions $\rho(x_j)$ and the interatomic function $P(x'_{ij})$, are Gaussian, but the half-width of a Patterson peak is the root mean square of the half-widths of the atomic functions of the corresponding atoms. This leads to chance overlap of vector peaks and tends to complicate the interpretation of the Patterson function. However, inversion centers and translations in the crystal give rise to anti-parallel and parallel vectors respectively. This results in exact superposition of certain maxima, thus increasing their weights relative to background, and is often useful in interpreting a vector map. If there are a small number of heavy atoms in the structure, the outstanding vectors in the Patterson map will be the heavy atom-heavy atom vectors and the heavy atom-light atom vectors. Peaks representing interactions between light atoms form a fluctuating background of vector density for a structure of moderate complexity. If an atom is heavy enough to be distinguished in the Patterson map, its position in the unit cell can be determined from a knowledge of

Table 1

Patterson Vectors for the Space Group $F2_12_12_1$

General Positions

- 1) x, y, z
- 2) $\frac{1}{2}-x, -y, \frac{1}{2}+z$
- 3) $\frac{1}{2}+x, \frac{1}{2}-y, -z$
- 4) $-x, \frac{1}{2}+y, \frac{1}{2}-z$

Vector Positions

- | | |
|---|--|
| 2-1) $\frac{1}{2}-2x, -2y, \frac{1}{2}$ | (1-2) $\frac{1}{2}+2x, 2y, \frac{1}{2}$ |
| 3-1) $\frac{1}{2}, \frac{1}{2}-2y, -2z$ | (1-3) $\frac{1}{2}, \frac{1}{2}+2y, 2z$ |
| 4-1) $-2x, \frac{1}{2}, \frac{1}{2}-2z$ | (1-4) $2x, \frac{1}{2}, \frac{1}{2}+2z$ |
| 4-2) $\frac{1}{2}, \frac{1}{2}+2y, -2z$ | (2-3) $-2x, \frac{1}{2}, \frac{1}{2}+2z$ |
| 4-3) $\frac{1}{2}-2x, 2y, \frac{1}{2}$ | (2-4) $\frac{1}{2}, \frac{1}{2}-2y, 2z$ |
| 3-2) $2x, \frac{1}{2}, \frac{1}{2}-2z$ | (3-4) $\frac{1}{2}+2x, -2y, \frac{1}{2}$ |

Projection on (010)

position	weight
$\frac{1}{2}-2x, \frac{1}{2}$	2
$\frac{1}{2}, -2z$	2
$-2x, \frac{1}{2}-2z$	1
$2x, \frac{1}{2}-2z$	1
$\frac{1}{2}+2x, \frac{1}{2}$	2
$\frac{1}{2}, 2z$	2
$2x, \frac{1}{2}+2z$	1
$-2x, \frac{1}{2}+2z$	1

the equivalent positions of the space group. For instance, the methiodide of 7-acetoxy-7-H-yohimbine crystallizes in the orthorhombic system with space group $P2_12_12_1$. There are four equivalent positions in the space group and four molecules in the unit cell. This should give 16 heavy atom heavy atom interactions with four superimposed at the origin and the other 12 distributed through the cell (Table 1). The symmetry elements present cause eight of these to coincide in pairs when viewed in projection, thus giving four double-weight and four single-weight peaks. In the unique area of the projection on (010) there are two double- and one single-weight peak:

$$\frac{1}{2}, 2z; \quad \frac{1}{2}-2x, \frac{1}{2}; \quad 2x, \frac{1}{2}-2z;$$

where x , and z are the fractional coordinates of the atom in the unit cell.

1.4. DECONVOLUTION OF THE PATTERSON FUNCTION

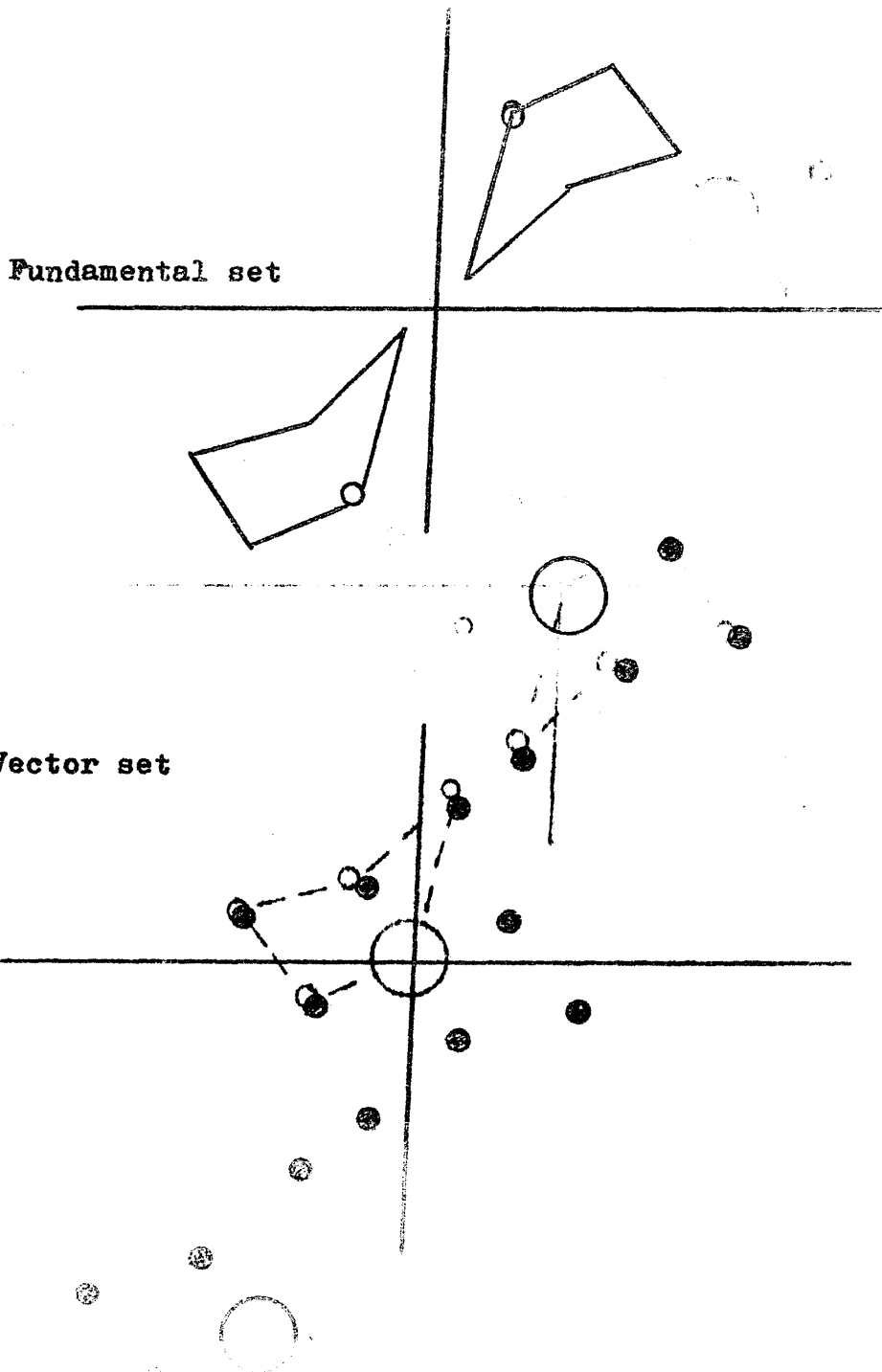
Several methods are described in the literature for deconvoluting the Patterson function to derive the crystal structure. These are reviewed briefly by Lipson and Cochran (1957) and by McLachlan (1957), and very comprehensively by Buerger (1959). These techniques are most easily understood as attempts to recover the fundamental set from the vector set by superimposing one vector map on the other so that the origin of the

first map lies on a peak in the second. The coincidences are marked and this gives a "reduced vector set". This is illustrated for a fundamental set of points and a vector set of points in figure 1.

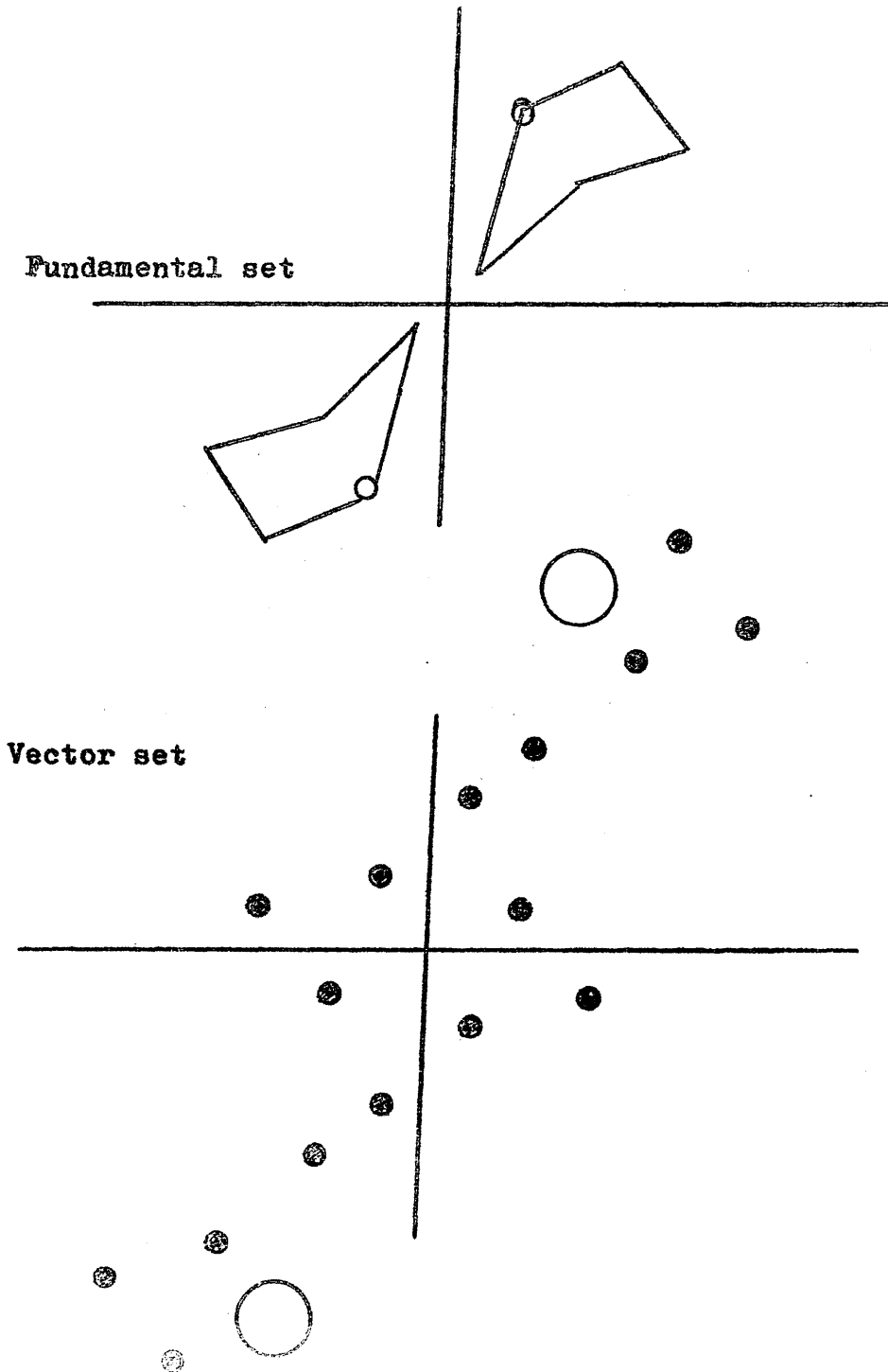
If a single weight peak is chosen as the origin for superposition, and if the fundamental set contains a centre of symmetry, that set can be recovered by a single superposition. If the fundamental set is not centrosymmetric, the "reduced vector set" contains the fundamental set and its mirror image. The penalty incurred for placing the origin of one vector map on an "n"-fold multiple peak in the second is to recover the fundamental set and its congruent image repeated "n" times in different positions. However, successive superpositions will reveal a single solution if sufficient single weight peaks are found.

This need for single weight peaks limits the applicability of the method. A Patterson peak representing a vector between like atoms will have a half-width $\sqrt{2}$ greater than that of the corresponding atomic peak. If there are N resolved peaks in the Fourier synthesis, the N^2 Patterson peaks are unlikely to be resolved in the same volume. In a structure of reasonable complexity with atoms of similar scattering power, the single weight peaks will be among the smallest

"Fundamental set" containing a heavy atom, and
the heavy atom interactions of the "vector set".



"Fundamental set" containing a heavy atom, and
the heavy atom interactions of the "vector set".



in the map and may be lost in a background of vector density. This lack of resolution causes a further difficulty when we attempt to recover the density function by superposition methods. If we record only the sites where the two functions overlap when superposed, chance coincidences will produce false atomic positions.

McLachlan suggests that the resolution of a superposition function can be improved if the vector density is summed or multiplied at each point in the cell. This method does enhance the correct peaks but it does not eliminate "ghost" peaks caused by chance coincidences. Buerger claims that the best resolution is obtained when the minimum value is recorded at each point. The resulting minimum function has a value only when both functions have a positive value and is zero elsewhere.

One of the simplest applications of the method is in the solution of structures with space group $\bar{P}1$. The parabromobenzoate of benzil monoxime will serve as an example. In a centrosymmetric space group, vectors between atoms not related by the inversion centre are constrained to overlap in pairs to give peaks of weight $2Z_i Z_j$ at $x_i - x_j$. Atoms related by the centre of symmetry give rise to vectors of weight Z_j^2 at $2x_j$. Thus the heavy atom-heavy atom interaction produces a single weight peak. Once this has been identified, the electron

density can be recovered in a single superposition. These conditions apply in the case of the benzil monoxime derivative. Two copies were made of a sharpened, three-dimensional Patterson map. The origin of one map was placed on the centre of the Br-Br vector peak of the second, and the various sections were paired to maintain this displacement. A "minimum function" was derived by tracing minimum contours where peaks in the two maps overlapped. The origin of coordinates in the "minimum function" was defined at the mid-point of the displacement; half-way along the bromine-bromine vector.

1.5. SUMMARY

The Patterson function is a very powerful tool in the early stages of structure analysis. It can be used with both centered and non-centered structures, for molecules with atoms of similar scattering power or for molecules with one or more heavy atoms. When combined with the heavy-atom method, it provides a very elegant means of circumventing the phase problem.

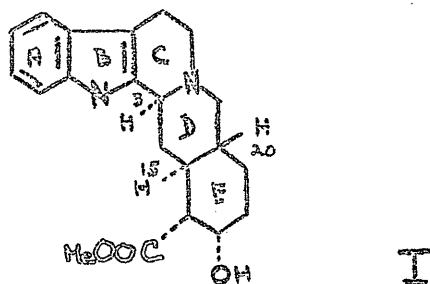
Part II

The Structure of Yohimbine:

The Crystal Structure of the
Methiodide of 7-Acetoxy-7-H-Yohimbine

2.1. INTRODUCTION

Yohimbine is an indole alkaloid isolated from yohimbe bark and used as an aphrodisiac in veterinary medicine. The structure of yohimbine, I, was first established by chemical means. The final link in the long chain of evidence leading to the complete structure was provided by Swan (1950), when he determined the position of the hydroxyl group. The work of Witkop (1949) and others established the stereochemistry of the molecule. This work is well documented in "The Alkaloids" (Manske, vols. II and VII).



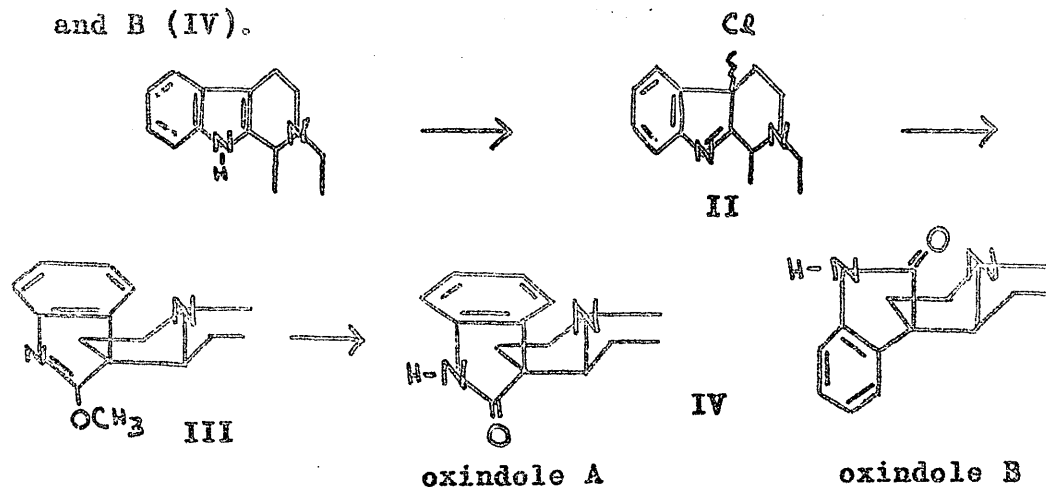
The molecule is characterized by five asymmetric centres. Hydrogen atoms at C(3), C(15) and C(20) are in axial positions; those at C(3) and C(15) are in the α configuration. Rings D and E are trans-fused.

The present structure analysis of the methiodide of 7-acetoxy-7-H-yohimbine confirms both the gross

structure and the stereochemistry of yohimbine itself. However, its importance lies in its connection with research on oxidative transformations of indole alkaloids by N. Finch and W.I. Taylor of the CIBA Pharmaceutical Company. By correlating the results of this X-ray analysis with their own optical rotatory dispersion data, Finch and Taylor have been able to draw some fairly definite conclusions about the mechanisms of certain reactions of indole alkaloids, and to establish the stereochemistry of several alkaloids of this class.

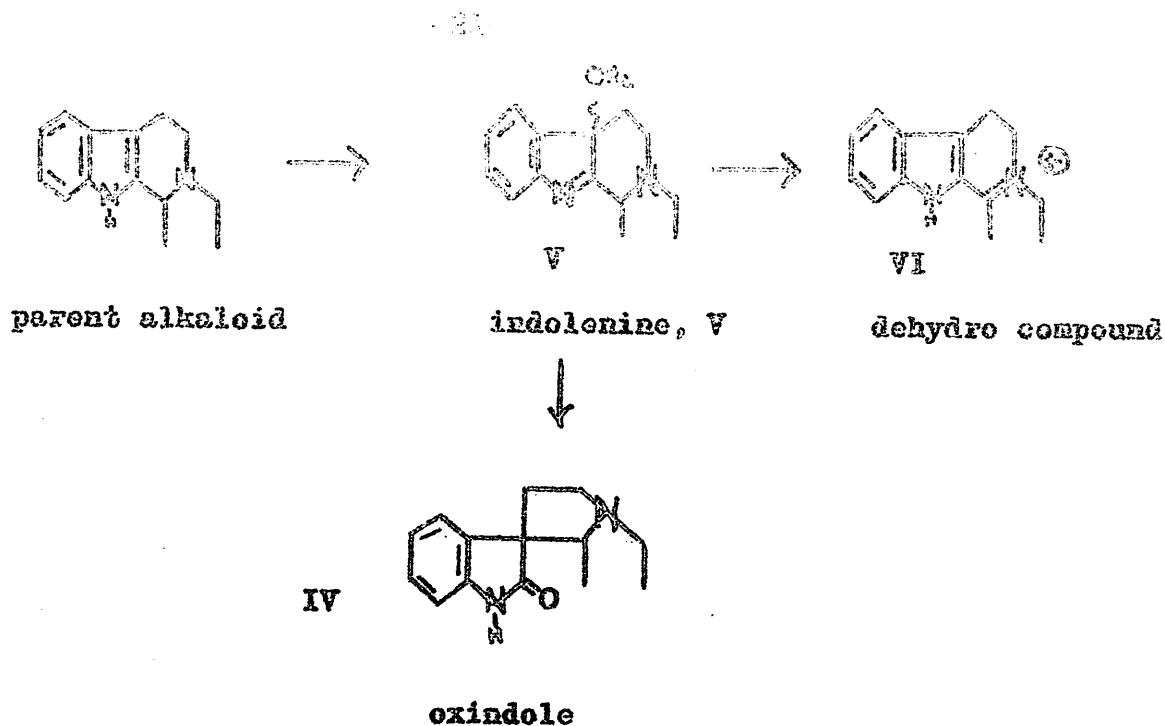
2.2. OXIDATIVE TRANSFORMATIONS OF YOHIMBINE AND RELATED ALKALOIDS

In 1962, Finch and Taylor reported that yohimbinoid alkaloids with trans-fused DE ring systems could be converted to their oxindole equivalents by methanolysis of the 7-chloro derivatives (II) to give the imino-ester (III). When the ester is refluxed with aqueous acetic acid, the reaction product is a mixture of oxindoles A and B (IV).



When this series of reactions was attempted using an alkaloid with cis fused DE rings, the results were markedly different. Instead of a 50:50 mixture of epimers of the chloro-derivative, there was a marked preponderance of the Cl-axial epimer. This resulted in a reduction from 40% to 4% in the yield of the methanolysis product, the imino-ester. Thus the method is of practical utility in synthesising only those oxindole alkaloids that have trans DE stereochemistry. (Finch and Taylor, 1962a; 1962b).

A later paper (Finch et al., 1963) explored the reactions of Pb^{IV} salts of carboxylic acids with alkaloids of the yohimbine type. In all cases, the reactions of the parent alkaloid in methylene chloride with one molar equivalent of the lead salt yielded the corresponding acyloxy-indolenine (V). When treated with acetic acid, the indolenines eliminated the acyloxy residue to give the dehydro-compound (VI). However, if the indolenines are refluxed with methanol and a few drops of acetic acid, those with trans-fused DE ring systems do not react but those with cis-fused systems rearrange to give the corresponding oxindole (IV).



Oxindole B is the major product of the rearrangement regardless of the stereochemistry at C(3). The reaction mechanism proposed depends on the protonation of the indolenine nitrogen and rearrangement of the resulting carbonium ion with neighbouring-group participation by the acetoxy moiety. This mechanism would explain why trans-fused systems do not rearrange under the conditions of the reaction. The bridge-head nitrogen is more basic than the indolenine nitrogen, therefore protonation cannot occur at the latter site. However, in acyloxy indolenines with cis DE stereochemistry there is a fairly large 1:3-diaxial interaction with the lone pair of the bridge-head nitrogen. This decreases the basicity of this nitrogen relative to the indolenine nitrogen and allows protonation of the latter site. (Winch et al., 1963).

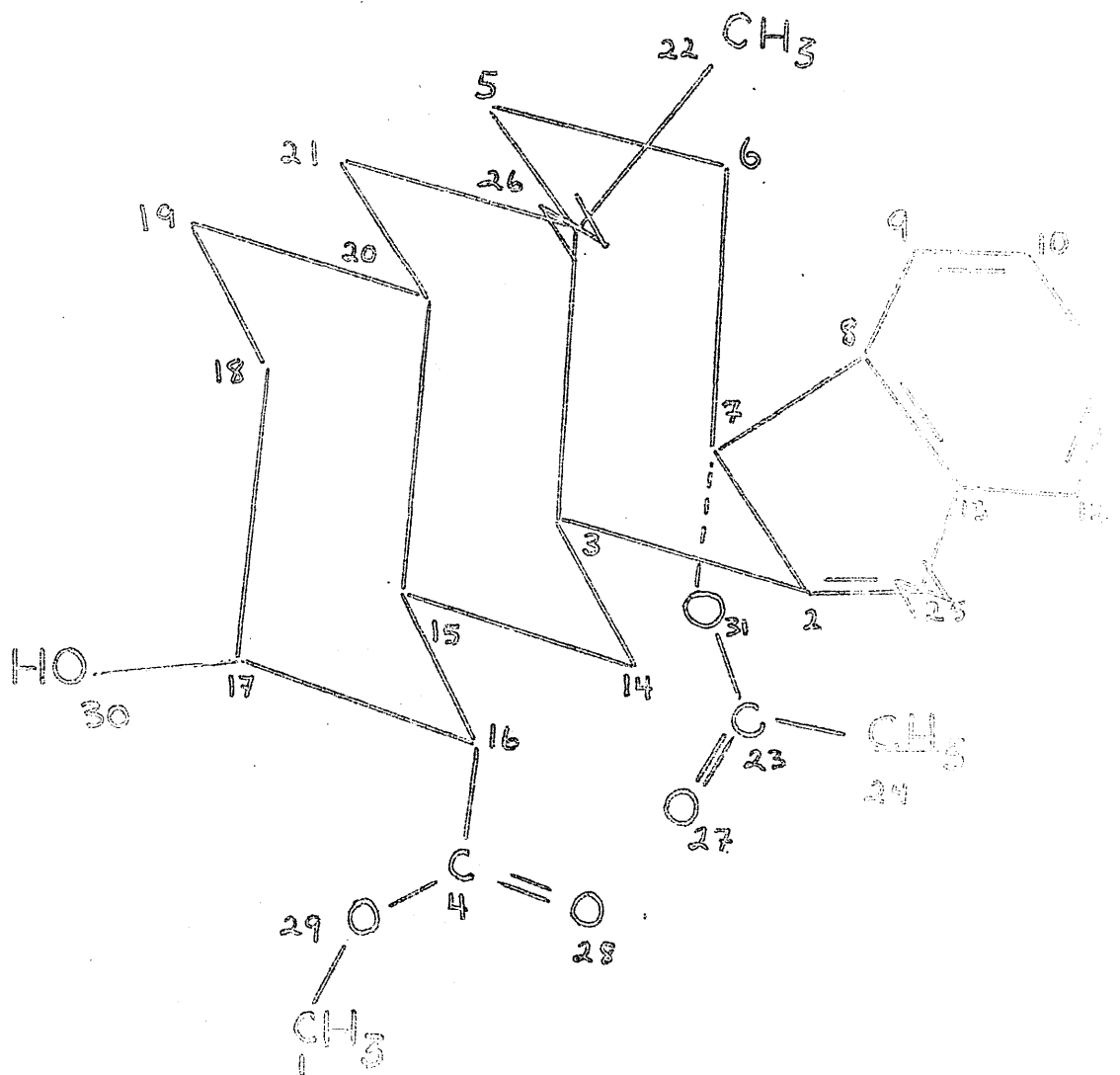
At this stage, it was uncertain whether the $\text{Pb}(\text{OAc})_3$ group is displaced with retention or inversion of configuration in the formation of the indolenine. On the basis of chemical evidence, it was not possible to determine whether the rearrangement at C(3) occurs before or after the formation of the oxindole. It was hoped that a detailed knowledge of the stereochemistry of at least one of the acyloxy indolenines would be helpful in interpreting these reactions. The compound chosen for X-ray analysis was the acetoxy-indolenine of yohimbine.

2.2.1. X-RAY ANALYSIS OF 7-ACETOXY-7-H-YOHIMBINE

The methiodide of 7-acetoxy-7-H-yohimbine was prepared by Dr. Taylor and his colleagues by dissolving 7-acetoxy-7-H-yohimbine in methyl iodide and recrystallising the resulting salt from methanol. This gave colourless, well-formed needles, suitable for X-ray crystallography, m.p. 225-227°C. The crystal structure of the methiodide was determined by the heavy atom method. Approximate locations of all light atoms, other than hydrogen, were found from a series of three-dimensional electron -

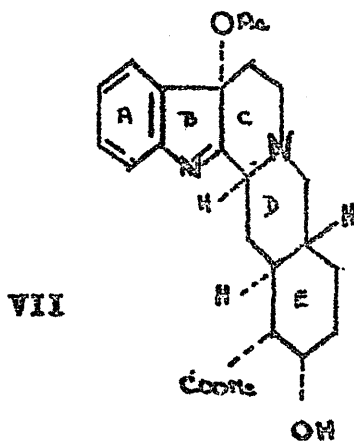
Figure 2

Stereochemistry of the Methide of 7-Acetoxy
7- β -Yohimbine.



density syntheses. Refinement of positional and thermal
of
parameters by the method of least squares gave a final
R-value of 14.60% over 1059 independent reflections.

Molecular dimensions are given in sub-section 2.4
after a description of the experimental work. The
stereochemistry of the methiodide of 7-acetoxy-7-*H*-
yohimbine is shown in Figure 2. The molecule has a
trans-fused DE ring system. The hydrogen at C(3) is
in the α position as is the acetoxy group at C(7). This
corresponds to the structural formula VII for the acetoxy
indolenine tentatively proposed by N. Finch at the
I.U.P.A.C. Symposium on the Chemistry of Natural Products
Prague, 1962.



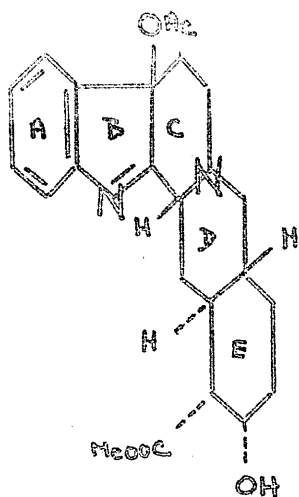
acetoxy indolenine of yohimbine

2.2.2. CORRELATION OF X-RAY RESULTS WITH O.R.D. DATA

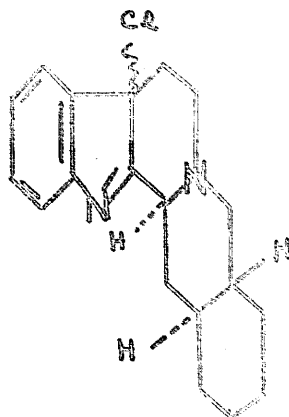
Optical rotatory dispersion curves are available for 7-acetoxy-7-H-yohimbine, VII; two epimeric 7-chloro-7-H-yohimbans, VIII; methyl-7-hydroxy-7-H-reserpate, IX; and the acetoxy indolenines of pseudoyohimbine, X, and aracine, XI. The formula of these compounds are shown in Figure 3; their O.R.D. curves are shown in Figures 4 and 5. These curves have been drawn from diagrams published by Finch et al. (1965).

The optical rotatory dispersion curves of the two chloro-yohimbans bear a mirror image relationship to one another (Figure 4). This indicates that the C(7) substituent makes the largest contribution to the Cotton effect. The two curves have nearly identical amplitude but opposite sign because the CD ring fusion constrains the C(7)-C(8) bond to be equatorial or pseudo-equatorial in the two epimers. This means that the substituent at C(7) is either axial or pseudo-axial and will have a large interaction with the chromophore group in either the α or the β configuration. Any contribution to the dispersion curve from the hydrogen at C(3) would be expected to be small.

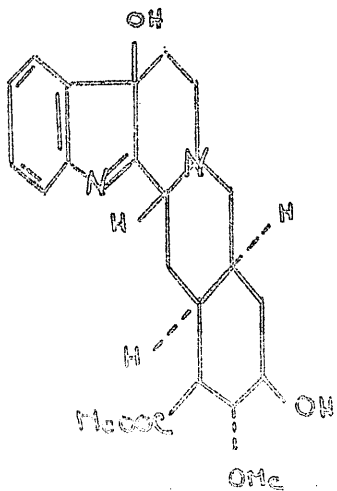
Figure 3



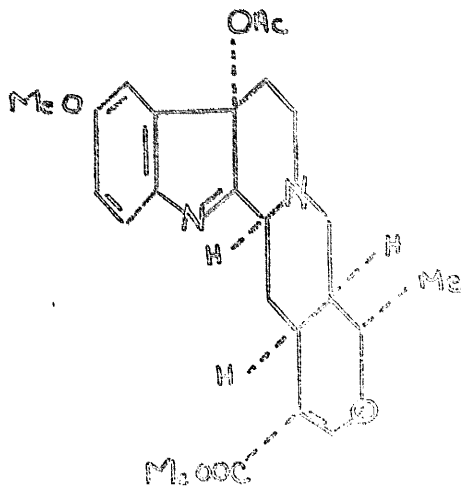
X, 7-β-acetoxy-7-H-pseudoyohimbine



VIII, 7-chloro-7-H-yohimban



IX, methyl-7-β-hydroxy-7-H-reserpate



XI, 7-α-acetoxy-7-H-ajicine

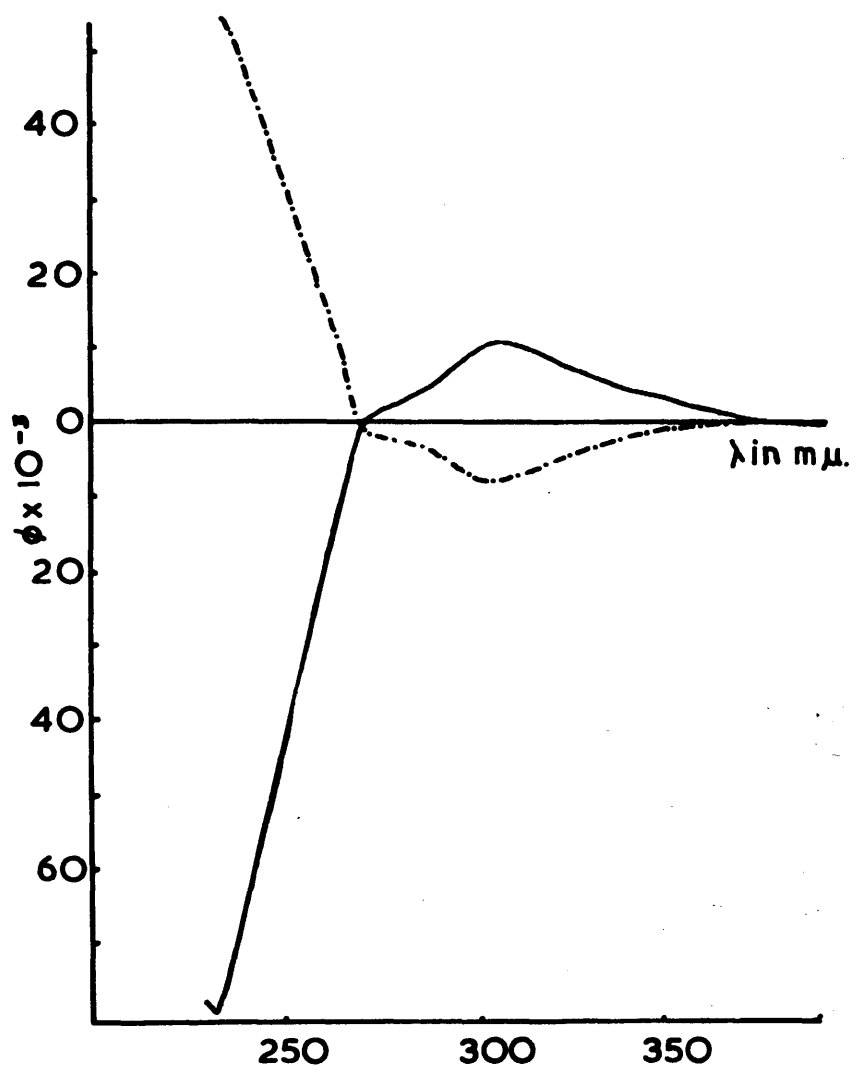
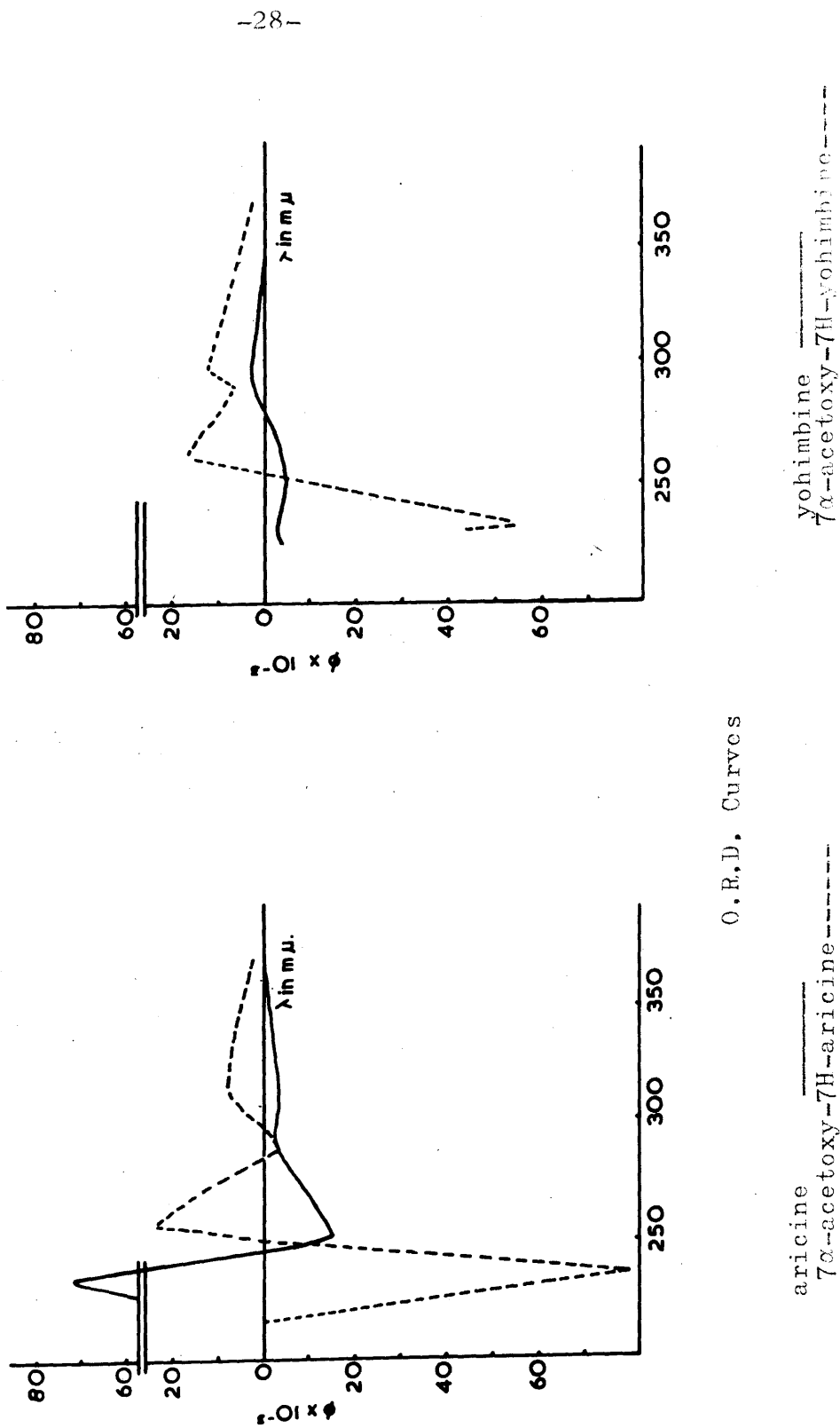


Figure 4

O.R.D. curves for epimeric 7-chloro-7-H-yohimbans

Figure 5



The dispersion curves of the yohimbine and aricine derivatives (Figure 5) have the same sign and similar shape in spite of different configurations at the DE ring junction. On the assumption that the C(7) contribution dominates the dispersion curve, the α configuration was assigned to the acetoxy group in the aricine derivative. The curves of the other two compounds have the sign expected of indolenines with a C(7) substituent in the β position. (Finch et al., 1965).

These results throw some light on the mechanism of the reactions of lead salts of carboxylic acids to give acyloxy indolenines. The configuration of the C(7) substituent can be predicted if the position of the hydrogen at C(3) is known. If that hydrogen has an α configuration, the C(7) substituent will do likewise. This observation is compatible with the reaction mechanism proposed in an earlier paper (Finch et al., 1963). Although the stereochemistry of this reaction is now established, there is still no information about the subsequent rearrangement of the indolenines to oxindolone. The fact that the observed product is the one expected to be most stable in the reaction mixture would seem to support the thesis that epimerization at C(3) occurs after rearrangement.

2.3. EXPERIMENTAL

The sample provided by Dr. Taylor consisted of small white needle-like crystals that showed sharp extinction along the needle direction. Rotation, Weissenberg and precession photographs were taken with the crystal mounted so that the needle axis was perpendicular to the X-ray beam. These photographs were used to determine cell dimensions and space group.

2.3.1. CRYSTAL DATA

Formula	$C_{24}H_{31}N_2O_5I$	F.W. 554.43 gm
System	orthorhombic	
Lattice constants	$a = 13.16 \text{ \AA}$	
	$b = 26.82 \text{ \AA}$	
	$c = 7.33 \text{ \AA}$	
Absent Spectra	$h00$ with $h = 2n + 1$	
	$0k0$ with $k = 2n + 1$	
	$00l$ with $l = 2n + 1$	
Space Group	$P2_12_12_1$	
Volume	2587 \AA^3	
Density (obs.)	1.44 gm./cc	
Molecules/cell	4	
Density(calc)	1.42 gm./c.c.	
$F(000)$	1128	
μ (CuK α)	110 cm^{-1}	

2.3.2. INTENSITY DATA

The 0kl zone was photographed by the precession method. Since the crystal-to-film distance is critical in this method (Buerger, 1944), only one film can be placed in the cassette at a time. A timed series of six photographs was taken using a stabilised power supply. Exposure times ranged from 11 minutes to 45 hours. Mo K α radiation was used. The precession angle was set at 30°. Lorentz-polarisation corrections were applied to these data by means of a template. (Buerger 1960, p 188).

Reflection data for the six reciprocal lattice nets hk0 to hk5 were collected using an equi-inclination Weissenberg camera and multiple film packs (Robertson, 1943). The intensities of these reflections were estimated visually by comparison with a calibrated intensity scale. The film factor between successive photographs of a series was 3.3 for the zero layer. This was modified for upper layers to allow for the increase in path length through the film with increasing inclination angle (Rossmann, 1956). These data were collected with CuK α radiation. They were corrected for Lorentz and polarization factors according to the formula

$$I(hk\ell) = \frac{1}{\sin 2\theta} \cdot \frac{1 + \cos^2 2\theta}{2} \cdot |F(hk\ell)|^2$$

The form of the Lorentz factor shown here applies only to zero layer photographs of rotating crystals. For upper layers, this must be modified by a factor $1/D$, where

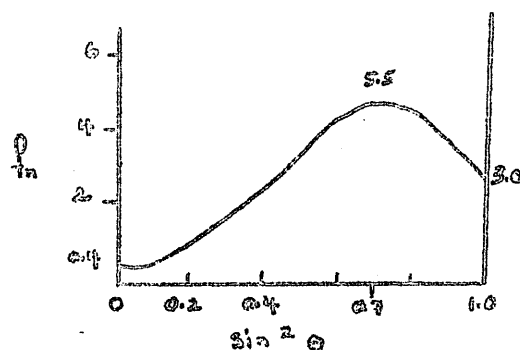
$$D = (\cos^2 \mu - \cos^2 \theta)^{\frac{1}{2}} / \sin \theta$$

the rotation factor defined by Tunnell (1939) for the equi-inclination case. The symbols μ and θ are the equi-inclination angle and the Bragg angle, respectively. No absorption corrections were applied.

The structure amplitudes, $|F_{\text{obs}}|$, were given the same relative scale by comparison of axial reflections from Weissenberg photographs with common reflections from precession photographs of the $0k\frac{1}{2}$ zone.

2.3.3. ANALYSIS OF THE STRUCTURE

Patterson coefficients, $|F|^2$, were modified by a



tabulated function of $\sin^2 \theta$ and used to compute sharpened Patterson projections, $F(VW)$ and $P(UV)$. The form of the modification function is shown in the figure on the left.

The $I-L$ vectors were easily recognized in the Patterson maps. Fractional coordinates of the iodide ion were;

$$x = 0.1772, \quad y = 0.1026, \quad z = 0.1453.$$

Structure factors were calculated with phases determined by the heavy atom alone. These were placed on absolute scale by comparison of the sums of the observed and calculated structure amplitudes.

$$R\Sigma|F_o| = \Sigma|F_c|$$

The R-value, $\Sigma\Delta/\Sigma|F_o|$, was 37%

The iodide ion is very "heavy" in comparison with the light atoms in the structure. Using Sim's (1957) method, it is possible to predict how well the phase of the heavy atom approximates to the true phase. Sim has calculated the fraction, $N(\alpha)$, of structure factors whose phase is within $\pm\alpha$ of the phase of the heavy atom for various values of r , where r is defined by

$$r = (\Sigma f_H^2 / \Sigma f_L^2)^{\frac{1}{2}}$$

In general, the probable phase-angle error decreases as r increases. For this compound, the value of r is 1.46. Interpolation on Sim's graph shows that only about 20% of the structure factors will have phase errors greater than $\pm 35^\circ$ in a calculation based on the position of the heavy atom alone. This estimate is probably a pessimistic one, as it is strictly valid only for space group $F1$. The space group $P2_12_12_1$ has three centrosymmetric projections. About 88% of the data from these zones can be expected to have the correct sign. Since nearly all

of the data were considered to be reliably phased by the heavy atom, only those planes with very small values $|F_{\text{obs}}|$ were excluded from the first Fourier calculation. The resulting map revealed the positions of all of the atoms in the skeleton of the molecule. When these were included in the next structure-factor calculation with scattering factors appropriate to carbon atoms, the R-value dropped to 29.6%. After the second Fourier synthesis all 31 light atoms, excluding hydrogen, had been located and identified. They were included in the next cycle of Fourier refinement with scattering curves corresponding to their chemical types.

After a third cycle, the coordinates were corrected for termination of series errors by a modification of Booth's (1946, 1947) method. This involves calculating two Fourier syntheses with the same coordinates. Observed structure amplitudes are used as coefficients in one map, and calculated structure amplitudes for the other. If a peak appears at a position, Σ_{p_j} , in the penultimate Fourier map, it should appear in the same position in an F_o map calculated from those coordinates. Termination of series errors cause these two sets of coordinates to differ by $\Delta \Sigma_j = \Sigma_j - \Sigma_j$ for the j^{th} atom. If x_{o_j} is the position of this atom in the final Fourier map, its correct position is given by $x_{o_j} - \Delta \Sigma_j$.

If atom peaks appear at different heights in the F_o and F_c syntheses, the temperature factor is in error by an amount that can be estimated from the difference in peak height. In the two maps calculated here, peaks in the F_o synthesis were systematically lower than those in the F_c synthesis, indicating that the temperature factor $\alpha = 2^{-B}$, had been underestimated. When α was increased from 2.8 to 3.0 ($B = 4.94 \text{ \AA}^2$), structure factors calculated with corrected coordinates showed an R-value of 21%.

2.3.4. REFINEMENT OF THE STRUCTURE

The computer program used for least-squares refinement on DEUCE was written by J.S. Rollett and is described elsewhere. (Rollett, 1961). The program uses a three by three matrix for the atomic positions and a six by six matrix for the atomic vibrations. In order to prevent overflow of the least-squares totals, it was necessary to reduce the scale of the scattering curves and of the structure amplitudes to 1/10 of the normal value. The weighting scheme used gives maximum value to the "average" structure amplitude:

$$\text{if } |F_o| < F^* / w = F_o / F^* ,$$

$$|F_o| > F^* / w = F^* / F_o ,$$

$$\text{where } F^* = 8 F_{\text{min}} = 4.64 .$$

There is no facility for isotropic refinement in this program. Anisotropic temperature parameters are defined by the equation

$$\exp(-B \sin^2 \theta / \lambda^2) = 2 \cdot (b_{11}h^2 + b_{22}k^2 + b_{33}l^2 + b_{12}hk + b_{23}kl + b_{13}hl).$$

After each cycle the b_{ii} terms from the diagonal of the vibration matrix were used to calculate the thermal parameters parallel to each of the principal axes using the following formulae:-

$$B_{11} = b_{11}/1.4427,$$

and

$$B_X = 4B_{11}/a^2.$$

(Rollett, 1961; Rossmann, Jacobson, Hirshfeld and Lipscomb, 1959; Cruickshank, 1956).

The components of B_g in the three axial directions were averaged to give an isotropic temperature factor for each atom. These individual temperature factors were averaged over all members of each chemical type. After two cycles of least squares, the R-value was 19.9%. The average B_g values for the different chemical types were

$$I = 4.79 \text{ \AA}^2 \quad C = 4.53 \text{ \AA}^2 \quad N = 3.74 \text{ \AA}^2 \quad O = 5.49 \text{ \AA}^2$$

No further calculations were done on DEUCE. The parameters obtained from the second cycle were used when refinement was continued on KDF9 eight months later.

When the Glasgow SFLS program became available, two rounds of isotropic least squares were calculated with unit weights. The R-value was 17.8%. Cruickshank's (1961) weighting scheme has been recommended for film data and seemed appropriate for this case. It has the form

$$w = 1 / (p_1 + |F| + p_2|F|^2 + p_3|F|^3).$$

The initial values given to these parameters were as follows:

$$p_1 = 17 \quad p_2 = 8.9 \times 10^{-3} \quad p_3 = 9.9 \times 10^{-5}$$

After a further three cycles of isotropic block-diagonal least squares, the R-value was 15.4% but the weighting analysis showed that some adjustment in the weighting parameters was required. One more cycle with the constants in the weighting function changed to 8, 1.04×10^{-2} , and 1.34×10^{-4} , respectively improved the weighting analysis but had little effect on the R-value. Reflection data for the short and median axes were sorted from the full, scaled, three-dimensional data that had been used for the least-squares refinement. These data were used in a "minimum residual" refinement by the method of Bhuiya and Stanley (1963). Three cycles of refinement reduced the R-value in the short-axis projection from 19.5 to 17.5%. A single refinement cycle in the median axis projection produced an improvement of about 1% to give a final R-value of 17.9%. When the new positions were used for a three-

dimensional structure-factor calculation, the R-value was 16.9% but this dropped to 14.6% after three further cycles of block-diagonal least squares. Refinement was terminated at this point. Unobserved reflections were not included in the analysis.

2.4. MOLECULAR DIMENSIONS AND DESCRIPTION OF THE STRUCTURE

Table 2 shows the final atomic coordinates and the isotropic temperature factor for light atoms other than hydrogen. The coordinates and anisotropic thermal parameters of the iodide ion are listed separately at the foot of the table. In all cases, the coordinates are expressed as fractions of the crystallographic axes. Their estimated standard deviations have been calculated from the least squares residuals and are shown in parentheses to the same number of places of decimal as the coordinates with which they are associated. Both the coordinates and their standard deviations are listed with one figure that is not significant to permit derived quantities to be calculated without loss of accuracy.

The isotropic temperature factor used here is the mean square vibration amplitude, U_{iso} , where

$$U_{iso} = \bar{u}^2 = B/8\pi^2$$

and B is the Debye factor. Both U and B have the units of \AA^2 . Anisotropic vibrational parameters are defined by the equation

$$\exp(-B \sin^2 \theta / \lambda^2) = \exp[-2\pi^2 (U_{11}h^2a^{*2} + U_{22}k^2b^{*2} + U_{33}l^2c^{*2} + 2U_{12}hka^*b^* + 2U_{23}k\ell b^*c^* + 2U_{13}h\ell a^*c^*)]$$

Bond lengths and angles with their estimated standard deviations are listed in Tables 3 and 4, respectively. Figure 6 is a schematic diagram showing bond lengths and the numbering used in the Tables. From Table 3, it can be seen that the average standard deviation of bond lengths in this structure is 0.05 to 0.06 Å. In the benzene ring, the average carbon-carbon bond length is 1.39 Å. The C(sp₃)-C(sp₃) single bonds in the CDE ring system have an average length of 1.55 Å as do the C(sp₂)-C(sp₃) bonds at the junction of that ring system with the indolenine moiety. These lengths do not differ significantly from the expected values of 1.397 Å, 1.525 Å, respectively quoted by Sutton et al. (1958).

The carbon-nitrogen bonds fall into three categories. The C=N bond is 1.24 ± 0.04 Å long, not appreciably different from the value of 1.27 Å found in dimethyl-glyoxime. The C(sp₂)-N bond length of 1.36 ± 0.05 Å is in agreement with the value of 1.371 Å reported in para-nitroaniline (Trueblood, Goldish and Donohue, 1961); 1.38 Å in 2-chloro-4-nitroaniline (McPhail and Sim, 1963); 1.39 and 1.40 Å in ibogain hydrobromide (Arai, Coppola and Jeffrey, 1960). The value of 1.54 Å for the average C(sp₃)-N bond length is somewhat longer than the value given by Sutton et al., but the long

bonds of this type are frequently reported in alkaloids. In the structure of schittamine (Hamilton, Hamor, Robertson and Sim, 1962) there are three bonds of this type with a length of 1.54 Å and a fourth with a length of 1.56 Å. After comparing the results of several analyses of different alkaloids, these authors suggest that 1.52 Å is perhaps a better estimate of the length of such a bond.

Positional standard deviations and thermal parameters are greater for atoms in the side chains of this molecule. The bond lengths show broad agreement with the expected values. The carbon-oxygen single bond lengths can be placed in two groups; those adjacent to a carbonyl group are shorter than those removed from the carbonyl. Average lengths are 1.33 and 1.44 Å respectively.

The equation of the weighted mean plane through the benzene ring is;

$$0.0195X + 0.1207Y - 0.9925Z = 0.1275$$

Each atom was given a weight inversely proportional to the estimated variance of the atomic position. Deviations from this "best" plane are shown diagrammatically in Figure 7. The units are Å x 10³. The only atom in the indoleine moiety that deviates significantly from the plane through the benzene ring is C(7). This is not a surprising result since resonance interaction between the benzene ring and the double bond would tend to keep the C-N group in the plane.

of the benzene ring. C(7) would be expected to deviate from this plane because it is fully substituted and cannot take part in resonance interactions.

Intermolecular contacts under 3.6 \AA are listed in Table 5 and are shown in the packing diagram (Figure 8). For clarity, these have been divided into two groups and are shown as contacts from molecules I and I'; i.e. the molecule whose coordinates are given in Table 2, and the same molecule in the next cell in the positive x -direction. The only contact of particular interest is that between N(25) and O(30). This is a hydrogen bond of the O-H...O type between the hydroxyl group at C(17) and the indole ring nitrogen of a symmetry-related molecule. The O...N distance ($2.90 \pm 0.03 \text{ \AA}$) is more reliable than many of the bond distances in the structure. The C-O...N angle is $103.8 \pm 1.7^\circ$.

Table 6 and Figure 9 show the environment of the iodide ion. The ion has more than a dozen contacts closer than the quaternary nitrogen with which it is formally associated. The $N^+ \dots I^-$ contacts are quite long and some of the contacts between the iodide ion and the atoms of the acetoxy group seem quite short.

The iodide ion is almost midway between the quaternary nitrogen of one molecule and that of the same molecule in the next cell in the negative x -direction. The $N^+ \dots I^-$

distances are 4.59 and 4.63 Å respectively. However, the iodide ion does not lie on a direct line between the two nitrogens but is displaced towards the acetoxy group. Its closest contacts are with C(23), C(24) and O(31) of that group. Expected van der Waals separations are shown in the table. These were calculated in the following way: Pauling's figures for the ionic radius of iodide (2.16 Å), oxygen (1.40 Å), and a methyl group (2.00 Å), predict a minimum separation of 4.15 Å between iodide and C(24), and 3.66 Å between the ion and O(31). The van der Waals radius of a carbon with no hydrogen would probably be about half the inter-layer spacing in graphite, or 1.70 Å. Thus C(23) would not be expected to be closer than about 3.86 Å. Several of the observed contacts are very near the expected van der Waals distances. The contact with C(24) is very short. In view of the high standard deviation in the experiment and the uncertainties in the van der Waals distances, very little significance can be placed on any individual observation.

Both the long $N^+ \dots I^-$ distances and the short contacts between iodide and carbon and iodide and oxygen seem to be observed frequently in compounds of this type. Silvers and Yuliansky (1963) report $N^+ \dots I^-$ distances of 4.6 and 5.1 Å in skatamine; and in cocaineine-II dimethide, McPhail (1963) reports four distances between 4.42 and 4.56 Å.

Short contacts of iodide with carbon and oxygen have been observed by Przybylska (1961 a, 1961 b), Jellinek (1957), McPhail, Robertson and Sim, (1963) and others. This is probably a steric effect. It is possible that with an ion the size of an iodide ion, packing consideration and van der Waals forces play a greater role in determining the environment of the ion than do forces of Coulombic attraction.

TABLE 2

Mathematical and Temperature Parameters

Final atomic coordinates and temperature factors.
Standard deviations are given in units in the last place.

Atom	x	y	z	U ₁₁			
C (1)	0.0005	(28)	0.1595	(16)	-0.3878	(73)	0.078
C (2)	0.0182	(21)	0.3257	(12)	0.4502	(36)	0.045
C (3)	-0.0574	(19)	0.3259	(11)	0.4525	(58)	0.057
C (4)	-0.0541	(16)	0.2134	(10)	-0.4596	(38)	0.031
C (5)	-0.1504	(24)	0.4131	(13)	0.3526	(65)	0.052
C (6)	-0.0745	(25)	0.4302	(20)	0.3740	(31)	0.030
C (7)	-0.0162	(19)	0.4203	(16)	0.2022	(57)	0.023
C (8)	0.0240	(28)	0.4765	(15)	0.1955	(70)	0.021
C (9)	0.1249	(25)	0.5230	(12)	0.2170	(62)	0.021
C (10)	0.2141	(27)	0.5232	(21)	0.2294	(31)	0.021
C (11)	0.2875	(25)	0.5081	(20)	0.2270	(58)	0.021
C (12)	0.2612	(22)	0.4459	(11)	0.1832	(59)	0.021
C (13)	0.1601	(23)	0.4351	(15)	0.1899	(66)	0.020
C (14)	-0.0323	(21)	0.3045	(11)	0.0923	(51)	0.021
C (15)	-0.1251	(18)	0.2706	(09)	0.1021	(51)	0.021
C (16)	-0.1010	(29)	0.2138	(16)	0.0333	(70)	0.020
C (17)	-0.1995	(20)	0.1822	(11)	0.0341	(56)	0.021
C (18)	-0.2542	(22)	0.1847	(12)	0.2505	(70)	0.021
C (19)	-0.2207	(20)	0.2339	(12)	0.2362	(73)	0.020
C (20)	-0.1770	(23)	0.2701	(11)	0.2620	(57)	0.020
C (21)	-0.2101	(22)	0.3270	(12)	0.2312	(60)	0.020
C (22)	-0.0220	(25)	0.3779	(21)	0.2103	(78)	0.020
C (23)	-0.0731	(25)	0.4725	(14)	-0.1107	(50)	0.021
C (24)	-0.1223	(31)	0.5031	(17)	-0.1937	(41)	0.022
C (25)	0.1115	(17)	0.3299	(08)	0.1250	(62)	0.020
C (26)	-0.1174	(18)	0.3307	(10)	0.2465	(68)	0.020
C (27)	-0.0102	(23)	0.4244	(13)	-0.1679	(53)	0.021
C (28)	-0.0316	(21)	0.2499	(12)	-0.2603	(52)	0.020
C (29)	-0.0232	(19)	0.1676	(11)	-0.2079	(75)	0.021
C (30)	-0.2556	(15)	0.1949	(09)	-0.1076	(43)	0.021
C (31)	-0.0793	(16)	0.4732	(09)	0.0565	(52)	0.020
C (32)	-0.3206	(21)	0.3252	(10)	-0.14516	(56)	

TEMPERATURE PARAMETERS OF THE NON-H CRY

U	U	U	U ₂₃	U ₃₁	U ₁₂
0.0131	0.0069	0.0755	0.0185	-0.0191	-0.0410

Table 3

Methiodide of 7-acetoxyl-N-Yohimbine:
Bond Lengths (Å). Estimated standard
deviations are given as units in the
last place.

C(1)	C(2)	1.372
C(2)	C(3)	1.374
C(3)	C(4)	1.374
C(4)	C(5)	1.374
C(5)	C(6)	1.374
C(6)	C(7)	1.374
C(7)	C(8)	1.374
C(8)	C(9)	1.374
C(9)	C(10)	1.374
C(10)	C(11)	1.374
C(11)	C(12)	1.374
C(12)	C(13)	1.374
C(13)	C(14)	1.374
C(14)	C(15)	1.374
C(15)	C(16)	1.374
C(16)	C(17)	1.374
C(17)	C(18)	1.374
C(18)	C(19)	1.374
C(19)	C(20)	1.374
C(20)	C(21)	1.374
C(21)	C(22)	1.374
C(22)	C(23)	1.374
C(23)	C(24)	1.374
C(24)	C(25)	1.374
C(25)	C(26)	1.374
C(26)	C(27)	1.374
C(27)	C(28)	1.374
C(28)	C(29)	1.374
C(29)	C(30)	1.374
C(30)	C(31)	1.374
C(31)	C(32)	1.374
C(32)	C(33)	1.374
C(33)	C(34)	1.374
C(34)	C(35)	1.374
C(35)	C(36)	1.374
C(36)	C(37)	1.374
C(37)	C(38)	1.374
C(38)	C(39)	1.374
C(39)	C(40)	1.374
C(40)	C(41)	1.374
C(41)	C(42)	1.374
C(42)	C(43)	1.374
C(43)	C(44)	1.374
C(44)	C(45)	1.374
C(45)	C(46)	1.374
C(46)	C(47)	1.374
C(47)	C(48)	1.374
C(48)	C(49)	1.374
C(49)	C(50)	1.374
C(50)	C(51)	1.374
C(51)	C(52)	1.374
C(52)	C(53)	1.374
C(53)	C(54)	1.374
C(54)	C(55)	1.374
C(55)	C(56)	1.374
C(56)	C(57)	1.374
C(57)	C(58)	1.374
C(58)	C(59)	1.374
C(59)	C(60)	1.374
C(60)	C(61)	1.374
C(61)	C(62)	1.374
C(62)	C(63)	1.374
C(63)	C(64)	1.374
C(64)	C(65)	1.374
C(65)	C(66)	1.374
C(66)	C(67)	1.374
C(67)	C(68)	1.374
C(68)	C(69)	1.374
C(69)	C(70)	1.374
C(70)	C(71)	1.374
C(71)	C(72)	1.374
C(72)	C(73)	1.374
C(73)	C(74)	1.374
C(74)	C(75)	1.374
C(75)	C(76)	1.374
C(76)	C(77)	1.374
C(77)	C(78)	1.374
C(78)	C(79)	1.374
C(79)	C(80)	1.374
C(80)	C(81)	1.374
C(81)	C(82)	1.374
C(82)	C(83)	1.374
C(83)	C(84)	1.374
C(84)	C(85)	1.374
C(85)	C(86)	1.374
C(86)	C(87)	1.374
C(87)	C(88)	1.374
C(88)	C(89)	1.374
C(89)	C(90)	1.374
C(90)	C(91)	1.374
C(91)	C(92)	1.374
C(92)	C(93)	1.374
C(93)	C(94)	1.374
C(94)	C(95)	1.374
C(95)	C(96)	1.374
C(96)	C(97)	1.374
C(97)	C(98)	1.374
C(98)	C(99)	1.374
C(99)	C(100)	1.374

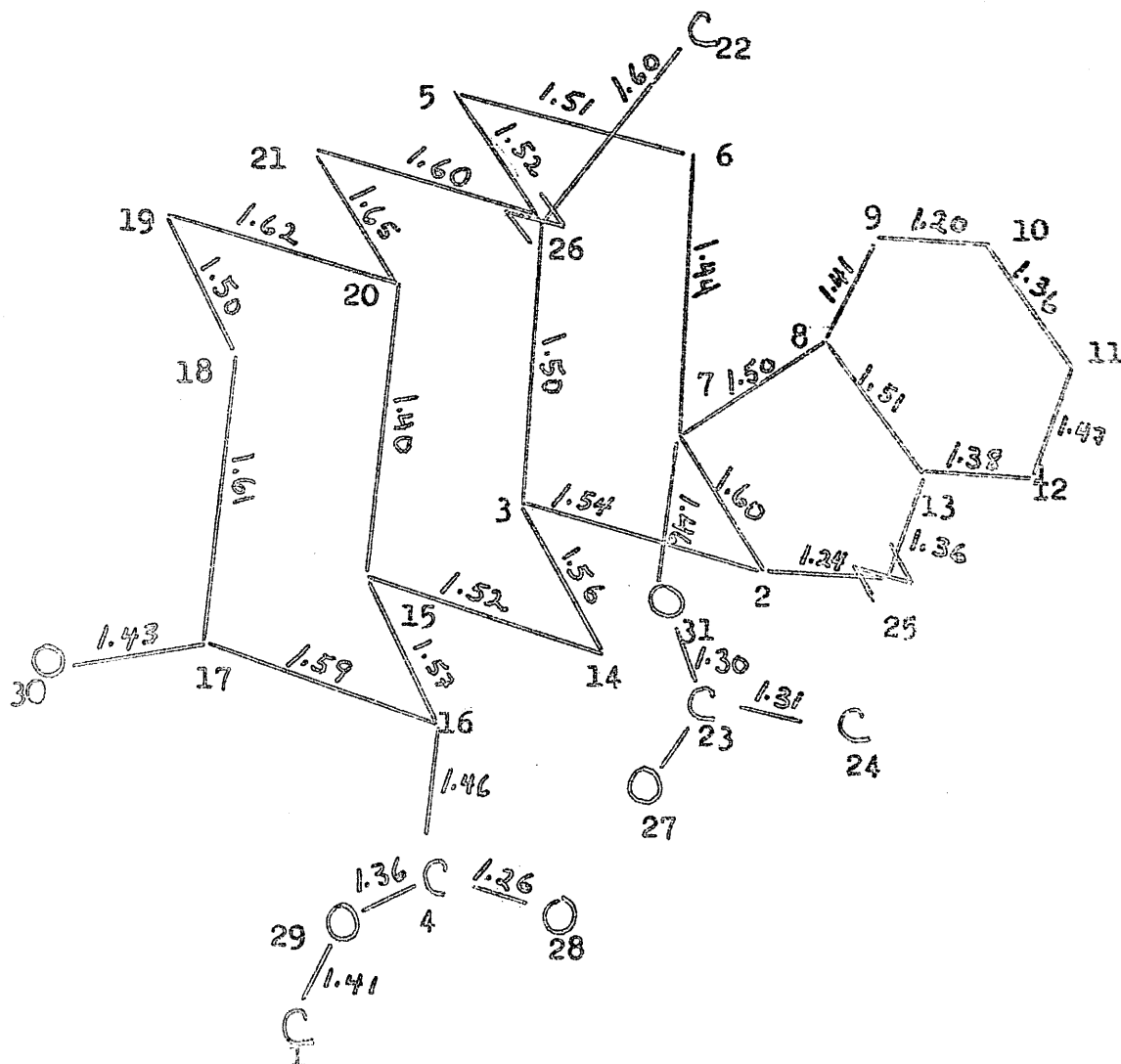
TABLE I

Boothide of 7-Acetyl-7-ethyl-Yohimbine

Bond angles and their standard deviations

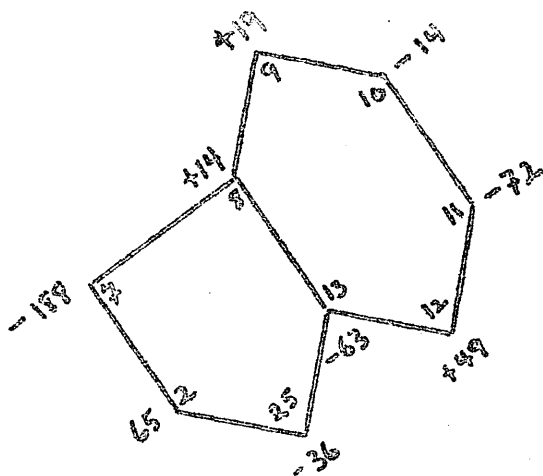
		(^o)	S. D.
C(2)	C(3)	109.2	0.2
C(3)	C(2)	114.4	0.2
C(2)	C(3)	118.2	0.2
C(3)	C(2)	109.5	0.2
C(2)	C(3)	112.4	0.2
C(3)	C(2)	121.0	0.2
C(2)	C(3)	124.0	0.2
C(3)	C(2)	138.2	0.2
C(2)	C(3)	103.7	0.2
C(3)	C(2)	110.6	0.2
C(2)	C(3)	111.5	0.2
C(3)	C(2)	111.8	0.2
C(2)	C(3)	123.5	0.2
C(3)	C(2)	123.7	0.2
C(2)	C(3)	120.3	0.2
C(3)	C(2)	115.7	0.2
C(2)	C(3)	107.0	0.2
C(3)	C(2)	115.0	0.2
C(2)	C(3)	112.8	0.2
C(3)	C(2)	110.4	0.2
C(2)	C(3)	113.8	0.2
C(3)	C(2)	95.0	0.2
C(2)	C(3)	107.1	0.2
C(3)	C(2)	107.1	0.2
C(2)	C(3)	111.4	0.2
C(3)	C(2)	111.0	0.2
C(2)	C(3)	133.5	0.2
C(3)	C(2)	111.3	0.2
C(2)	C(3)	116.3	0.2
C(3)	C(2)	131.5	0.2
C(2)	C(3)	127.8	0.2
C(3)	C(2)	115.7	0.2
C(2)	C(3)	122.0	0.2
C(3)	C(2)	116.3	0.2
C(2)	C(3)	102.7	0.2

Figure 6



Bond lengths in the Methiodide of 7-Acetoxy-7-H-Yohimbine.

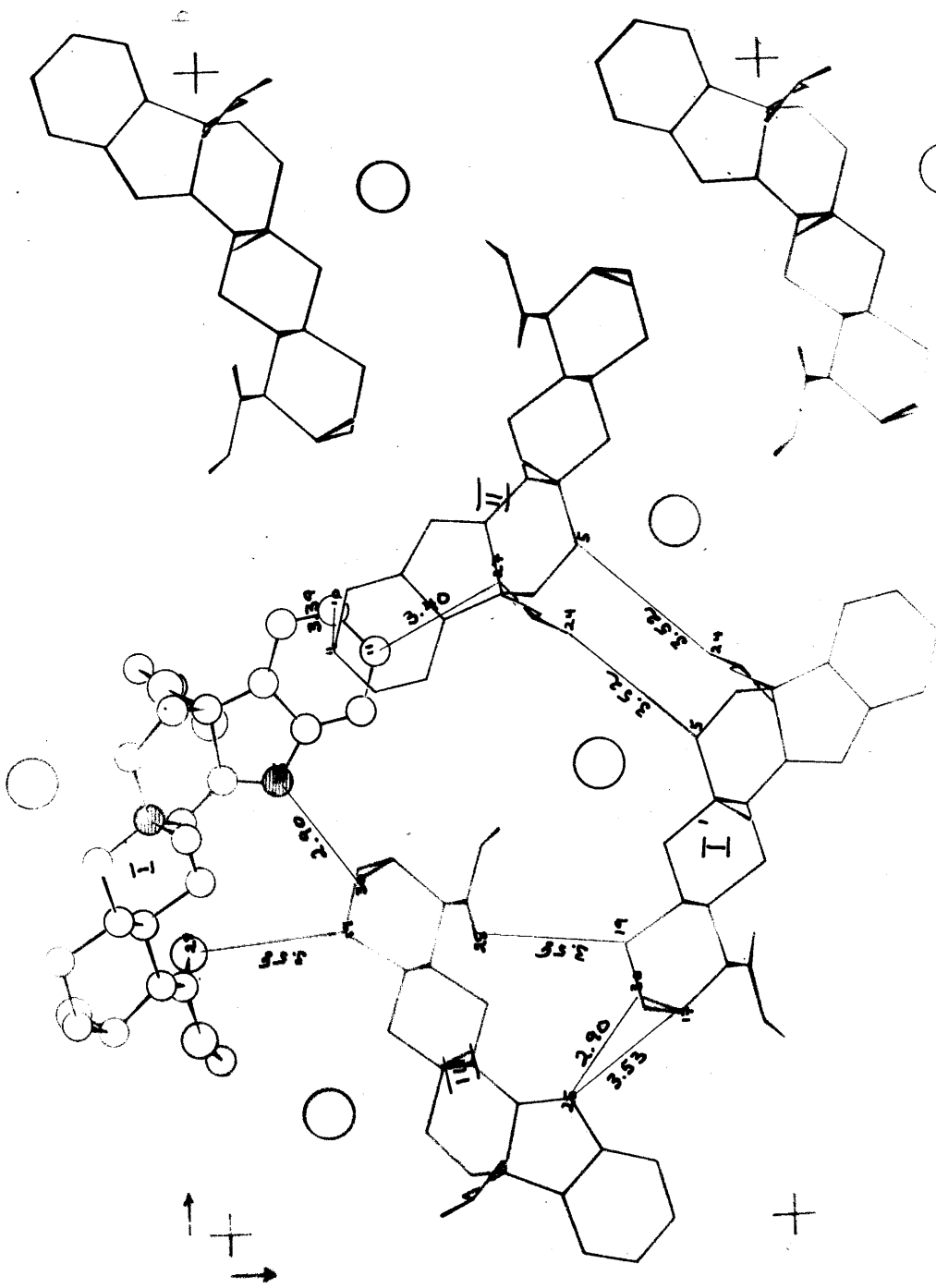
Figure 7



Methiodide of 7-Acetoxy-7-N-Yohimbine:
Weighted Mean Plane.

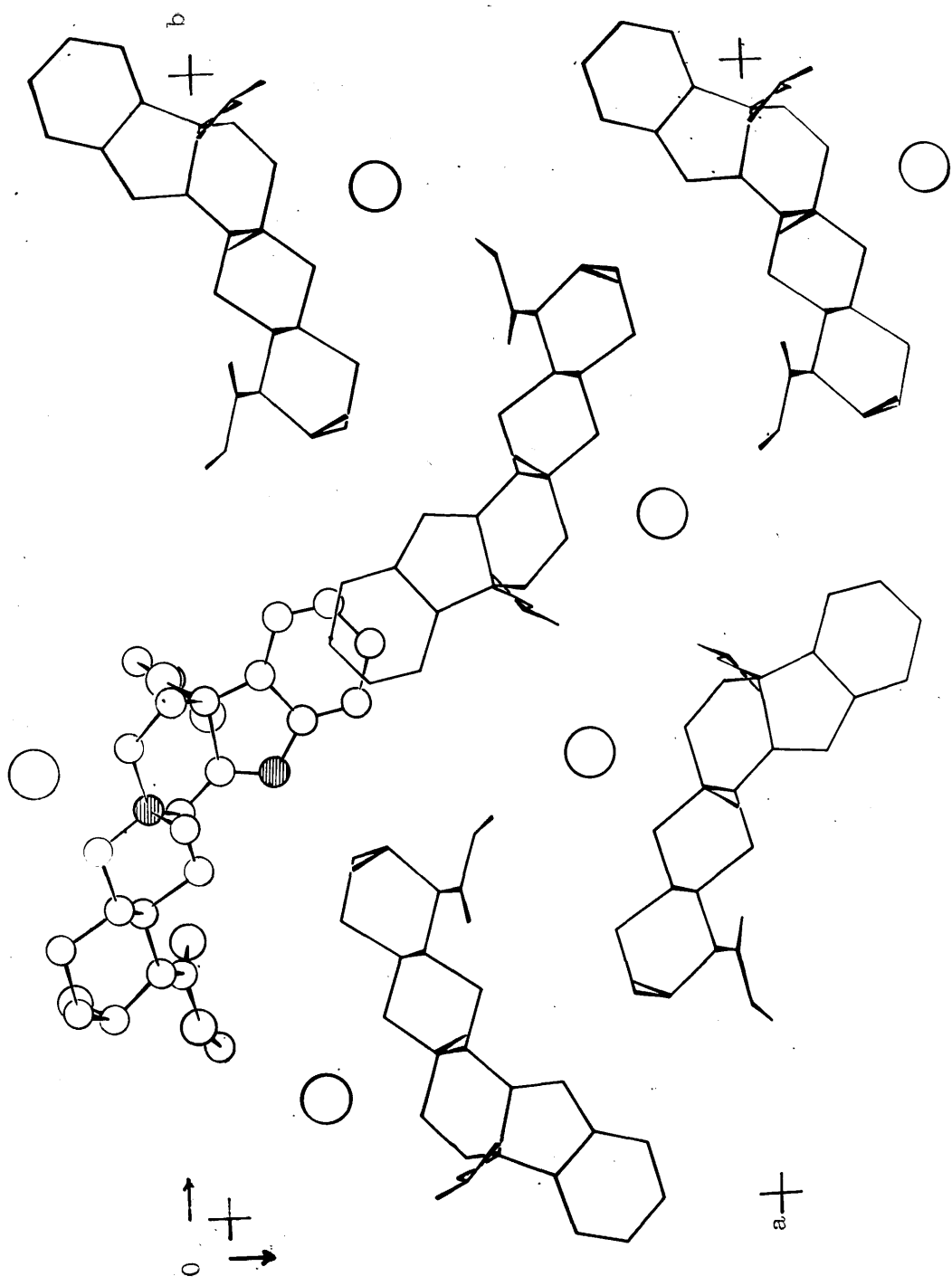
Deviations from the plane through the benzene ring are shown in units of angstroms x 10⁻³.
The equation of the plane is:

$$0.0195 X + 0.1207 Y - 0.9925 Z = 0.1275 .$$



Inter-molecular contacts. Roman numerals refer to equivalent position numbers.

-50-
Figure 8



Crystal Structure of the methiodide of 7α-acetoxy-7H-yehiibine
viewed in projection along the c-axis.

Table 5

Methiodide of 7-Acetoxy-7-II-Yohimbine:

Inter-molecular Non-bonded Distances (A)

Contacts are between the molecule in equivalent position 1 and a molecule related to it by the symmetry operation indicated by the equivalent position number. The three numbers in brackets refer to the unit cell translations. The equivalent positions are given.

1. $x, y, z,$
2. $1/2-x, -y, 1/2+z$
3. $1/2+x, 1/2-y, -z$
4. $-x, 1/2+y, 1/2-z$

Contacts from equivalent position 1, cell (0,0,0).

		e.p.	cell	
C(10)	C(11)	2	(0,1,0)	3.39 A
C(11)	O(27)	2	(0,1,0)	3.40
N(25)	O(30)	3	(0,0,0)	2.90
O(28)	C(19)	3	(0,0,0)	3.58

Contacts from equivalent position 1, cell (1,0,0).

C(5)	C(24)	2	(0,1,0)	3.52 A
C(17)	N(25)	3	(0,0,0)	3.53
C(19)	O(28)	3	(0,0,0)	3.58
C(24)	C(5)	2	(0,1,-1)	3.52
O(30)	N(25)	3	(0,0,0)	2.90

Figure 9

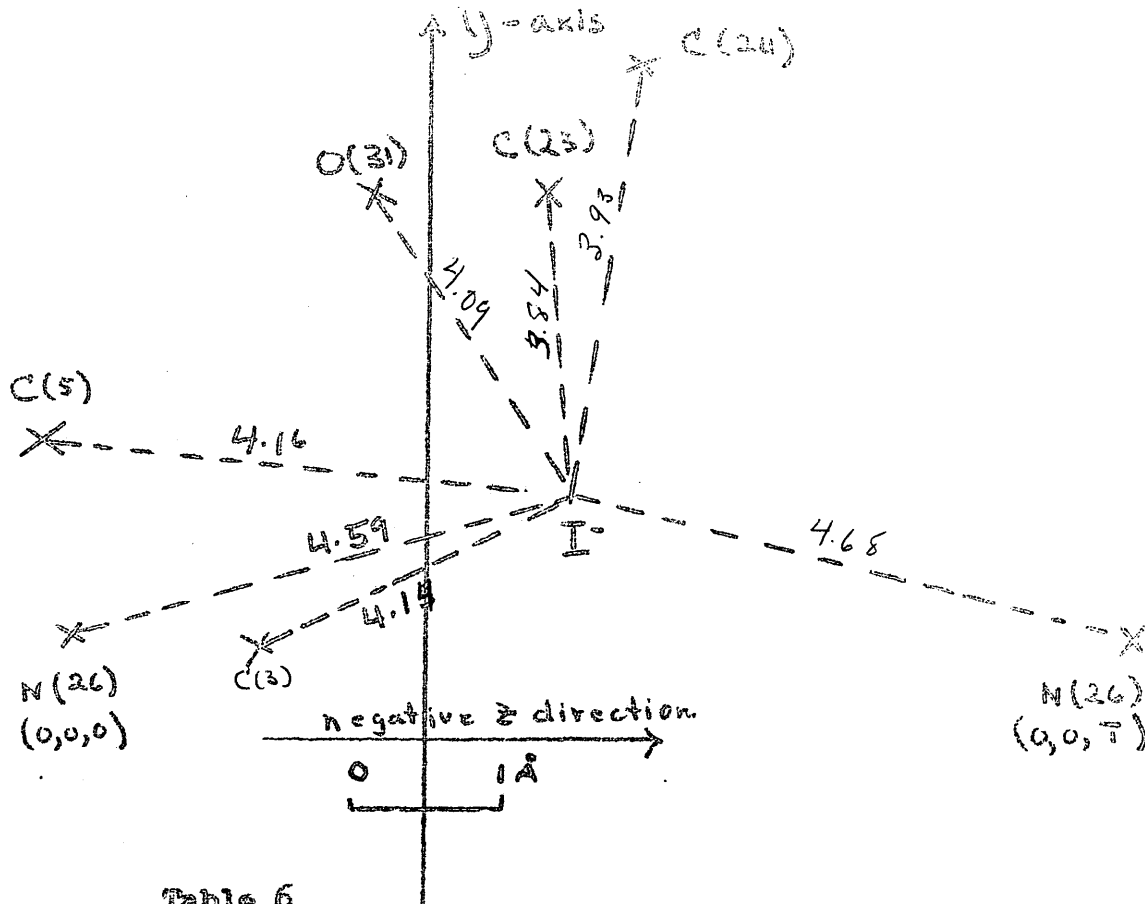


Table 6

Environment of the Iodide Ion:

Contacts with atoms in equivalent position 1.

C(23)	3.84 Å
C(24)	3.93
O(31)	4.09
C(3)	4.14
C(5)	4.16

Closest contacts with quaternary nitrogen

N(26)	1	(0,0,0)	4.59
N(26)	1	(0,0,-1)	4.68

Table 7

The methiodide of 7-acetoxy-7-yohimbine

H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc
2	27	0	10.3	13.8	6	30	0	16.0	11.9	12	4	0	12.2	13.6	3	4	1	108.9	117.2	7	11	1	28.6	27.7
2	28	0	11.2	9.0	7	1	0	83.2	99.6	12	5	0	9.6	2.3	3	5	1	96.2	91.7	7	12	1	41.5	41.8
2	29	0	10.0	11.9	7	2	0	8.8	9.3	12	6	0	12.1	15.3	3	6	1	113.3	111.5	7	13	1	51.6	50.7
2	30	0	18.2	16.8	7	3	0	38.5	43.7	12	7	0	27.5	32.2	3	7	1	14.6	20.8	7	14	1	19.0	14.0
2	31	0	7.9	10.5	7	4	0	29.6	26.3	12	8	0	18.7	18.3	3	8	1	57.4	55.5	7	15	1	15.7	18.3
2	32	0	71.9	75.6	7	5	0	68.5	71.6	12	9	0	10.3	14.5	3	9	1	52.6	52.9	7	16	1	29.1	34.1
2	2	2	124.6	145.9	7	7	0	49.4	51.4	12	13	0	22.9	20.6	3	10	1	38.5	37.7	7	18	1	26.2	27.1
2	3	0	25.6	33.1	7	9	0	55.4	63.9	12	14	0	11.3	8.6	3	11	1	35.3	37.7	7	19	1	16.3	15.5
2	4	0	113.1	130.3	7	9	0	85.3	75.1	12	17	0	18.3	20.7	3	12	1	25.1	20.9	7	23	1	10.7	19.3
2	5	0	83.6	110.4	7	10	0	29.1	22.6	12	20	0	8.9	8.1	3	13	1	63.4	56.6	7	21	1	23.0	22.1
2	6	0	54.6	54.5	7	11	0	42.8	45.4	12	22	0	8.4	9.7	3	14	1	75.9	75.6	7	22	1	35.0	27.1
2	7	0	21.8	28.8	7	12	0	16.8	18.2	13	1	0	8.4	11.6	3	15	1	37.8	48.1	7	23	1	15.7	13.3
2	8	0	75.2	75.8	7	13	0	73.1	60.9	13	2	0	13.3	13.1	3	16	1	44.8	46.8	7	27	1	28.2	23.7
2	9	0	70.6	74.7	7	14	0	22.2	14.6	13	4	0	13.2	15.8	3	18	1	54.0	45.4	8	0	1	21.2	23.9
2	10	0	51.4	42.0	7	15	0	52.9	47.4	13	5	0	14.7	15.3	3	19	1	35.3	41.4	8	0	1	26.7	27.3
2	11	0	48.2	47.9	7	16	0	47.8	47.9	13	9	0	12.8	16.2	3	20	1	19.1	23.2	8	8	1	61.3	60.1
2	12	0	115.1	149.0	7	18	0	15.0	17.1	13	10	0	9.9	5.9	3	23	1	29.6	29.7	8	8	3	46.6	47.7
2	14	0	18.2	19.3	7	19	0	39.8	42.8	13	12	0	15.2	18.3	3	24	1	33.2	32.9	8	4	1	39.5	33.7
2	15	0	50.1	48.8	7	21	0	24.0	22.7	13	13	0	9.3	11.1	3	25	1	21.4	19.8	8	5	1	17.0	20.6
2	16	0	58.4	60.0	7	23	0	26.7	28.3	13	14	0	10.2	12.5	3	26	1	12.5	9.0	8	5	1	64.1	51.4
2	18	0	50.0	53.6	7	24	0	16.0	21.3	0	3	1	80.5	86.5	4	4	0	108.5	131.4	8	7	1	31.8	26.0
2	19	0	41.2	36.3	8	25	0	15.4	12.9	0	4	1	93.8	100.2	4	4	1	88.9	87.8	8	8	1	49.5	46.6
2	20	0	29.2	28.2	8	26	0	10.9	55.5	0	5	1	31.3	32.1	4	8	1	31.3	31.1	8	9	1	69.1	23.3
2	21	0	33.8	29.6	8	2	0	12.1	2.9	0	6	1	92.6	95.9	4	3	3	25.1	18.9	8	10	1	13.1	19.3
2	22	0	71.9	57.3	8	8	2	10.9	8.6	0	7	1	57.6	61.2	4	4	4	57.1	48.0	8	11	1	30.4	28.7
2	23	0	9.6	7.1	8	8	3	40.3	43.4	0	8	1	91.9	99.7	4	5	1	96.4	113.3	8	12	1	56.2	43.6
2	24	0	13.6	8.4	8	8	4	40.7	43.2	0	9	1	66.6	60.9	4	6	1	61.0	60.5	8	13	1	37.3	41.9
2	25	0	12.1	11.7	8	8	6	47.7	48.0	0	10	1	31.1	21.2	4	7	1	43.0	35.4	8	15	1	32.5	21.5
2	26	0	21.0	28.4	8	8	7	18.2	11.9	0	11	1	132.0	132.9	4	8	1	29.5	30.1	8	16	1	31.1	32.4
2	28	0	8.7	9.5	8	8	8	40.9	34.9	0	12	1	54.7	116.8	4	9	1	125.2	124.1	8	18	1	33.2	32.4
2	29	0	22.2	18.2	8	8	9	12.2	16.4	0	13	1	41.2	48.1	4	10	1	39.5	29.7	8	22	1	61.3	28.7
2	30	0	7.7	11.2	8	8	10	50.2	48.2	0	14	1	16.6	17.3	4	11	1	61.3	69.7	8	27	1	22.8	19.4
2	32	0	9.8	9.0	8	11	0	31.7	33.6	0	15	1	31.7	29.2	4	12	1	31.0	35.4	9	0	1	27.2	26.5
2	0	0	22.6	16.8	8	12	0	22.3	19.7	0	16	1	53.6	51.9	4	13	1	36.1	31.6	9	1	1	29.0	27.5
2	1	0	162.2	163.7	8	14	0	43.1	41.9	0	17	1	44.6	43.6	4	14	1	43.1	39.4	9	2	1	36.8	35.2
2	2	0	41.8	55.0	8	15	0	9.4	8.0	0	18	1	28.6	34.1	4	15	1	63.3	77.7	9	3	1	34.6	30.1
2	3	0	100.0	119.8	8	16	0	39.5	37.9	0	21	1	35.5	32.2	4	16	1	51.9	47.4	9	4	1	41.7	39.8
2	4	0	10.3	11.7	8	18	0	21.2	16.4	0	22	1	21.1	24.2	4	17	1	29.5	29.7	9	5	1	35.5	35.2
2	5	0	58.7	44.9	8	20	0	21.3	15.9	0	24	1	10.3	8.2	4	18	1	26.6	22.2	9	6	1	35.5	35.2
2	6	0	87.9	96.3	8	23	0	8.9	1.8	0	25	1	24.5	26.3	4	19	1	44.7	41.6	9	7	1	28.8	26.9
2	8	0	118.4	118.9	8	24	0	8.6	12.2	0	26	1	39.5	27.0	4	20	1	22.4	29.5	9	8	1	15.1	15.7
2	9	0	17.8	8.5	8	25	0	8.1	11.1	0	27	1	38.6	21.9	4	21	1	14.4	17.6	9	9	1	27.8	28.5
2	10	0	36.9	39.2	8	26	0	17.1	15.4	0	28	1	12.0	13.5	4	22	1	26.5	25.8	9	10	1	17.4	20.3
2	11	0	49.7	63.5	8	27	0	17.7	17.9	0	32	1	8.8	8.8	4	24	1	20.4	19.8	9	11	1	35.9	32.4
2	12	0	12.3	12.3	8	28	0	12.8	17.7	0	33	1	112.2	126.3	4	26	1	15.9	10.5	9	13	1	32.7	24.9
2	13	0	42.8	53.7	8	29	0	7.8	8.8	1	1	1	55.1	30.6	5	0	1	35.2	41.9	9	14	1	21.4	17.3
2	14	0	19.0	16.3	9	1	0	43.5	47.8	1	2	1	80.7	83.0	5	1	1	61.5	65.5	9	15	1	23.5	25.2
2	15	0	21.5	14.9	9	2	0	48.9	49.0	1	3	1	181.6	177.5	5	2	1	51.1	46.6	9	18	1	25.6	23.7
2	16	0	42.8	33.4	9	3	0	14.9	18.0	1	4	1	143.9	135.9	5	3	1	62.9	76.8	9	19	1	22.6	21.7
2	17	0	56.0	60.9	9	4	0	12.0	7.4	1	5	1	54.2	51.1	5	4	1	75.2	61.6	9	20	1	14.0	14.9
2	19	0	29.1	27.8	9	5	0	44.6	43.4	1	6	1	123.6	100.5	5	5	1	32.7	32.5	9	24	1	13.1	13.8
2	20	0	48.3	26.8	9	6	0	48.3	43.4	1	7	1	146.6	130.1	5	6	1	67.4	69.8	9	23	1	18.6	13.6
2	21	0	62.3	26.8	9	8	0	51.3	49.9	1	8	1	56.4	65.7	5	7	1	39.9	43.8	9	24	1	13.3	12.7
2	22	0	6.8	12.9	9	9	0	20.0	16.3	1	9	1	71.1	82.1	5	8	1	36.8	41.2	10	0	1	24.9	23.3
2	23	0	12.2	15.8	9	10	0	22.8	19.7	1	10	1	80.9	80.4	5	9	1	53.8	58.0	10	1	1	31.1	32.9
2	24	0	19.2	16.5	9	11	0	33.4	31.1	1	11	1	67.9	78.4	5	10	1	76.6	56.0	10	2	1	15.3	12.1
2	27	0	28.8	30.1	9	12	0	38.9	35.6	1	12	1	71.7	77.1	5	11	1	65.4	62.8	10	3	1	17.3	18.3
2	28	0	12.0	13.8	9	13	0	27.0	19.6	1	13	1	32.0	39.2	5	12	1	32.7	33.5	10	4	1	11.7	13.0
2	29	0	62.3	26.8	9	14	0	55.3	47.9	1	14	1	55.3	55.9	5	13	1	24.0	26.7	10	5	1	51.4	42.4
2	30	0	124.6	127.0	9	16	0	40.1	40.8	1	15	1	45.9	38.5	5	14	1	56.8	51.6	10	6	1	28.5	29.7
2	3	0	69.5	76.9	9	17	0	10.7	10.2	1	16	1	53.9	46.2	5	15	1	58.7	63.5	10	7	1	15.8	13.8
2	4	0	54.9	58.8	9	18	0	13.5	11.1	1	17	1	59.3	60.3	5	16	1	48.2	46.1	10	8	1	14.2	17.3
2	5	0	31.0	39.9	9	19	0	14.9	13.8	1	18	1	34.0	33.8	5	17	1	27.3	25.1	10	9	1	46.0	47.7
2	6	0	105.4	100.0	9	20	0	13.1	16.2	1	19	1	37.7	27.3	5	18	1	28.0	25.5	10	10	1	18.0	

Table 7

The methiodide of 7-acetoxy-7-H-yohimbine Ctd.

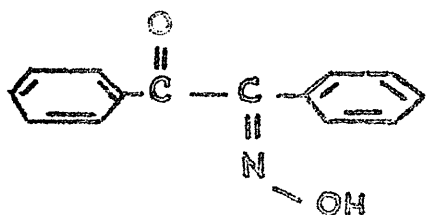
H	K	L	[Po]	[Pc]	H	K	L	[Po]	[Pc]	H	K	L	[Po]	[Pc]	H	K	L	[Po]	[Pc]	H	K	L	[Po]	[Pc]
0	7	8	12	8	4	18	2	55.1	44.1	10	6	2	32.9	47.7	4	11	3	30.6	30.3	1	11	4	42.0	39.4
0	8	9	15.7	48.9	4	19	2	17.1	15.9	10	7	2	19.5	20.1	4	12	3	20.1	22.5	1	13	4	29.2	20.0
0	9	10	28.2	66.2	4	20	2	17.4	22.2	10	8	2	22.0	31.5	4	13	3	19.8	23.4	1	14	4	45.2	31.9
0	10	11	28.4	85.0	4	21	2	19.7	21.5	10	9	2	19.7	14.0	4	14	3	34.1	43.9	1	15	4	27.3	30.7
0	11	12	78.4	74.2	4	22	2	36.0	42.9	10	10	2	22.3	32.4	4	15	3	31.5	27.7	2	16	4	27.1	26.0
0	12	13	20.0	15.3	5	0	0	50.2	52.6	11	1	2	12.5	20.9	4	16	3	34.4	32.3	2	17	4	30.9	34.6
0	13	14	34.3	17.5	5	1	0	64.8	65.9	11	2	2	27.9	38.2	4	17	3	20.3	27.7	2	18	4	27.3	23.5
0	14	15	46.1	40.9	5	2	0	36.0	36.5	11	3	2	6.8	16.3	4	18	3	23.2	28.4	3	19	4	49.5	63.5
0	15	16	68.7	79.1	5	3	0	74.2	85.0	11	4	2	8.9	11.9	4	19	3	23.4	26.4	4	20	4	47.7	50.2
0	16	17	16.8	13.0	5	4	0	56.6	56.6	11	5	2	6.9	18.1	4	20	3	33.9	29.2	5	21	4	21.4	25.1
0	17	18	26.6	25.7	5	5	0	38.1	32.9	11	6	2	15.4	24.7	4	21	3	53.4	55.0	6	22	4	39.0	42.4
0	18	19	43.1	48.1	5	6	0	71.0	58.6	11	7	2	185.5	110.6	4	22	3	48.1	42.2	7	23	4	46.6	60.1
0	19	20	13.8	20.3	5	7	0	81.8	68.5	11	8	2	19.8	19.1	4	23	3	32.9	26.7	8	24	4	23.3	24.3
0	20	21	25.8	23.6	5	8	0	75.4	59.6	11	9	2	144.9	148.7	4	24	3	37.5	33.3	9	25	4	31.0	36.8
0	21	22	22.9	26.7	5	9	0	53.9	48.8	11	10	2	55.2	65.7	4	25	3	50.8	50.4	10	26	4	29.1	38.3
0	22	23	192.5	250.8	10	0	0	92.5	42.8	11	11	2	48.6	42.0	4	26	3	17.9	15.7	11	27	4	34.4	38.2
1	1	1	85.9	80.0	10	1	0	51.3	48.1	11	12	2	21.1	6.8	4	27	3	23.7	14.7	12	28	4	25.7	28.1
1	1	1	66.8	82.0	10	2	0	18.3	17.3	11	13	2	66.1	51.1	4	28	3	33.6	31.7	13	29	4	37.5	45.7
1	1	1	97.9	130.6	10	3	0	49.1	42.1	11	14	2	89.7	98.7	4	29	3	66.6	71.0	14	30	4	27.8	31.3
1	1	1	111.3	130.6	10	4	0	111.3	130.6	11	15	2	41.0	33.7	4	30	3	45.6	49.9	15	31	4	32.2	36.1
1	1	1	59.3	53.4	10	5	0	36.1	24.8	11	16	2	62.3	49.2	4	31	3	54.3	49.0	16	32	4	21.0	18.1
1	1	1	63.9	85.1	10	6	0	42.9	38.1	11	17	2	12.0	10.2	4	32	3	62.7	51.5	17	33	4	27.0	35.4
1	1	1	65.0	78.7	10	7	0	19.1	19.7	11	18	2	79.2	70.3	4	33	3	23.3	23.5	18	34	4	29.3	30.2
1	1	1	36.2	35.3	10	8	0	39.7	29.3	11	19	2	16.8	12.6	4	34	3	21.3	27.3	19	35	4	64.5	58.0
1	1	1	56.0	38.2	10	9	0	24.9	22.0	11	20	2	44.3	48.5	4	35	3	40.9	49.5	20	36	4	56.0	47.7
1	1	1	76.0	66.6	10	10	0	22.0	20.7	11	21	2	25.0	25.0	4	36	3	15.3	14.1	21	37	4	44.8	41.0
1	1	1	33.5	62.1	10	11	0	31.1	32.1	11	22	2	38.4	41.1	4	37	3	35.3	37.4	22	38	4	34.2	32.2
1	1	1	43.3	39.0	10	12	0	19.3	17.2	11	23	2	27.0	32.9	4	38	3	12.7	29.3	23	39	4	48.1	45.0
1	1	1	43.3	53.8	10	13	0	31.1	32.1	11	24	2	58.8	74.1	4	39	3	36.3	35.0	24	40	4	29.8	33.4
1	1	1	82.5	81.3	10	14	0	81.4	91.2	11	25	2	15.5	14.8	4	40	3	90.8	69.8	25	41	4	74.9	66.9
1	1	1	26.6	15.7	10	15	0	57.0	66.2	11	26	2	18.3	20.4	4	41	3	12.2	28.9	26	42	4	25.5	36.3
1	1	1	34.6	31.6	10	16	0	45.6	47.7	11	27	2	49.1	59.2	4	42	3	37.7	33.5	27	43	4	47.0	53.7
1	1	1	38.4	35.3	10	17	0	35.4	35.5	11	28	2	29.3	29.3	4	43	3	55.2	57.8	28	44	4	52.7	42.3
1	1	1	16.8	16.8	10	18	0	48.0	48.0	11	29	2	51.7	66.2	4	44	3	19.1	18.3	29	45	4	39.8	33.3
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1	1	1	14.6	15.6	10	21	0	12.9	14.0	11	32	2	41.3	39.4	4	47	3	30.8	34.4	32	48	4	28.0	24.3
1	1	1	20.5	29.5	10	22	0	42.5	39.9	11	33	2	84.7	79.4	4	48	3	35.3	31.2	33	49	4	46.1	43.1
1	1	1	20.5	25.7	10	23	0	21.7	24.3	11	34	2	26.3	30.0	4	49	3	28.9	32.7	34	50	4	21.3	30.3
1	1	1	17.7	15.4	10	24	0	31.9	34.9	11	35	2	44.7	40.0	4	50	3	40.9	49.5	35	51	4	17.7	17.7
1	1	1	68.4	75.5	10	25	0	16.4	16.4	11	36	2	34.1	36.9	4	51	3	16.4	19.8	36	52	4	53.3	57.7
1	1	1	51.2	66.0	10	26	0	15.1	13.1	11	37	2	56.5	47.8	4	52	3	32.9	41.1	37	53	4	49.1	51.5
1	1	1	69.7	90.3	10	27	0	13.6	7.4	11	38	2	79.4	72.9	4	53	3	25.1	28.0	38	54	4	20.2	21.3
1	1	1	18.2	21.2	10	28	0	39.8	40.4	11	39	2	33.6	32.9	4	54	3	21.3	19.5	39	55	4	39.8	25.9
1	1	1	81.0	91.1	10	29	0	34.1	34.2	11	40	2	22.4	31.3	4	55	3	63.7	73.3	40	56	4	28.2	27.1
1	1	1	58.1	47.9	10	30	0	14.0	10.6	11	41	2	19.5	19.5	4	56	3	33.7	33.3	41	57	4	80.0	85.8
1	1	1	69.2	78.1	10	31	0	15.7	17.1	11	42	2	27.2	41.9	4	57	3	32.2	36.5	42	58	4	46.6	36.0
1	1	1	103.6	98.6	10	32	0	17.2	21.5	11	43	2	16.8	16.3	4	58	3	9.8	12.8	43	59	4	47.6	39.9
1	1	1	48.1	47.9	10	33	0	16.2	16.0	11	44	2	32.6	37.8	4	59	3	46.8	43.4	44	60	4	38.9	37.3
1	1	1	92.1	81.6	10	34	0	34.5	62.1	11	45	2	20.0	17.7	4	60	3	53.4	48.3	45	61	4	35.4	35.7
1	1	1	39.2	35.3	10	35	0	30.7	31.6	11	46	2	18.3	14.5	4	61	3	23.2	31.9	46	62	4	51.6	45.8
1	1	1	27.0	31.0	10	36	0	24.5	20.5	11	47	2	18.7	25.9	4	62	3	13.2	29.9	47	63	4	42.7	46.7
1	1	1	78.0	71.9	10	37	0	17.1	17.3	11	48	2	133.3	131.2	4	63	3	13.5	16.6	48	64	4	27.5	29.2
1	1	1	50.5	43.7	10	38	0	50.9	65.0	11	49	2	37.3	45.7	4	64	3	12.2	13.0	49	65	4	25.3	27.0
1	1	1	25.6	15.0	10	39	0	17.7	75.8	11	50	2	65.9	48.4	4	65	3	16.6	24.7	50	66	4	49.8	40.9
1	1	1	36.5	35.3	10	40	0	12.9	10.9	11	51	2	103.9	95.6	4	66	3	33.3	33.4	51	67	4	44.5	33.2
1	1	1	29.3	29.9	10	41	0	37.9	36.9	11	52	2	40.6	34.5	4	67	3	16.8	22.1	52	68	4	31.6	27.9
1	1	1	17.0	19.4	10	42	0	13.9	18.7	11	53	2	61.2	86.3	4	68	3	16.7	16.3	53	69	4	41.9	39.9
1	1	1	17.3	19.3	10	43	0	56.6	54.4	11	54	2	77.3	79.4	4	69	3	30.7	27.7	54	70	4	31.9	27.7
1	1	1	24.8	26.1	10	44	0	23.9	23.2	11	55	2	44.8	40.8	4	70	3	40.8	38.1	55	71	4	45.5	41.7
1	1	1	17.1	17.2	10	45	0	43.2	42.3	11	56	2	29.5	37.3	4	71	3	30.9	37.7	56	72	4	26.3	26.3
1	1	1	43.8	32.3	10	46	0	27.1	25.9	11	57	2	62.3	68.9	4	72	3	26.4	24.6	57	73	4	38.3	32.8
1	1	1	60.6	67.6	10	47	0	44.6	42.0	11	58	2	17.4	17.1	4	73	3	23.8	23.0	58	74	4	27.8	22.3

Part III

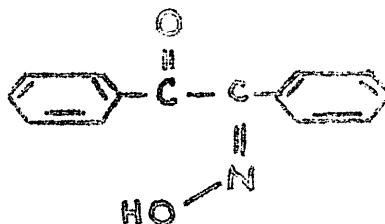
The Structure of Benzilmonoxime:
The Crystal Structure of the
Para-Bromobenzoate of Benzilmonoxime

3.1 INTRODUCTION

The currently accepted structures of the two monoximes of benzil are shown here

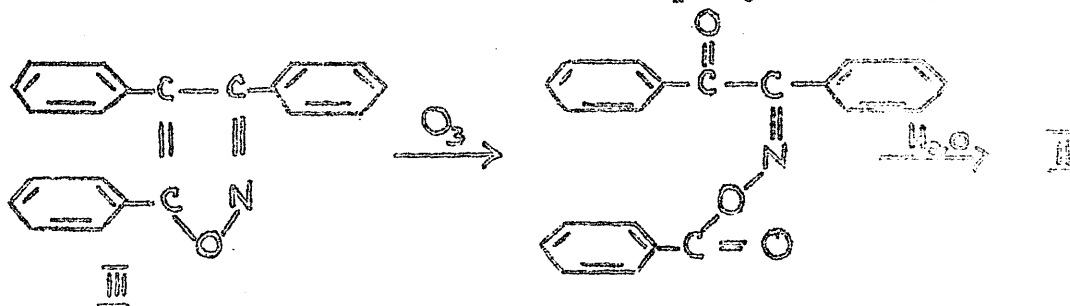


α m.p. 138°C I



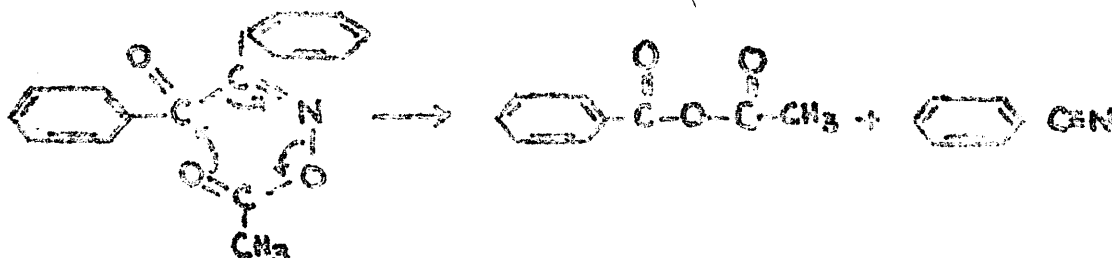
β m.p. 107°C II
(114°C polymorph)

This configurational assignment, made by Meisenheimer (1921), was based on the fact that the benzoate of the β -oxime, IV, is formed from the ozonisation of triphenylsloxazole, III.



There is some evidence that the configurations should be reversed. The acetate of β -benzilmonoxime is stable at 190°C, whereas that of the α -oxime undergoes thermal decomposition at 140°C. When the reaction products are treated with water, acetic acid, benzoic acid and benzonitrile are formed, indicating that the initial products were probably benzonitrile and acetic-benzoic anhydride. Such thermal decompositions often occur by intramolecular

cyclic mechanisms, viz.



This would imply that the hydroxyl group in the α -oxime was gig to the carbonyl group.

Some spectroscopic evidence would seem to support this argument. An investigation of some α - and β -oxime by infra-red spectroscopy (Palm and Vexlin, 1959), shows that with each isomeric pair, the β -oxime has the higher melting point and the higher conductivity. This increase in acidity of the O-H bond is accompanied by a shift to lower frequencies of the I.R. absorption bands associated with the O-H stretching mode. In most isomeric oximes, the β -oxime absorbs at lower frequency than the α -isomer. Benzilmonoxime is an exception on two counts. In solutions of benzene and chloroform, the β -oxime absorbs at higher frequencies than the α -oxime. The melting point of the ~~para-benzilmonoxime of the~~ β -oxime is only 107°C as compared with 136°C in the case of the α isomer.

Other anomalies in the behaviour of these two compounds include an abnormal Beckmann rearrangement (Blatt, 1951) and abnormal reactions of the α -oxime with trifluoroacetic acid and with polyphosphoric acid (Ferre, 1951; Under and Kilmickel, 1959 - 1959).

N.M.E. studies undertaken at Ohio State University seemed to support a change in the configuration but these were not conclusive (M.S. Newman, private communication). Professor Newman requested an X-ray analysis and sent a sample alleged to be the para-bromobenzoate of β -benzilmonoxime (m.p. 105-106°C). Barnes (1938) reports melting points of the para-bromobenzoates of α and β -monoxime. His values are as follows:

α form 105°C ,

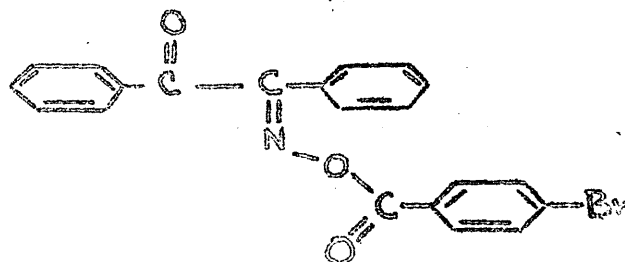
β form 147°C .

That the sample provided was the para-bromobenzoate of the α -form of the oxime was further confirmed by the fact that attempted hydrolysis of the para-bromobenzoate did not yield the parent oxime. (M.S. Newman, private communication). This behaviour is typical of derivatives of α -oximes.

The crystals provided by Professor Newman were long white needles and showed sharp extinction along the direction of the needle axis. Unfortunately, they showed a very strong tendency to grow together as twins. The specimen chosen for analysis had one individual of the twin sufficiently smaller than the other to allow accurate indexing. It had been intended to solve the structure by the usual heavy atom method. This approach failed. The structure was finally solved by combining a heavy

atom-phased Fourier synthesis with a "minimum function" derived from a three-dimensional, sharpened Patterson function. Three electron density distributions were calculated to refine the atomic positions. Further improvement of the positional and thermal parameters was obtained by the method of least squares. The final R-value was 16.0% for 1258 independent, non-zero reflections.

The results of this analysis show that the para-bromobenzoate of the α -benzilmonoxime (m.p. 105-106°C) has the para-bromobenzoate group trans relative to the carbonyl group (Figure 10).



This confirms Meisenheimer's assignment of structure to the oximes, I and II, of benzil.

3.2. EXPERIMENTAL

The para-bromobenzoate of α -benzilmonoxime yields long colourless, needle-shaped crystals when crystallised from ethanol solutions. These melt sharply at 106°C on a hot-stage microscope. When examined under a polarising microscope, they show sharp extinction along the needle direction.

3.2.1. CHOICE OF CRYSTAL

Preliminary photographs showed that the compound crystallises in the triclinic system, with space group $P1$ or $P\bar{1}$. (Subsequent calculations have shown that the space group is $P\bar{1}$). Weissenberg photographs of the $hk0$ zone were taken in the usual manner with multiple film packs (Robertson, 1943). Photographs of the hkl and $hk2$ reciprocal lattice nets showed that the crystals were twinned. This twinning occurred in such a way that there was exact registry of points in the zero layer of the reciprocal lattice, but complete disregistry in other layers. In the crystal chosen for analysis, one individual of the twin was sufficiently larger than the other to allow accurate indexing. The spot shape of the reflections in the photographs was very poor.

3.2.2. CHOICE OF UNIT CELL

In the triclinic system, the choice of unit cell is not dictated by symmetry considerations. For this reason there are an infinite number of ways of choosing a primitive cell in a triclinic lattice. However, for any crystal, there is one cell that is unique. This particular cell is called the "reduced cell" and is defined by the three shortest, non-coplanar translations in the lattice. In such a cell, each cell edge is shorter than

reciprocal lattice points into the sphere of reflection. Thus both the $\underline{a^*b^*}$ plane and the $\underline{b^*g^*}$ plane can be photographed with the same mounting. The dihedral angle between the planes is the complement of the angle between the plane normals: e.g. $[\underline{a^*b^*}] \perp \underline{g}$ and $[\underline{a^*g^*}] \perp \underline{b}$; therefore, the angle between the planes is $180-\alpha$. In the monoclinic case and in systems of higher symmetry $\alpha^* = 180-\alpha$. In the triclinic case

$$\alpha^* = \cos^{-1} (\cos \beta^* \cos \gamma^* - \sin \beta^* \sin \gamma^* \cos \alpha).$$

Thus the six reciprocal cell dimensions are available from a single mounting. The real cell dimensions can then be calculated from the formulae given by Buerger (1942, p.360).

3.2.4. CRYSTAL DATA

Formula	$C_{21}NO_3Br$	F.W. 395.23 gm.
System	triclinic	
Space group	H or $P\bar{1}$	
Lattice constants	$a = 8.78 \text{ \AA}$	
	$b = 17.01 \text{ \AA}$	
	$c = 6.27 \text{ \AA}$	
	$\alpha = 95.11^\circ$	
	$\beta = 91.08^\circ$	
	$\gamma = 102.34^\circ$	

reciprocal lattice points into the sphere of reflection. Thus both the a^*b^* plane and the b^*c^* plane can be photographed with the same mounting. The dihedral angle between the planes is the complement of the angle between the plane normals: e.g. $[a^*b^*] \perp a$ and $[b^*c^*] \perp b$; therefore, the angle between the planes is $180-\alpha$. In the monoclinic case and in systems of higher symmetry $\alpha^* = 180-\alpha$. In the triclinic case

$$d^* = \cos^{-1} (\cos \beta^* \cos \gamma^* - \sin \beta^* \sin \gamma^* \cos \alpha).$$

Thus the six reciprocal cell dimensions are available from a single mounting. The real cell dimensions can then be calculated from the formulae given by Buerger (1942, p.360).

3.2.4. CRYSTAL DATA

Formula	$C_{21}NO_3Br$	F.W. 395.23 gm.
System	triclinic	
Space group	R or $R\bar{1}$	
Lattice constants	$a = 8.78 \text{ \AA}$	
	$b = 17.01 \text{ \AA}$	
	$c = 6.27 \text{ \AA}$	
	$\alpha = 95.11^\circ$	
	$\beta = 91.08^\circ$	
	$\gamma = 102.34^\circ$	

Volume	911 Å
Density(obs)	1.47 gm/cc
Molecules/ cell	2
Density(calc)	1.49 gm/cc
F(000)	416
$\mu(\text{Cu K}\alpha)$	35.9 cm ⁻¹

3.2.5. INTENSITY DATA

Intensity data were obtained for the reciprocal lattice nets hk0 to hk5 using a Nonius Weissenberg camera and multiple film packs. To avoid the spot shape correction necessary if data are obtained from upper and lower halves of the same film, two sets of photographs were taken of each layer, one displaced from the other by a rotation of 180°. The two sets were scaled to one another by comparison of common reflections. The various reciprocal lattice nets were given a common relative scale by comparison by axial reflections from the Weissenberg data with common reflections from the precession photographs of the hkl zone. Lorentz and polarization corrections had been calculated on the DEUCE computer for the Weissenberg data. These corrections had been applied to the precession data by means of a template. (Beurger, 1960, p.185).

3.2.6. ANALYSIS OF THE STRUCTURE

Projected Patterson syntheses, $P(UV)$ and $P(VW)$, were calculated to determine three coordinates of the heavy atom. The median axis projection, calculated from precession data, showed two large peaks. Neither of these gave satisfactory agreement with the major peak in the short axis projection. A three-dimensional vector map, $P(UVW)$, showed two major peaks corresponding to the two peaks in the projected function, $P(UV)$. The larger of these was a diffuse peak, very much extended in a direction roughly parallel to the median axis. Since the smaller peak was the usual Gaussian shape it was assumed to represent the Br-Br interaction. The bromine coordinates derived from this peak were;

$$x = 0.3246, y = 0.0826, z = 0.3573.$$

Structure factors were calculated with phases determined by the heavy atom. Sim's ratio, r , for this compound

$$\text{is } r = \frac{\left(\frac{\sum F_o^2}{\sum F_c^2}\right)^{\frac{1}{2}}}{\left(\frac{1225}{1012}\right)^{\frac{1}{2}}} = 1.10.$$

indicating that about 20% of the data will have the wrong sign in a calculation based on the heavy atom alone.

(Sim, 1957). To avoid introducing errors in to the electron density distribution by including terms with the wrong sign, all structure factors with

$$|F_o| - |F_c| > \frac{1}{2}|F_o|$$

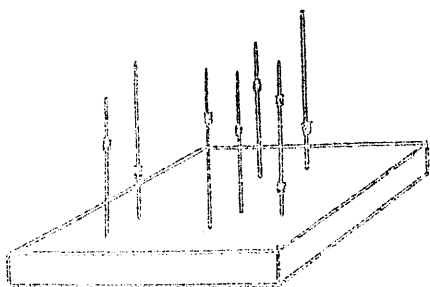
were excluded from the direct Fourier synthesis. The

R-value for the terms actually included in the summation was 53.7%.

The first Fourier map was calculated in sections along the median axis. There were 40 peaks with heights between two and six electrons/ \AA^3 . Other than the bromine atom, none of the known features of the molecule could be recognized.

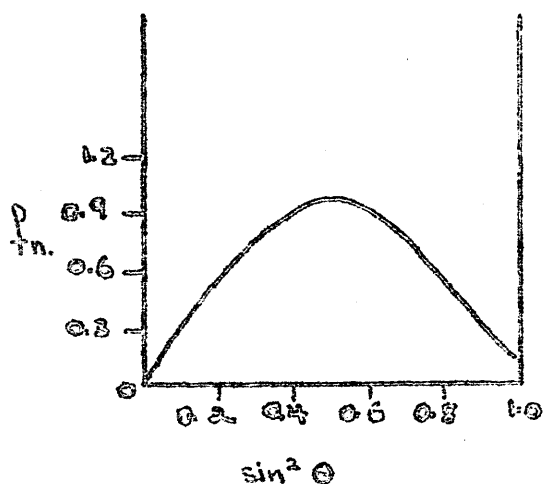
A minimum function was derived from the three-dimensional Patterson function in the manner described in Part I (1.4). Both the Fourier map and the minimum function were plotted on glass sheets, but the only obvious similarities between them were the positions of the bromine atoms and very pronounced streaks of electron density running diagonally across the maps. Although the minimum function was not well resolved, it was possible to reduce the number of atomic sites by comparing the two functions, one section at a time. Only 27 of the 40 peaks in the Fourier map corresponded to positive regions in the minimum function.

These 27 peaks were plotted in three-dimensions using long wires supported by a cork base. Two coordinates were defined by the positions of the wires relative to a grid drawn on the base. The vertical coordinate was marked by a small piece of fibre-glass electrode.



insulation sheathing. Tentative locations were assigned to 9 of the atoms in the molecule. Five of these were used in a structure-factor calculation with $hk0$ reflection data. The R -value increased. The three-dimensional data were "sharpened" by applying the Lorentz-polarisation function to the Patterson coefficients, $|P|$. This gives

a sharpening function of the form shown on the left. The sharpened Patterson function was used to give an improved position for the bromine atom and to derive a minimum function. The bromine-phased electron-density calculation and the Patterson synthesis were carried in sections along the ab



axis because it seemed likely that the streaking observed in the previous maps represented unresolved, steeply tilted benzene rings. The resolution of the minimum function had been much improved by the sharpening process.

The electron density distribution and the minimum function were superimposed by plotting equivalent sections from the two functions on the same glass sheet. This composite map revealed the positions of the 25 light atoms. Since the structure of the molecule was known, all of the atoms could be identified. When they were isolated

in the next structure factor calculation with scattering factors appropriate to their chemical types, the R-value dropped to 34.8%. Another cycle of Fourier refinement gave a further improvement to 30.1%. No further calculations were done on the DEUCE computer at Glasgow University.

3.2.7. REFINEMENT OF THE STRUCTURE

The Glasgow Structure Factor Least Squares program was available before the Fourier program. For this reason refinement was continued by least squares in spite of the high R-value.

The first cycle of least squares used unit weights for each reflection and served to provide a rough over all scale factor for the data. The R-value was 35.7%. For all subsequent calculations, Cruickshank's (1961) weighting scheme was used:

$$1/w = (1/p_1 + F + p_2 F^2 + p_3 F^3)^{1/2}$$

The initial values of the parameters were

$$p_1 = 6.0, p_2 = 9.9 \times 10^{-3}, p_3 = 1.25 \times 10^{-4}$$

These values were adjusted slightly during the course of refinement. The final values were

$$p_1 = 4.2, p_2 = 1.6 \times 10^{-2}, p_3 = 3.8 \times 10^{-4}$$

After four cycles of isotropic block-diagonal least-squares refinement, the R-value had been reduced to 26.6%. The data were rescaled. After two more cycles the R-value was 24.6%.

A program for refining projections by the "minimum residual" method of Bhuiya and Stanley (1963) was written by K.W. Muir of this Department. When this program became available, it was used to refine the projections along the short and median axes. The R-values for the two zones improved from 25% to 23% and from 27% to 22% respectively. Three-dimensional structure factors based on positions from the two projections showed an R-value of 21.9%. Three cycles of least squares reduced this to 18.2%. Anisotropic temperature factors were allowed to refine for 4 cycles of block diagonal least squares. Refinement was terminated at this point. The R-value was 16.0% for 1258 independent reflections. Unobserved reflections were not included in the analysis.

3.3 MOLECULAR DIMENSIONS AND DESCRIPTION OF THE STRUCTURE

The configuration of the para-bromobenzoate of benzil monoxime is best displayed when the crystal structure is viewed in projection along the short axis (Figure 10). Van der Waals forces are important in determining the packing arrangement of the molecule, but a very short contact between the oxygen of the carbonyl group and the carbon of the carboxyl group would seem to indicate that dipole-dipole interactions may also be important. Intermolecular contacts are seen most easily in the projection along the median axis (Figure 11). Figure 14 is an enlargement of a portion of the median axis projection.

Table 8 lists the final atomic coordinates and their standard deviations as given by the least-squares refinement. The coordinates and their standard deviations are quoted with the same number of significant figures. No significance should be attached to the last figure as it has been included only to allow derived quantities to be calculated without loss of accuracy. It should be noted that the standard deviations of the heavy atom coordinates are an order of magnitude smaller than those of the light atoms.

Anisotropic temperature parameters are shown in Table 9. These are the parameters of the equations:

$$\exp \left(-B \sin^2 \theta / \lambda^2 \right) = \exp \left[-2 \kappa^2 \left(U_{11} h^2 a^{*2} + U_{22} k^2 b^{*2} + U_{33} l^2 c^{*2} \right. \right. \\ \left. \left. + 2U_{23} k l b^* c^* + 2U_{31} l a^* c^* + 2U_{12} h k a^* b^* \right) \right]$$

The factor of 2 has been included in the cross-terms.

Bond lengths and their standard deviations are given in Table 10; bond angles are shown in Table 11. These are shown diagrammatically in Figure 12.

Table 12 is a list of ρ_{ij} , the correlation coefficients between the coordinates of each atom. The correlation coefficient between the i^{th} and j^{th} parameters in a least-squares calculation is defined as

$$\rho_{ij} = (a^{-1})_{ij} / [(a^{-1})_{ii}(a^{-1})_{jj}]^{1/2},$$

where $(a^{-1})_{ij}$ is the j^{th} element of the i^{th} row of the matrix inverse to a_{ij} , the matrix of the normal equations. The intra-atomic correlations arise primarily from non-orthogonality of the axes and from anisotropy of the thermal parameters.

Standard deviations of bond lengths in this structure lie between 0.03 and 0.04 Å. Most of the observed bond lengths are very near their expected values. Individual bond lengths in the benzene rings vary between 1.30 and 1.51 Å, with an average length of 1.41 Å. The C-Br bond length is 1.85 Å. Both of these values compare favourably with the expected values of 1.397 and 1.85 Å, respectively (Sutton, 1965). There are three C(sp₂)-C(sp₂) single bonds in the

structure with an average length of 1.49 \AA , and two C=O bonds with an average length of 1.21 \AA . These measurements are in agreement with the values of 1.492 and 1.210 \AA , respectively for bonds of these types in benzil (Brown and Sadanaga, 1965). The C-O bond length of 1.38 \AA is close to the value of 1.358 \AA reported for the long bond in esters and acids (Sutton et al., 1958). The N-O bond length of 1.42 \AA is similar to the value 1.408 \AA , reported in formaldehyde (Levine, 1963), and to the value 1.415 \AA , reported in formaldehyde oxime (Hall, 1965).

In a summary of oxime structures, Hamilton (1961) lists various lengths for the C-N bond between 1.225 and 1.28 \AA . The value predicted from Pauling's radii is 1.265 \AA . The measured bond length of $1.35 \pm 0.03 \text{ \AA}$ seems to be significantly longer than the predicted length. This may be real or it may reflect an under-estimate of the standard deviation of the bond length. Hodgson and Rollett (1962) have shown that when the variances of parameters are calculated from the inverses of the block matrices, standard deviations of bond lengths can be underestimated by as much as 80%. If the difference is real, it implies that this is not a pure double bond but that it is taking part in some sort of resonance interaction. This is not the case. Resonance interaction with the benzene ring is precluded because both H11 and H12 are 180° out of phase significantly from the plane through the

ring (Table 13). Resonance with the carbonyl group is impossible. The root mean square distance from the "best" plane through C(8), C(9) N(22) and O(23) is more than $\pm 0.25 \text{ \AA}$. This would indicate that the difference is not real and that the standard deviation of this bond length and been underestimated.

"Best" planes have been calculated for the three benzene rings and for the carboxyl group. Deviations from these planes are shown in Table 13 and Figure 13. It is interesting to note that C(17) and Br(26) lie in the plane of benzene ring A, but that the oxygens of the carboxyl group have been displaced from this plane by a rotation of 13° .

Pauling's figures for the Van der Waals radii of carbon (1.70 \AA), nitrogen (1.50 \AA) and oxygen (1.40 \AA) predict that the shortest inter-molecular contacts will be as follows:

C...C	3.40 \AA	C...N	3.20 \AA
C...O	3.12 \AA	O...O	2.80 \AA

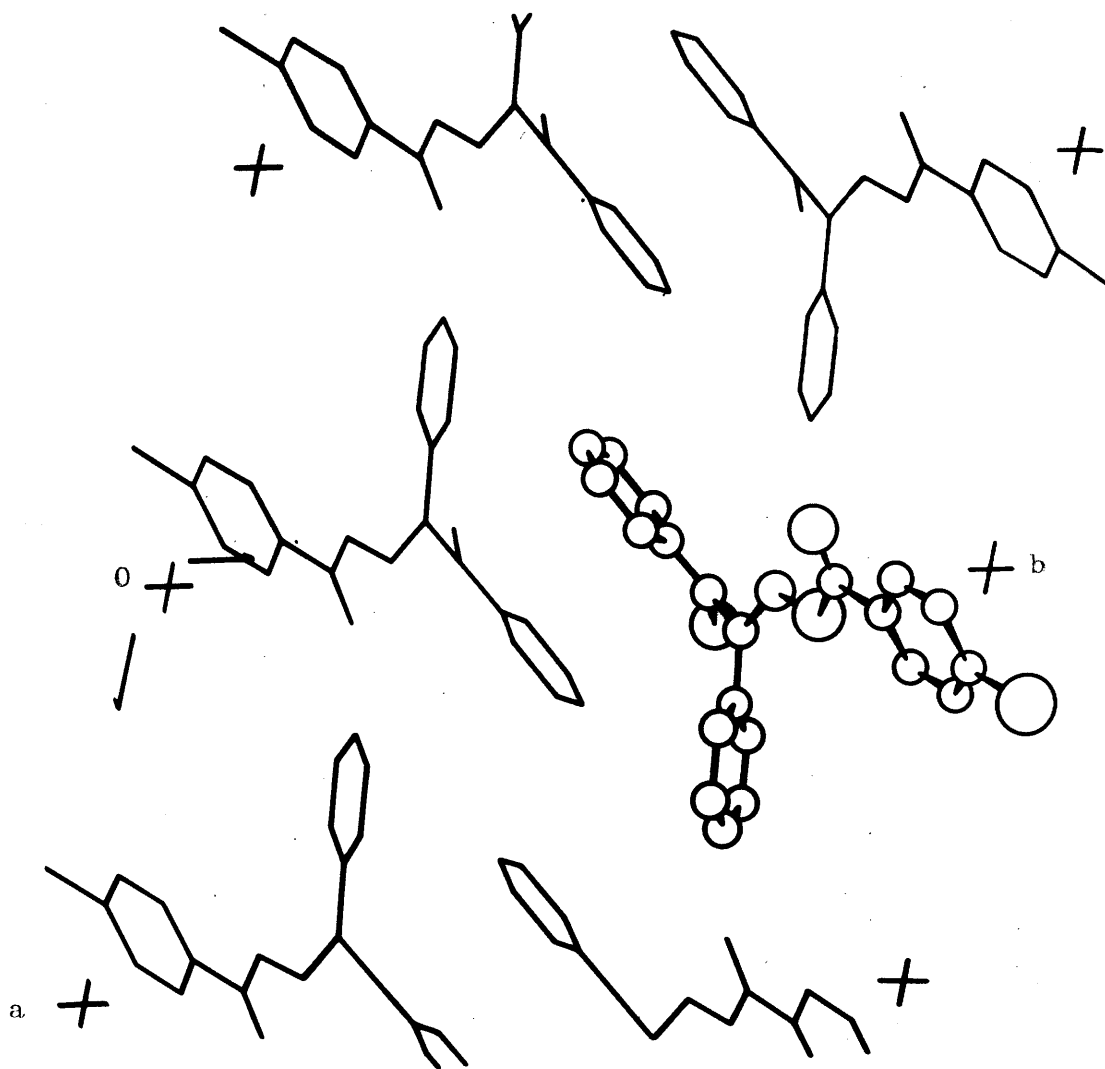
The only contact that is significantly shorter than the corresponding Van der Waals radius is that between O(24) and C(7). The calculated standard deviation of this measurement is just under 0.03 \AA , but the difference between the observed and calculated distances is large enough to be significant even if the standard deviation were as high as 0.04 \AA . Contacts as short as 2.77 \AA between sp_2 -hybridized carbon and the oxygen of a carbonyl group have been observed as well.

compounds where there is no possibility for hydrogen bonding. (Davies and Blum, 1955); Chn. Jeffrey and Sakurai, 1962; Bolton, 1963, 1964, 1965). Bolton has studied this effect and has suggested that it occurs when dipole-dipole or dipole-induced-dipole interactions are quite strong. The shortest contacts occur when the bond of the carbonyl group is approximately collinear with the π -bond of a carbon atom in an adjacent molecule. In the compounds studied by Bolton, the carbon atom is in an aromatic system with at least two carbonyl substituents. The average length of the O...C contact is 2.82 Å, and the average value of the C=O...C angle is 15f. In the present study, the carbon atom involved in the contact is an sp_2 -carbon in a carboxyl group. There is no large inductive effect present, but neither of the C=O groups in the molecule seem to be involved in any appreciable resonance interaction. It seems reasonable to expect a certain amount of dipole-dipole interaction between the two groups. This interaction would not be as strong as in alloxan and the other molecules studied by Bolton: the C=O...C angle (116°) is less favourable and the contact is longer than those he observed.

Kihara (1962) has shown that the structures of multipolar molecular crystals are governed by electrostatic forces between the molecules. He has determined correct crystal structures for a number of relatively simple molecules by building models with shape and magnetization appropriate

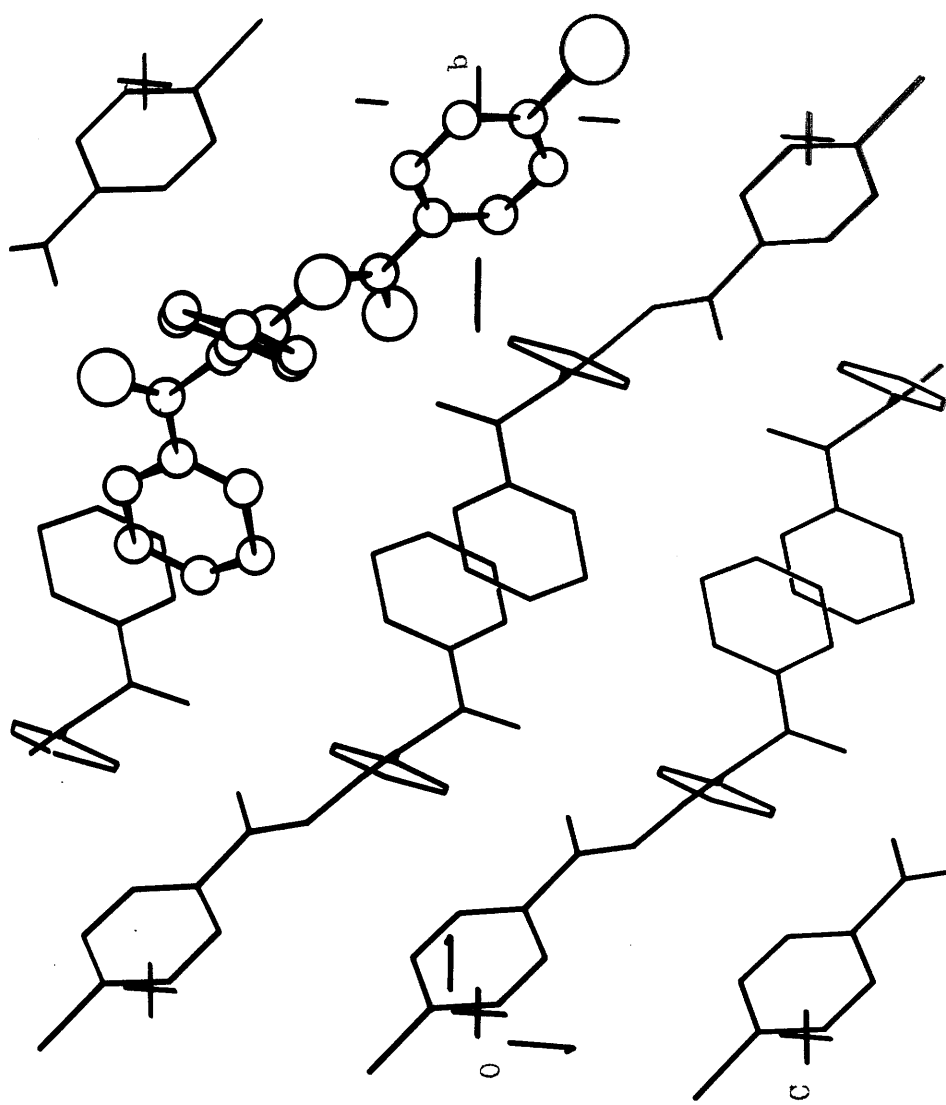
to the particular molecule and allowing the models to "crystallise". This would seem to indicate, that electrostatic forces play an important role in determining the crystal structures of a fairly broad category of molecules. This general observation, coupled with the fact that the carboxyl group in the present structure has been rotated out of the plane of the benzene ring in a direction that would make a more favourable angle between the π -orbital of the carbon atom and the σ -orbital of the oxygen, suggests that the close contact is real.

Figure 10



Packing diagram of the para-bromobenzoate of benzil monoxime, viewed along the c-axis

Figure 11



Molecular packing of the para-bromobenzate of benzylmagnesium bromide viewed in projection along the a-axis.

para-Diethylbenzene of Benzilmonoxime.

Final fractional atomic coordinates with their estimated standard deviations. No significance should be attached to the last figure as it has been included only to allow derived quantities to be calculated without loss of accuracy.

C(1)	0.2293 (35)	-0.0007 (16)	1.1903 (35)
C(2)	0.3015 (30)	-0.0077 (16)	0.9457 (40)
C(3)	0.2333 (29)	-0.0692 (17)	0.7899 (42)
C(4)	0.0974 (27)	-0.1240 (14)	0.8362 (37)
C(5)	0.0197 (31)	-0.1105 (17)	1.0341 (41)
C(6)	0.0534 (31)	-0.0550 (18)	1.2051 (29)
C(7)	0.0109 (26)	-0.1892 (14)	0.6821 (31)
C(8)	0.1373 (25)	-0.2880 (13)	0.2282 (32)
C(9)	0.0604 (27)	-0.3420 (14)	0.0462 (35)
C(10)	-0.0801 (27)	-0.4029 (13)	0.0837 (37)
C(11)	-0.1029 (29)	-0.4368 (12)	0.2827 (40)
C(12)	-0.2271 (30)	-0.5044 (16)	0.3188 (50)
C(13)	-0.3294 (31)	-0.5339 (18)	0.1435 (50)
C(14)	-0.3040 (30)	-0.5019 (20)	-0.0518 (51)
C(15)	-0.1702 (29)	-0.4330 (17)	-0.0857 (51)
C(16)	0.3110 (25)	-0.2745 (14)	0.2499 (36)
C(17)	0.4064 (24)	-0.2459 (14)	0.0701 (37)
C(18)	0.5563 (33)	-0.2456 (16)	0.0935 (40)
C(19)	0.6244 (28)	-0.2642 (16)	0.2791 (36)
C(20)	0.5243 (26)	-0.2907 (15)	0.4507 (43)
C(21)	0.3674 (26)	-0.2909 (15)	0.4399 (40)
N(22)	0.0373 (21)	-0.2576 (11)	0.3570 (28)
O(23)	0.1123 (18)	-0.3256 (10)	-0.1263 (25)
O(24)	0.1148 (16)	-0.2029 (9)	0.5287 (23)
O(25)	-0.1135 (19)	-0.2309 (10)	0.7229 (28)

Br(26) 0.31968(41)

0.08090(18)

1.3570(49)

Table 9

Para-bromobenzoate of Benzilmonoxime: Anisotropic Temperature Parameters

	U_{11}	U_{22}	U_{33}	$2U_{23}$	$2U_{31}$	$2U_{12}$
C(1)	0.156	0.073	0.063	-0.064	0.035	0.085
C(2)	0.079	0.086	0.116	-0.010	-0.009	0.028
C(3)	0.060	0.103	0.139	0.010	-0.004	0.009
C(4)	0.074	0.079	0.082	0.006	0.030	0.045
C(5)	0.095	0.110	0.087	0.014	0.024	0.069
C(6)	0.115	0.063	0.082	-0.007	0.025	-0.007
C(7)	0.072	0.087	0.079	-0.039	0.038	0.023
C(8)	0.067	0.075	0.064	-0.007	0.022	0.029
C(9)	0.087	0.083	0.052	0.041	0.040	-0.004
C(10)	0.083	0.074	0.076	0.011	0.078	0.039
C(11)	0.088	0.046	0.114	0.005	0.041	0.022
C(12)	0.120	0.077	0.126	0.001	0.022	-0.022
C(13)	0.090	0.089	0.153	-0.008	0.013	0.022
C(14)	0.108	0.123	0.120	0.002	0.012	0.050
C(15)	0.053	0.138	0.077	-0.062	-0.023	-0.005
C(16)	0.054	0.080	0.086	-0.020	-0.020	0.016
C(17)	0.047	0.088	0.084	0.019	-0.001	-0.044
C(18)	0.105	0.092	0.076	0.020	-0.016	-0.036
C(19)	0.083	0.097	0.069	-0.018	-0.010	0.001
C(20)	0.050	0.089	0.116	-0.026	-0.002	-0.003
C(21)	0.061	0.090	0.097	-0.020	0.026	0.012
N(22)	0.074	0.071	0.070	0.022	0.010	0.005
O(23)	0.080	0.110	0.088	0.015	0.009	-0.003
O(24)	0.059	0.093	0.083	-0.042	0.010	-0.004
O(25)	0.090	0.098	0.110	-0.020	0.065	-0.004
Br(26)	0.136	0.096	0.102	-0.060	-0.011	0.014.

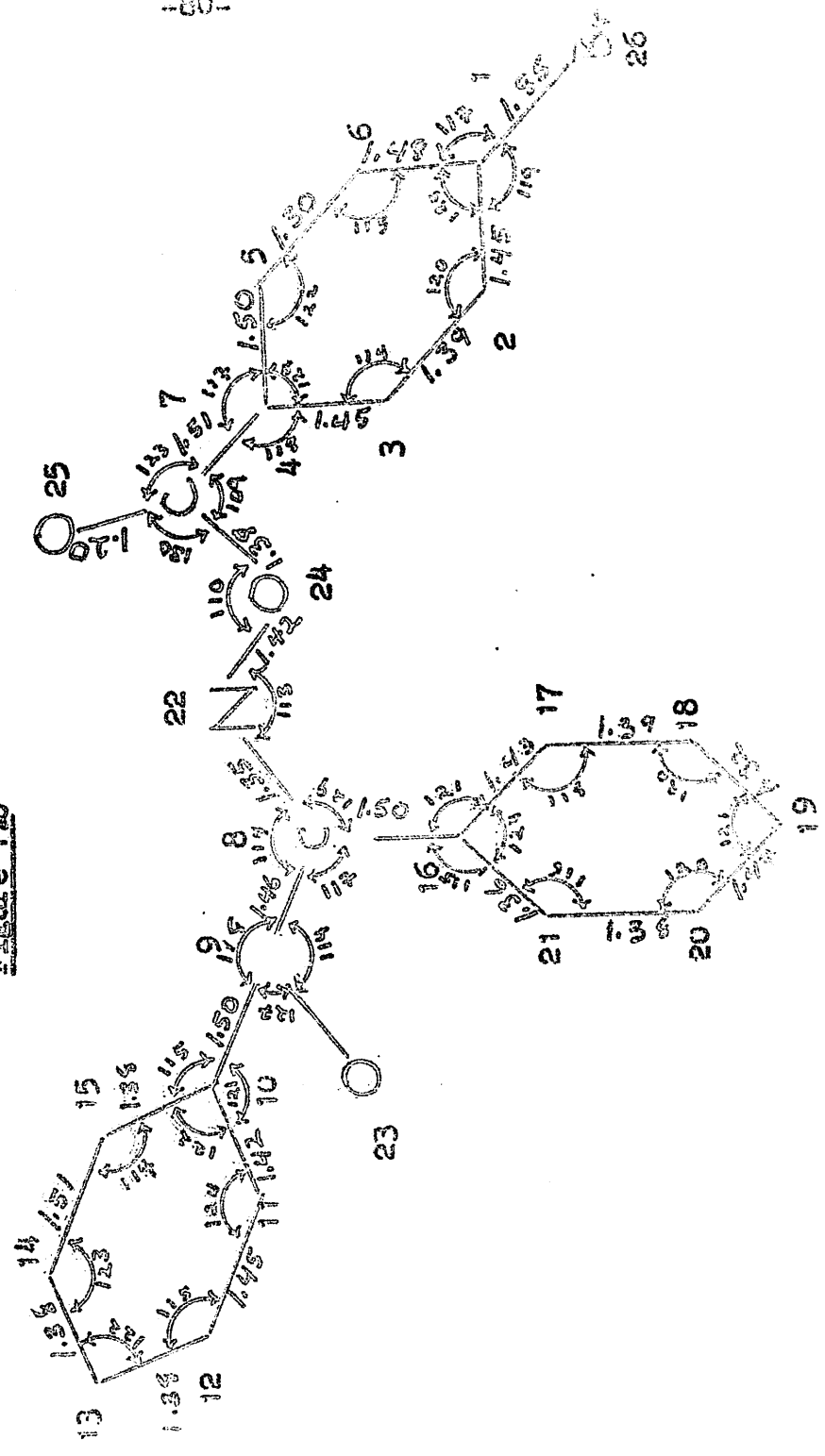
Table 10

Para-bromobenzoate of Benzilmonoxime:

Bond lengths in angstroms, with standard deviations given as units in the last place.

Br(26)	C(1)	1.845(25)
C(1)	C(2)	1.448(37)
C(2)	C(3)	1.390(39)
C(3)	C(4)	1.446(35)
C(4)	C(5)	1.449(35)
C(5)	C(6)	1.301(36)
C(6)	C(1)	1.477(39)
C(4)	C(7)	1.507(33)
C(7)	O(25)	1.197(28)
C(7)	O(24)	1.377(27)
O(24)	N(22)	1.417(23)
N(22)	C(8)	1.354(28)
C(8)	C(16)	1.495(30)
C(16)	C(17)	1.433(32)
C(17)	C(18)	1.387(35)
C(18)	C(19)	1.316(35)
C(19)	C(20)	1.475(35)
C(20)	C(21)	1.377(32)
C(21)	C(16)	1.357(34)
C(8)	C(9)	1.456(30)
C(9)	O(23)	1.217(26)
C(9)	C(10)	1.469(32)
C(10)	C(11)	1.420(34)
C(11)	C(12)	1.447(38)
C(12)	C(13)	1.375(44)
C(13)	C(14)	1.380(46)
C(14)	C(15)	1.511(41)
C(15)	C(10)	1.323(33)

Figure 12



para-Bromobenzoate of Benzilmonoxime: Bond Lengths and Angles.

Table II

Poly-Bromoterephthalate of Hexamethylene:

Bond angles with their estimated standard deviations given in units in the last place.

			ANGLE	E. S. D.
Br(26)	C(1)	C(2)	119.5	2.0
C(1)	C(2)	C(3)	119.7	2.4
C(2)	C(3)	C(4)	115.3	2.5
C(3)	C(4)	C(5)	124.5	2.3
C(4)	C(5)	C(6)	117.7	1.1
C(5)	C(6)	Br(26)	116.9	2.0
C(6)	Br(26)	C(1)	121.8	2.5
Br(26)	C(1)	C(2)	115.4	2.4
C(1)	C(2)	C(3)	116.7	2.2
C(2)	C(3)	C(4)	123.7	2.2
C(3)	C(4)	C(5)	122.9	2.1
C(4)	C(5)	C(6)	108.7	1.8
C(5)	C(6)	Br(26)	129.9	2.5
C(6)	Br(26)	C(1)	110.0	2.5
Br(26)	C(1)	C(2)	113.7	2.1
C(1)	C(2)	C(3)	129.3	2.0
C(2)	C(3)	C(4)	115.0	2.0
C(3)	C(4)	C(5)	120.7	2.0
C(4)	C(5)	C(6)	117.6	1.1
C(5)	C(6)	Br(26)	120.2	2.4
C(6)	Br(26)	C(1)	120.8	2.4
Br(26)	C(1)	C(2)	120.5	2.3
C(1)	C(2)	C(3)	115.8	2.3
C(2)	C(3)	C(4)	124.3	2.0
C(3)	C(4)	C(5)	117.0	2.0
C(4)	C(5)	C(6)	114.5	2.0
C(5)	C(6)	Br(26)	126.7	2.0
C(6)	Br(26)	C(1)	118.3	2.0
Br(26)	C(1)	C(2)	121.3	2.0
C(1)	C(2)	C(3)	123.6	2.0
C(2)	C(3)	C(4)	114.0	2.0
C(3)	C(4)	C(5)	121.8	2.0
C(4)	C(5)	C(6)	123.1	2.0
C(5)	C(6)	Br(26)	114.3	2.0
C(6)	Br(26)	C(1)	122.3	2.0

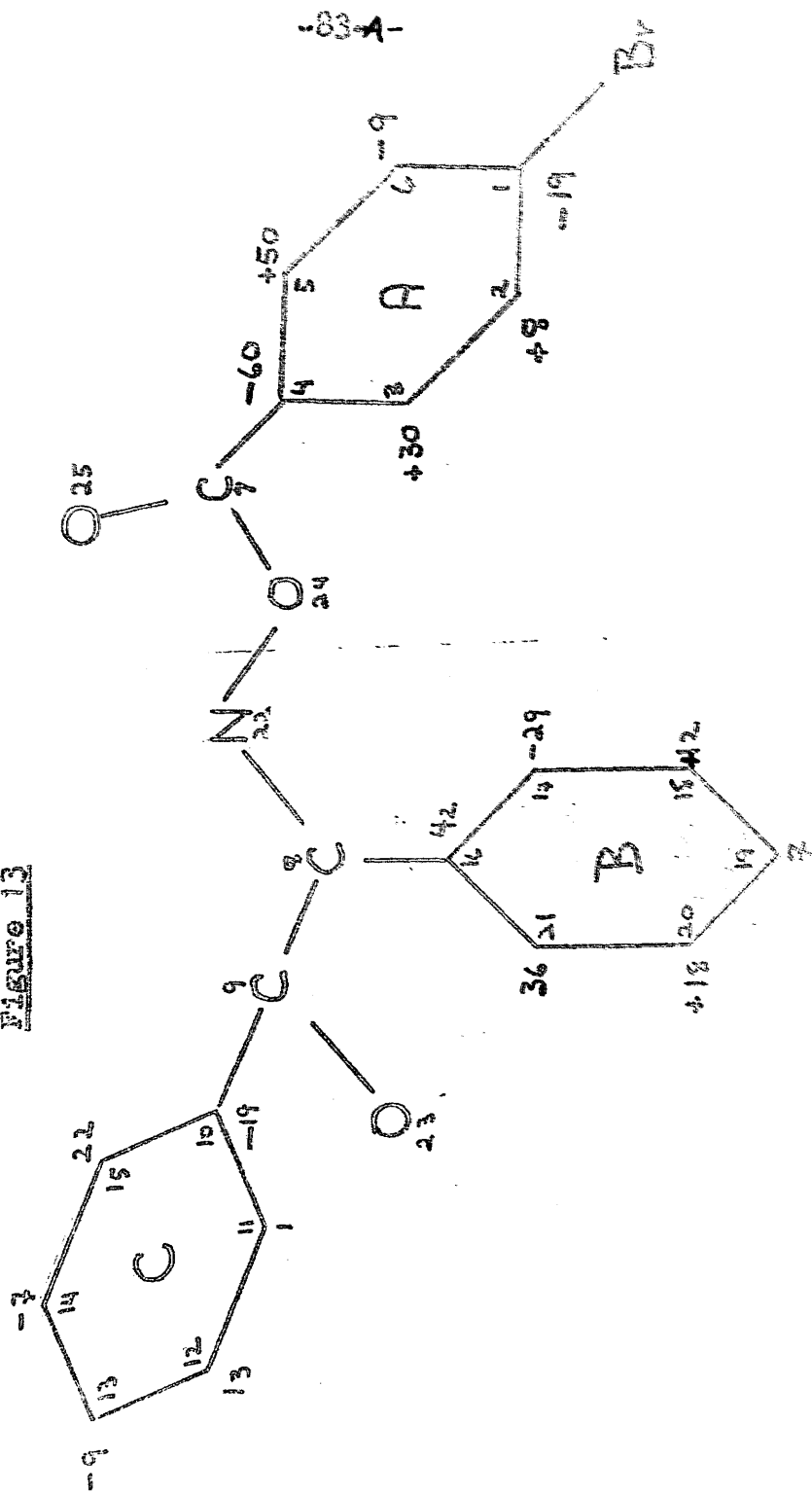
Table 12

Para-Bromobenzoate of Benzilmonoxime:

Correlation Coefficients

C(1)	0.29998	0.14586	-0.16655
C(2)	0.11987	0.06582	0.01953
C(3)	0.08878	0.04642	0.07248
C(4)	0.18409	0.17467	0.07385
C(5)	0.25684	0.12183	0.09247
C(6)	0.00550	0.12405	0.05836
C(7)	0.13869	0.18289	-0.07714
C(8)	0.16573	0.12154	0.03710
C(9)	0.01546	0.16854	0.21003
C(10)	0.18099	0.30766	0.08628
C(11)	0.11643	0.18964	0.06149
C(12)	-0.02487	0.09343	0.09301
C(13)	0.12420	0.09403	0.05286
C(14)	0.17601	0.07558	0.06221
C(15)	0.03727	-0.03303	-0.13106
C(16)	0.09594	-0.00379	0.00554
C(17)	-0.10309	0.05197	0.13689
C(18)	-0.09101	-0.02412	0.15814
C(19)	0.05061	-0.01672	0.01787
C(20)	0.05333	0.07842	-0.03691
C(21)	0.07686	0.13350	-0.01227
N(22)	0.07201	0.09923	0.13674
O(23)	0.05114	0.08290	0.11462
O(24)	0.04473	0.08099	-0.06906
O(25)	0.04323	0.24747	0.02315
Br(26)	0.04472	0.00410	-0.16339

Figure 13



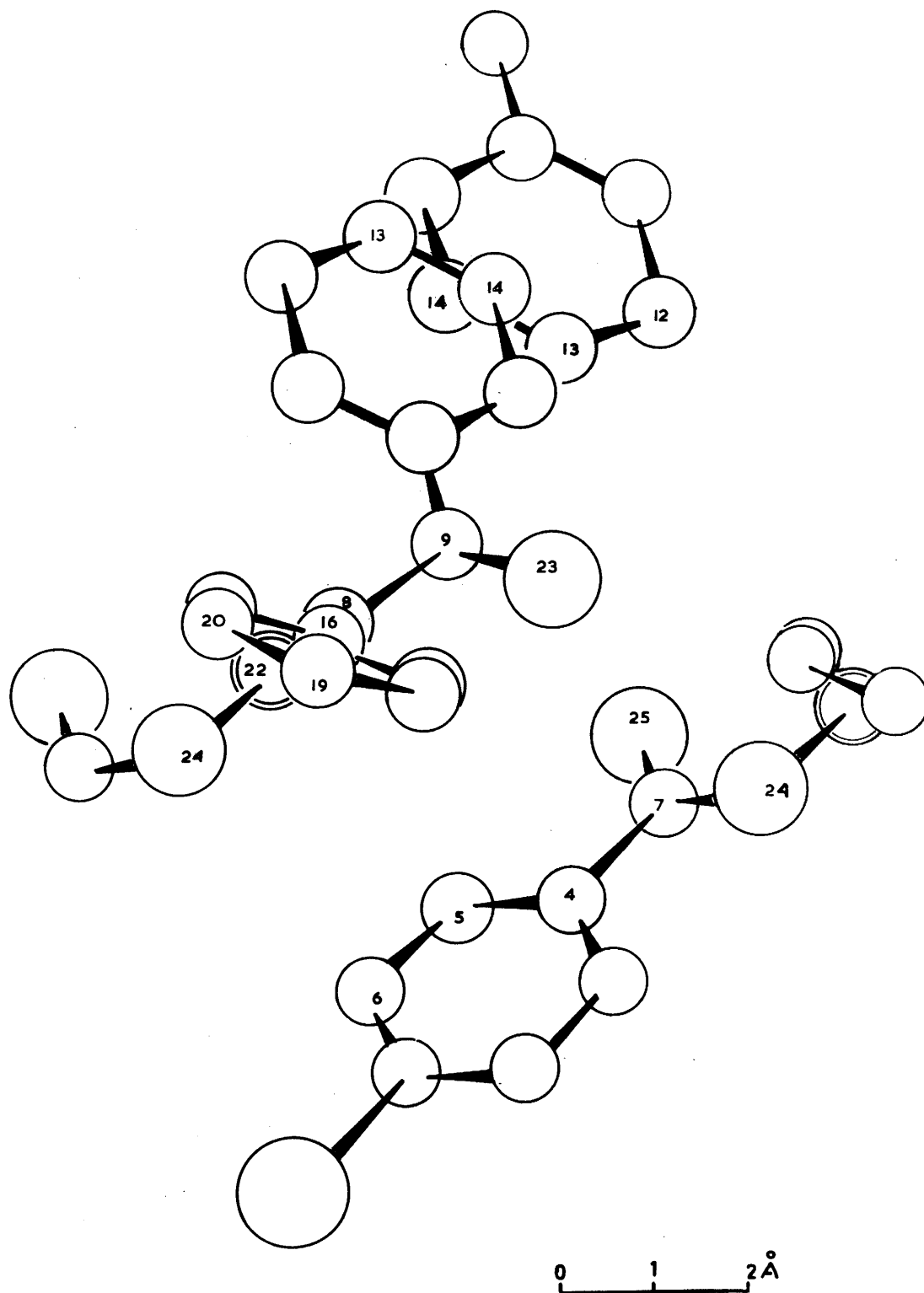
p-bromobenzoyl bromide Deviations from the benzene planes through the benzene rings are given in units of \AA $\times 10^{-8}$

Table 14

Para-Bromobenzoate of BenzilprooximeInter-molecular Contacts < 3.60 A.

Equivalent position 1 refers to the molecule whose coordinates are listed in Table 5; position 2 refers to the molecule related to this one by a centre of inversion.

C(9)	O(25)	1	(0, 0, -1)	3.42
C(13)	C(14)	2	(-1, -1, 0)	3.48
C(14)	C(14)	2	(-1, -1, 0)	3.53
C(19)	O(25)	1	(1, 0, 0)	3.53
C(20)	O(25)	1	(1, 0, 0)	3.49
H(22)	C(5)	1	(0, 0, -1)	3.30
O(23)	C(4)	1	(0, 0, -1)	3.47
	C(7)	1	(0, 0, -1)	3.49
	C(12)	2	(0, -1, 0)	3.43
	C(13)	2	(0, -1, 0)	3.52
	C(21)	1	(0, 0, -1)	3.56
	O(24)	1	(0, 0, -1)	3.17
O(25)	1	(0, 0, -1)	3.08	
O(24)	O(6)	1	(0, 0, -1)	3.42



Inter-molecular contacts (median axis projection)

Table 15 Ctd.

H	K	L	Fe	Po	H	K	L	Fe	Po	H	K	L	Fe	Po	H	K	L	Fe	Po	H	K	L	Fe	Po
4	5	1	27.2	30.6	3	9	2	25.4	-23.0	0	0	2	34.0	27.9	5	10	2	14.1	-18.3	0	0	2	14.1	-18.3
4	4	1	16.5	-15.3	3	8	2	14.9	-13.8	0	0	2	27.1	22.3	5	11	2	7.2	-7.3	0	0	2	7.2	-7.3
4	4	3	36.9	-37.4	3	7	2	12.4	-13.6	0	0	2	7.4	-6.4	6	1	2	9.8	-9.6	0	0	2	9.8	-9.6
4	4	2	12.7	-12.1	3	6	2	29.3	29.5	0	0	2	2.9	-4.5	6	2	2	5.5	-4.2	0	0	2	5.5	-4.2
4	4	1	20.3	-19.0	3	5	2	20.3	20.3	0	0	2	30.8	-31.6	6	3	2	7.6	-5.7	0	0	2	7.6	-5.7
4	4	0	20.6	-23.4	3	4	2	32.2	40.9	0	0	2	28.9	-31.0	6	4	2	12.2	-13.1	0	0	2	12.2	-13.1
4	4	0	19.5	17.2	3	3	2	8.5	7.9	0	0	2	45.4	-54.7	6	5	2	7.8	8.9	0	0	2	7.8	8.9
4	4	3	18.1	17.9	3	2	2	10.9	-9.0	0	0	2	52.8	52.6	6	6	2	8.9	8.5	0	0	2	8.9	8.5
4	4	4	10.4	13.0	3	1	2	9.9	11.7	0	0	2	34.7	36.2	6	7	2	10.5	10.5	0	0	2	10.5	10.5
4	4	5	22.2	26.4	3	0	2	6.4	-8.2	0	0	2	17.5	15.2	6	8	2	6.0	-6.9	0	0	2	6.0	-6.9
4	4	6	16.5	17.9	3	3	3	35.5	-42.2	0	0	2	31.6	34.8	6	9	2	18.0	-15.2	0	0	2	18.0	-15.2
4	4	7	8.5	10.0	3	4	3	29.2	-29.9	0	0	2	55.7	51.6	6	10	2	12.6	-14.8	0	0	2	12.6	-14.8
4	4	8	15.7	-18.6	3	5	3	17.0	20.0	0	0	2	18.7	-16.7	6	11	2	15.9	-14.6	0	0	2	15.9	-14.6
4	4	9	6.4	5.8	3	6	3	5.5	4.2	0	0	2	22.8	-21.2	6	12	2	18.1	28.1	0	0	2	18.1	28.1
4	4	10	5.9	4.6	3	7	3	4.5	3.4	0	0	2	17.1	-17.9	6	13	2	8.7	-10.1	0	0	2	8.7	-10.1
4	4	11	6.7	-7.0	3	8	3	11.5	-3.7	0	0	2	12.9	-14.2	6	14	2	16.4	-15.0	0	0	2	16.4	-15.0
4	4	12	7.4	-10.4	4	1	3	14.5	-11.0	0	0	2	17.3	-16.4	6	15	2	32.9	40.3	0	0	2	32.9	40.3
4	4	13	7.1	-9.4	4	2	3	13.0	-8.5	0	0	2	10.0	-9.5	6	16	2	21.7	25.0	0	0	2	21.7	25.0
4	4	14	17.5	-22.9	4	3	3	4.6	-8.0	0	0	2	4.6	4.6	6	17	2	9.4	8.0	0	0	2	9.4	8.0
4	4	15	5.3	-6.8	4	4	3	11.4	-9.3	0	0	2	4.2	3.6	6	18	2	6.4	6.0	0	0	2	6.4	6.0
4	4	16	13.1	-14.0	4	5	3	7.8	-5.4	0	0	2	9.0	8.0	6	19	2	12.8	11.9	0	0	2	12.8	11.9
4	4	17	6.5	-5.8	4	6	3	6.7	-7.8	0	0	2	24.6	29.1	6	20	2	13.4	11.5	0	0	2	13.4	11.5
4	4	18	10.3	-9.3	4	7	3	9.1	-16.3	0	0	2	44.2	39.2	6	21	2	16.2	-15.7	0	0	2	16.2	-15.7
4	4	19	7.3	7.1	4	8	3	17.0	18.3	0	0	2	9.6	12.0	6	22	2	15.7	-17.4	0	0	2	15.7	-17.4
4	4	20	21.4	21.0	4	9	3	4.0	4.0	0	0	2	9.7	-6.0	6	23	2	51.1	-47.4	0	0	2	51.1	-47.4
4	4	21	28.0	28.5	4	10	3	11.1	-9.5	0	0	2	14.5	-19.0	6	24	2	51.0	-46.3	0	0	2	51.0	-46.3
4	4	22	3.9	-7.7	4	11	3	21.0	-21.8	0	0	2	25.0	-21.5	6	25	2	15.0	-14.4	0	0	2	15.0	-14.4
4	4	23	4.7	2.9	4	12	3	20.9	-21.7	0	0	2	19.1	-15.5	6	26	2	8.2	-5.9	0	0	2	8.2	-5.9
4	4	24	9.6	-9.2	4	13	3	26.5	-31.2	0	0	2	13.5	12.2	6	27	2	23.7	22.0	0	0	2	23.7	22.0
4	4	25	18.1	-23.1	4	14	3	9.2	-8.0	0	0	2	25.7	24.7	6	28	2	19.2	16.9	0	0	2	19.2	16.9
4	4	26	10.2	-13.9	4	15	3	4.7	-7.0	0	0	2	13.0	14.9	6	29	2	27.3	23.9	0	0	2	27.3	23.9
4	4	27	13.2	-16.2	4	16	3	6.2	4.1	0	0	2	9.5	8.9	6	30	2	20.5	19.3	0	0	2	20.5	19.3
4	4	28	6.6	6.5	4	17	3	9.1	8.5	0	0	2	32.5	-27.7	6	31	2	24.4	-20.3	0	0	2	24.4	-20.3
4	4	29	10.5	9.6	4	18	3	14.3	16.5	0	0	2	57.3	-45.3	6	32	2	13.5	12.2	0	0	2	13.5	12.2
4	4	30	10.0	11.3	4	19	3	5.9	3.4	0	0	2	18.7	-16.9	6	33	2	7.1	-11.0	0	0	2	7.1	-11.0
4	4	31	6.0	-5.3	4	20	3	8.1	-8.7	0	0	2	25.9	25.0	6	34	2	10.2	-10.4	0	0	2	10.2	-10.4
4	4	32	8.6	-8.2	4	21	3	7.7	-7.4	0	0	2	31.0	25.8	6	35	2	26.2	22.9	0	0	2	26.2	22.9
4	4	33	18.2	-20.5	4	22	3	7.7	-5.9	0	0	2	18.2	18.3	6	36	2	30.4	29.7	0	0	2	30.4	29.7
4	4	34	8.3	-6.4	4	23	3	5.0	-3.0	0	0	2	11.9	12.8	6	37	2	26.6	24.5	0	0	2	26.6	24.5
4	4	35	9.1	-9.4	4	24	3	3.8	2.4	0	0	2	14.1	16.0	6	38	2	8.5	-3.6	0	0	2	8.5	-3.6
4	4	36	8.3	-7.4	4	25	3	3.7	-2.2	0	0	2	8.3	-9.9	6	39	2	12.7	-13.0	0	0	2	12.7	-13.0
4	4	37	10.5	-12.5	4	26	3	2.5	-6.6	0	0	2	11.7	-12.6	6	40	2	6.5	-6.9	0	0	2	6.5	-6.9
4	4	38	14.7	-13.3	4	27	3	7.6	3.2	0	0	2	11.7	12.5	6	41	2	12.5	-11.0	0	0	2	12.5	-11.0
4	4	39	11.0	-8.7	4	28	3	3.6	3.2	0	0	2	28.7	25.5	6	42	2	14.5	-17.2	0	0	2	14.5	-17.2
4	4	40	8.3	8.0	4	29	3	12.8	9.8	0	0	2	34.1	33.5	6	43	2	9.2	-10.8	0	0	2	9.2	-10.8
4	4	41	13.3	10.9	4	30	3	7.9	5.9	0	0	2	25.0	20.8	6	44	2	40.3	42.2	0	0	2	40.3	42.2
4	4	42	46.4	44.7	4	31	3	14.0	12.3	0	0	2	13.0	12.9	6	45	2	21.9	21.8	0	0	2	21.9	21.8
4	4	43	33.8	35.7	4	32	3	12.8	-13.0	0	0	2	19.8	20.7	6	46	2	10.5	-12.9	0	0	2	10.5	-12.9
4	4	44	34.7	33.9	4	33	3	25.3	-23.5	0	0	2	3.2	4.4	6	47	2	19.0	-18.9	0	0	2	19.0	-18.9
4	4	45	17.2	19.2	4	34	3	45.4	-13.5	0	0	2	19.4	14.4	6	48	2	21.1	-18.3	0	0	2	21.1	-18.3
4	4	46	8.5	-15.9	4	35	3	8.4	-6.2	0	0	2	19.8	-20.3	6	49	2	20.8	-17.3	0	0	2	20.8	-17.3
4	4	47	17.2	-15.0	4	36	3	3.7	3.7	0	0	2	39.7	37.1	6	50	2	7.1	-10.1	0	0	2	7.1	-10.1
4	4	48	15.0	-23.0	4	37	3	13.4	13.6	0	0	2	17.8	17.0	6	51	2	5.8	-2.2	0	0	2	5.8	-2.2
4	4	49	50.8	-61.3	4	38	3	13.5	12.6	0	0	2	7.2	7.2	6	52	2	9.2	11.4	0	0	2	9.2	11.4
4	4	50	27.4	-54.4	4	39	3	4.3	4.2	0	0	2	17.1	17.9	6	53	2	7.1	9.3	0	0	2	7.1	9.3
4	4	51	12.7	-15.3	4	40	3	14.4	5.8	0	0	2	29.6	-29.0	6	54	2	5.8	4.2	0	0	2	5.8	4.2
4	4	52	16.6	21.5	4	41	3	14.4	16.1	0	0	2	27.5	-29.0	6	55	2	10.0	-6.1	0	0	2	10.0	-6.1
4	4	53	11.8	-11.7	4	42	3	11.9	-12.6	0	0	2	7.8	-8.0	6	56	2	14.3	-17.7	0	0	2	14.3	-17.7
4	4	54	11.5	9.5	4	43	3	6.9	-4.4	0	0	2	8.0	1.1	6	57	2	14.3	-16.0	0	0	2	14.3	-16.0
4	4	55	48.8	43.1	4	44	3	5.6	-5.8	0	0	2	6.9	-6.1	6	58	2	5.3	-3.1	0	0	2	5.3	-3.1
4	4	56	33.8	31.9	4	45	3	4.6	-5.3	0	0	2	23.7	-21.2	6	59	2	11.7	9.8	0	0	2	11.7	9.8
4	4	57	5.3	-5.2	4	46	3	6.5	-5.6	0	0	2	13.4	-12.3	6	60	2	7.4	6.6	0	0	2	7.4	6.6
4	4	58	5.0	3.9	4	47	3	4.9	4.2	0	0	2	6.2	-6.8	6	61	2	5.7	5.4	0	0	2	5.7	5.4
4	4	59	21.2	-17.2	4	48	3	4.8	5.2	0	0	2	9.0	-8.9	6	62	2	11.4	13.4	0	0	2	11.4	13.4
4	4	60	6.2	-2.4	4	49	3	4.5	3.5	0	0	2	14.6	-16.4	6	63	2	11.4	11.4	0	0	2	11.4	11.4
4	4	61	7.7	-5.3	4	50	3	4.6	-4.3	0	0	2	11.7	13.7	6	64	2	6.0	-5.7	0	0	2	6.0	-5.7

Table 15 Ctd.

H	K	L	Fe	Fe	H	K	L	Fe	Fe	H	K	L	Fe	Fe	H	K	L	Fe	Fe
22.0	-1	-4	-5	5.2	5	-2	-1	13.1	14.8	3	-7	-4	5.5	6.1	4	3	4	13.1	14.8
11.5	-4	-5	-7	5.4	5	-2	-1	6.0	7.0	3	-7	-4	14.2	14.9	4	4	4	6.0	7.0
9.2	-5	-6	-7	5.4	5	-2	-1	10.5	10.8	5	-11	-4	16.6	16.0	4	4	4	10.5	10.8
11.3	-6	-7	-8	5.7	5	-2	-1	8.4	8.4	5	-12	-4	4.2	-3.4	4	4	4	8.4	8.4
8.9	-7	-8	-9	8.0	5	-2	-1	6.1	6.1	5	-13	-4	5.2	-5.9	4	4	4	6.1	6.1
9.0	-8	-9	-10	7.3	5	-2	-1	6.1	6.1	5	-14	-4	6.5	-9.0	4	4	4	6.1	6.1
9.2	-9	-10	-11	7.9	5	-2	-1	6.1	6.1	5	-15	-4	9.3	-13.0	4	4	4	6.1	6.1
10.1	-10	-11	-12	4.5	5	-2	-1	6.1	6.1	5	-16	-4	10.8	-15.2	4	4	4	6.1	6.1
7.7	-11	-12	-13	5.0	5	-2	-1	6.1	6.1	5	-17	-4	6.4	-8.7	4	4	4	6.1	6.1
10.1	-12	-13	-14	4.5	5	-2	-1	6.1	6.1	5	-18	-4	8.5	-10.4	4	4	4	6.1	6.1
10.1	-13	-14	-15	3.8	5	-2	-1	6.1	6.1	5	-19	-4	4.5	6.2	4	4	4	6.1	6.1
10.1	-14	-15	-16	17.6	5	-2	-1	6.1	6.1	5	-20	-4	10.0	8.3	4	4	4	6.1	6.1
10.1	-15	-16	-17	22.5	5	-2	-1	6.1	6.1	5	-21	-4	5.7	5.4	4	4	4	6.1	6.1
10.1	-16	-17	-18	26.4	5	-2	-1	6.1	6.1	5	-22	-4	6.7	-6.8	4	4	4	6.1	6.1
10.1	-17	-18	-19	14.4	5	-2	-1	6.1	6.1	5	-23	-4	4.2	-4.1	4	4	4	6.1	6.1
10.1	-18	-19	-20	10.6	5	-2	-1	6.1	6.1	5	-24	-4	4.2	-4.0	4	4	4	6.1	6.1
10.1	-19	-20	-21	28.3	5	-2	-1	6.1	6.1	5	-25	-4	6.0	-5.9	4	4	4	6.1	6.1
10.1	-20	-21	-22	28.1	5	-2	-1	6.1	6.1	5	-26	-4	5.3	-6.1	4	4	4	6.1	6.1
10.1	-21	-22	-23	34.7	5	-2	-1	6.1	6.1	5	-27	-4	4.2	-1.6	4	4	4	6.1	6.1
10.1	-22	-23	-24	9.2	5	-2	-1	6.1	6.1	5	-28	-4	6.6	-5.9	4	4	4	6.1	6.1
10.1	-23	-24	-25	30.6	5	-2	-1	6.1	6.1	5	-29	-4	7.5	-7.7	4	4	4	6.1	6.1
10.1	-24	-25	-26	16.4	5	-2	-1	6.1	6.1	5	-30	-4	6.0	6.0	4	4	4	6.1	6.1
10.1	-25	-26	-27	36.2	5	-2	-1	6.1	6.1	5	-31	-4	7.1	7.8	4	4	4	6.1	6.1
10.1	-26	-27	-28	12.7	5	-2	-1	6.1	6.1	5	-32	-4	9.6	8.2	4	4	4	6.1	6.1
10.1	-27	-28	-29	16.2	5	-2	-1	6.1	6.1	5	-33	-4	8.2	-8.9	4	4	4	6.1	6.1
10.1	-28	-29	-30	21.5	5	-2	-1	6.1	6.1	5	-34	-4	4.0	-4.4	4	4	4	6.1	6.1
10.1	-29	-30	-31	17.6	5	-2	-1	6.1	6.1	5	-35	-4	6.7	7.2	4	4	4	6.1	6.1
10.1	-30	-31	-32	13.6	5	-2	-1	6.1	6.1	5	-36	-4	8.7	8.2	4	4	4	6.1	6.1
10.1	-31	-32	-33	25.1	5	-2	-1	6.1	6.1	5	-37	-4	9.0	9.4	4	4	4	6.1	6.1
10.1	-32	-33	-34	15.9	5	-2	-1	6.1	6.1	5	-38	-4	6.8	-6.4	4	4	4	6.1	6.1
10.1	-33	-34	-35	7.1	5	-2	-1	6.1	6.1	5	-39	-4	7.8	-6.1	4	4	4	6.1	6.1
10.1	-34	-35	-36	12.7	5	-2	-1	6.1	6.1	5	-40	-4	8.0	-8.9	4	4	4	6.1	6.1
10.1	-35	-36	-37	7.7	5	-2	-1	6.1	6.1	5	-41	-4	8.0	5.5	4	4	4	6.1	6.1
10.1	-36	-37	-38	6.2	5	-2	-1	6.1	6.1	5	-42	-4	9.0	7.4	4	4	4	6.1	6.1
10.1	-37	-38	-39	12.5	5	-2	-1	6.1	6.1	5	-43	-4	9.7	-8.1	4	4	4	6.1	6.1
10.1	-38	-39	-40	28.1	5	-2	-1	6.1	6.1	5	-44	-4	8.3	-9.5	4	4	4	6.1	6.1
10.1	-39	-40	-41	20.9	5	-2	-1	6.1	6.1	5	-45	-4	17.5	-17.7	4	4	4	6.1	6.1
10.1	-40	-41	-42	30.6	5	-2	-1	6.1	6.1	5	-46	-4	13.1	-11.1	4	4	4	6.1	6.1
10.1	-41	-42	-43	6.9	5	-2	-1	6.1	6.1	5	-47	-4	4.4	5.3	4	4	4	6.1	6.1
10.1	-42	-43	-44	18.5	5	-2	-1	6.1	6.1	5	-48	-4	8.5	6.3	4	4	4	6.1	6.1
10.1	-43	-44	-45	6.5	5	-2	-1	6.1	6.1	5	-49	-4	20.1	-4.1	4	4	4	6.1	6.1
10.1	-44	-45	-46	27.5	5	-2	-1	6.1	6.1	5	-50	-4	9.5	8.1	4	4	4	6.1	6.1
10.1	-45	-46	-47	35.3	5	-2	-1	6.1	6.1	5	-51	-4	13.8	12.4	4	4	4	6.1	6.1
10.1	-46	-47	-48	8.6	5	-2	-1	6.1	6.1	5	-52	-4	5.3	4.8	4	4	4	6.1	6.1
10.1	-47	-48	-49	13.9	5	-2	-1	6.1	6.1	5	-53	-4	4.3	2.5	4	4	4	6.1	6.1
10.1	-48	-49	-50	11.1	5	-2	-1	6.1	6.1	5	-54	-4	9.8	-10.3	4	4	4	6.1	6.1
10.1	-49	-50	-51	18.2	5	-2	-1	6.1	6.1	5	-55	-4	20.5	-21.4	4	4	4	6.1	6.1
10.1	-50	-51	-52	11.5	5	-2	-1	6.1	6.1	5	-56	-4	9.0	6.4	4	4	4	6.1	6.1
10.1	-51	-52	-53	15.6	5	-2	-1	6.1	6.1	5	-57	-4	6.4	5.3	4	4	4	6.1	6.1
10.1	-52	-53	-54	14.7	5	-2	-1	6.1	6.1	5	-58	-4	6.1	6.4	4	4	4	6.1	6.1
10.1	-53	-54	-55	5.6	5	-2	-1	6.1	6.1	5	-59	-4	6.1	6.1	4	4	4	6.1	6.1
10.1	-54	-55	-56	17.7	5	-2	-1	6.1	6.1	5	-60	-4	7.6	8.0	4	4	4	6.1	6.1
10.1	-55	-56	-57	10.0	5	-2	-1	6.1	6.1	5	-61	-4	7.7	8.4	4	4	4	6.1	6.1
10.1	-56	-57	-58	3.8	5	-2	-1	6.1	6.1	5	-62	-4	9.9	12.8	4	4	4	6.1	6.1
10.1	-57	-58	-59	6.2	5	-2	-1	6.1	6.1	5	-63	-4	11.3	11.7	4	4	4	6.1	6.1
10.1	-58	-59	-60	6.3	5	-2	-1	6.1	6.1	5	-64	-4	7.2	-8.9	4	4	4	6.1	6.1
10.1	-59	-60	-61	7.1	5	-2	-1	6.1	6.1	5	-65	-4	8.4	6.4	4	4	4	6.1	6.1
10.1	-60	-61	-62	10.1	5	-2	-1	6.1	6.1	5	-66	-4	10.2	-11.0	4	4	4	6.1	6.1
10.1	-61	-62	-63	12.8	5	-2	-1	6.1	6.1	5	-67	-4	7.1	-9.2	4	4	4	6.1	6.1
10.1	-62	-63	-64	5.8	5	-2	-1	6.1	6.1	5	-68	-4	4.7	6.1	4	4	4	6.1	6.1
10.1	-63	-64	-65	11.7	5	-2	-1	6.1	6.1	5	-69	-4	6.9	-6.6	4	4	4	6.1	6.1
10.1	-64	-65	-66	23.7	5	-2	-1	6.1	6.1	5	-70	-4	11.4	-16.6	4	4	4	6.1	6.1
10.1	-65	-66	-67	20.1	5	-2	-1	6.1	6.1	5	-71	-4	11.0	20.0	4	4	4	6.1	6.1
10.1	-66	-67	-68	22.2	5	-2	-1	6.1	6.1	5	-72	-4	9.4	11.6	4	4	4	6.1	6.1
10.1	-67	-68	-69	22.5	5	-2	-1	6.1	6.1	5	-73	-4	14.3	-5.4	4	4	4	6.1	6.1
10.1	-68	-69	-70	15.7	5	-2	-1	6.1	6.1	5	-74	-4	13.7	13.7	4	4	4	6.1	6.1
10.1	-69	-70	-71	13.9	5	-2	-1	6.1	6.1	5	-75	-4	8.5	-6.9	4	4	4	6.1	6.1
10.1	-70	-71	-72	11.0	5	-2	-1	6.1	6.1	5	-76	-4	7.1	-6.9	4	4	4	6.1	6.1
10.1	-71	-72	-73	5.5	5	-2	-1	6.1	6.1	5	-77	-4	8.1	-7.4	4	4	4	6.1	6.1
10.1	-72	-73	-74	5.1	5	-2	-1	6.1	6.1	5	-78	-4	10.3	-12.4	4	4	4	6.1	6.1
10.1	-73	-74	-75	13.0	5	-2	-1	6.1	6.1	5	-79	-4	7.0	-5.8	4	4	4	6.1	6.1
10.1	-74	-75	-76	7.2	5	-2	-1	6.1	6.1	5	-80	-4	6.1	17.0	4	4	4	6.1	6.1
10.1	-75	-76	-77	10.8	5	-2	-1	6.1	6.1	5	-81	-4	6.9	11.5	4	4	4	6.1	6.1
10.1	-76	-77	-78	4.5	5	-2	-1	6.1	6.1	5	-82	-4	6.8	5.7	4	4	4	6.1	6.1
10.1	-77	-78	-79	4.7	5	-2	-1	6.1	6.1	5	-83	-4	6.4	-4.5	4	4	4	6.1	6.1
10.1	-78	-79	-80	4.4	5	-2	-1	6.1	6.1	5	-84	-4	9.5	-10.6	4	4	4	6.1	6.1
10.1	-79	-80	-81	4.4	5	-2	-1	6.1	6.1	5	-85	-4	11.1	-12.2	4	4	4	6.1	6.1
10.1	-80	-81	-82	3.3	5	-2	-1	6.1	6.1	5	-86	-4	6.5	-2.6	4	4	4	6.1	6.1
10.1	-81	-82	-83	3.1	5	-2	-1	6.1	6.1	5	-87	-4	7.4	3.7	4	4	4	6.1	6.1
10.1	-82	-83	-84	3.6	5	-2	-1	6.1	6.1	5	-88	-4	7.5	7.9	4	4	4	6.1	6.1
10.1	-83	-84	-85	3.4	5	-2	-1	6.1	6.1	5	-89	-4	8.2	6.6	4	4	4	6.1	6.1
10.1	-84	-85	-86	5.4	5	-2	-1	6.1	6.1	5	-90	-4	7.5	-7.9	4	4	4	6.1	6.1
10.1	-85	-86	-87	7.1	5	-2	-1	6.1	6.1	5	-91	-4	8.2	-					

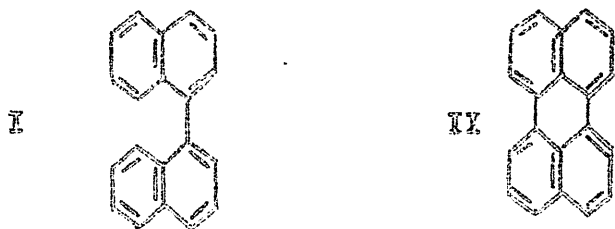
SECTION II

A Comparative Study of
1,1'-Binaphthyl and the
 β -form of Perylene

INTRODUCTION

Coulson has pointed out that a bond in a conjugated system may be regarded as a superposition of a σ - bond and a fractional π - bond. For a particular type of hybridization, the length of a bond can be expected to vary with change in π - bond order. Various empirical relationships have been suggested to correlate bond length with bond order. Since most of the accurate structure determinations have involved bonds of intermediate π - bond order, agreement between theory and experiment is very good in this region but becomes less reliable for extreme bond orders.

Binaphthyl, I, and perylene, II, are examples of molecules that contain bonds of very low π - bond order. Both of these molecules can be regarded as composed of two naphthalene residues connected by long bonds.



There are nine unexcited Kekulé structures for each molecule, representing the various combinations of three Kekulé structures of each naphthalene nucleus. None of these structures attributes double bond character to the linking bonds. Thus simple valence bond theory predicts that the long bonds are pure sp_2 - sp_2 single bonds with no π - bond character. Molecular orbital theory predicts

that the π -bond order will be low, but it cannot be zero since this theory assumes that there is always some interaction between π -bonds on adjacent atoms.

A study of these two compounds provides a fairly stringent test of the hypothesis that the aromatic character of the perylene molecule is localized in the naphthalene residues. Perylene is planar but binaphthyl is not. The angle between the planes of the naphthalenic residues is about 68° , sufficiently large to preclude any π -bonding interaction between the two ring systems. If there are significant differences between the dimensions of the two molecules, this could be interpreted as evidence for interaction between the naphthalenic residues in perylene.

THE CRYSTAL AND MOLECULAR STRUCTURE OF 1,1'-BINAPHTHYL

The crystal structure of 1,1'-binaphthyl (m.p. 148°C) was first established from projection data by Brown, Trotter and Robertson (1961). They have reported that the molecular configuration is cis with an angle of about 73° between the planes of the two ring systems, and have suggested that the variety of melting points reported in the literature corresponds to different crystalline forms. Brown has refined the structure in two projections by differential synthesis but his results have not been published.

Badar, Chua, Cook and Harris (1965) have shown that the compound has at least two distinct crystalline forms. The higher melting form (m.p. 157-159°C) was prepared by deamination of (+)-4,4'-naphthidine and is optically active. When melted and resolidified it is converted into the low melting form (m.p. 145°C). Infra-red spectra of the two forms are identical in carbon disulphide, in carbon tetrachloride and in cyclohexane, but there is an extra peak in the solid state spectrum of the low melting form. The authors attribute this to C-H out-of-plane vibrations caused by H...H and C...H interactions inherent in the cis configuration. They suggest that the higher melting form corresponds to the trans configuration of the molecule.

1.1. EXPERIMENTAL

Dr. Harris kindly provided samples of two crystalline modifications of 1,1'-binaphthyl. The high-melting form

does not show extinction when examined under a polarizing microscope. When a "single" crystal is photographed by the usual rotation method the result is a rather "spotty" powder diagram, typical of coarse, randomly-oriented crystallites. Recrystallisation from light petroleum did not improve the crystals.

The low-melting form crystallises as colourless plates that show sharp extinction and can be cut without apparent deformation. Weissenberg and precession photographs of one of these crystals mounted about its unique axis were used to check cell dimensions and space group. Cell dimensions were further confirmed by measuring reciprocal lattice spacings of high angle reflections on the linear diffractometer.

1.1.1. CRYSTAL DATA

Formula	$C_{20}H_{14}$	F.W.	254.3
System	monoclinic		
Lattice constants	$a = 20.97_g \overset{\circ}{\text{A}}$		
	$b = 6.35_1 \overset{\circ}{\text{A}}$		
	$c = 10.12_g \overset{\circ}{\text{A}}$		
	$\beta = 105.17^\circ$		
Absent Spectra	$hk\ell$ if $h+k$ odd		
	$h0\ell$ if ℓ odd		
	$0k0$ if k odd		
Space group	Cc or C2/c		
Volume	$1302.3 \overset{\circ}{\text{A}}^3$		

420

Density(obs.)	1.299 gm./cc
Molecules/cell	4
Density(calc.)	1.297 gm./cc
Linear absorption coefficient, $\mu = 0.86 \text{ cm}^{-1}$	
(Mo K α , $\lambda = 0.7107 \text{ \AA}$)	

1.1.2. REFLECTION DATA

Reflection data from eight reciprocal lattice nets, $h0^{\pm}$ to $h7^{\pm}$, were collected with Mo K α radiation on a Hilger-Watts linear diffractometer of the type described by Arndt and Phillips (1961). Peak height and background were measured at least twice for each reflection. Planes with intensities smaller than ten counts per minute were considered unobserved and were not included in the refinement. Lorentz-polarization corrections were applied to the observed data by computer program using the formula appropriate to equi-inclination Weissenberg geometry.

1.1.3. THE HILGER-WATTS DIFFRACTOMETER

The diffractometer is of the moving-crystal-stationary-counter type and can be described as a mechanical model of the reciprocal lattice. Its main components are three slides driven by a system of tapes, pulleys and gears. The motions of the crystal and the slides are coupled in such a way that any plane, hk^{\pm} , can be brought into the reflecting position by setting the coordinates of the corresponding reciprocal-lattice point, h_0^{\pm} , k_0^{\pm} , l_0^{\pm} , on

on the three slides. The oscillation motor then rotates the crystal through a small arc. During each oscillation, three counts are recorded. The first and third are background counts (time, t) recorded at the two extremes of the oscillation. The middle count (time, $2t$) is recorded while the crystal oscillates through a small, pre-set arc. A minimum of two oscillations, one with each of the SrO-ZrO_2 balanced filters, is required to measure peak and background for a single reflection. This type of counting mechanism should produce similar standard deviations for most measurements. However, spring and backlash in the mechanisms cause setting errors that increase with decrease in the cylindrical coordinate, ψ . This has two consequences: in terms of operation of the diffractometer, it means that reflections from planes with low $\sin \theta$ values must be measured by hand; in terms of quality of the data, it implies that absolute errors in the intensity data will show some systematic dependence on $\sin \theta$.

1.2. REFINEMENT OF THE STRUCTURE

Brown's (1961, unpublished) coordinates from differential synthesis refinement of two projections were used as starting point for the three-dimensional refinement of 1,1'-binaphthyl. The initial R-value was 36.2%, but this was reduced to 18.7% by three cycles of isotropic full-matrix least squares refinement with Cruickshank's (1961)

weighting scheme;

$$\sqrt{w} = (1/p_1 + p_0 + p_2 F^2 + p_3 F^3)^{-1/2}$$

The values of the parameters were:

$$p_1 = 4.20, p_2 = 0.011, p_3 = 0.$$

One further isotropic refinement cycle improved R to 14.6%.

The coordinates of the seven hydrogen atoms were calculated by hand. When these were included in the structure-factor calculation, the R-value was 15.3%. One further refinement cycle reduced R to 14.6%. At this point, the parameters of the weighting scheme were changed slightly:

$$p_1 = 2.56, p_2 = 0.011, p_3 = 0.$$

The refinement continued with isotropic temperature factors for hydrogen atoms and anisotropic temperature factors for carbon atoms. Because of the size of the matrix involved, the block-diagonal approximation was used. After five cycles, the refinement seemed to be near convergence: the R-value had fallen to 9.37%; indicated parameter shifts for carbon atoms were between 1/5 and 1/10 of the corresponding estimated standard deviations; and the weighting analysis showed that the average value of $\sum u_i^2$ was reasonably constant when analysed as a function of either $\sin \theta/\lambda$ or $|F_{obs}|$. However, some of the hydrogen atoms showed unlikely temperature factors and had moved to unacceptable locations.

Stewart, Davidson and Simpson (1965) have pointed out that if hydrogen atom form factors calculated from a simple 1s wave function are used in least-squares refinement, isotropic temperature factors will be abnormally low and the C-H bond distances will be short. The scattering factors usedⁱⁿ the present calculations are those tabulated by Hanson, Herman, Lea and Skillman (1964). Another possible source of difficulty in the refinement of the hydrogen parameters is the choice of weighting scheme. Cruickshank's (1961) weighting scheme was designed for film data and is probably not applicable in this case. The particular parameters used tend to give high relative weights to planes with small intensities.

The positions of the hydrogen atoms were recalculated by a computer program. Each atom was placed $1 \cdot 10 \text{ \AA}$ along the external bisector of the angle subtended at the corresponding carbon atom. Further refinement was attempted using these calculated positions with a scattering curve appropriate to bonded hydrogen (Stewart et al., 1965) and a new weighting scheme of the form,

$$\sqrt{w} = [1 - \exp(-2 \sin^2 \theta / \lambda^2)]^{\frac{1}{2}}$$

The anisotropic refinement of carbon parameters converged in two cycles to an R-value of 9.36%, but the weighting analysis was not satisfactory. A more successful weighting scheme was devised after examination of the

unweighted structure factors as a function of the magnitude of F_{obs} , the magnitude of $\sin \theta/\lambda$ and the index of the reciprocal lattice net, as shown in Table 1. The R-value is high for reflections with small values of $|F_{obs}|$. The increase in R with increase in layer index is caused by the large number of weak reflections observed at high angles. The absolute error in the structure factors is constant for most categories of reflections, but increases for large values of $|F_{obs}|$ and for low values of $\sin \theta/\lambda$. The weighting scheme adopted after this examination of the data gives unit weights to most reflections, but

$$\text{if } \sin^2 \theta / \lambda^2 < 0.16 \quad \text{then } w = (\sin^2 \theta / \lambda^2) / 0.16;$$

$$\text{if } |F_{obs}| > 20 \quad \text{then } w = 20 / |F_{obs}|.$$

Reflections with $|F_{calc}| < 1/3 |F_{obs}|$ were considered to be unreliably phased and were given a weight of zero. This effectively removes 45 of the 1315 reflections from the refinement. The value of $\sum w \Delta^2$ was 1545.86.

After two full-matrix calculations involving only the parameters of the carbon atoms, R was 9.30%. Positions and isotropic temperature factors of hydrogen atoms were allowed to refine for four block-diagonal cycles. Refinement was terminated at this point with R = 8.50% and $\sum w \Delta^2 = 1097.69$. Indicated shifts of carbon parameters were less than 1/10 of the corresponding estimated standard deviations; shifts of hydrogen parameters were between 1/3 and 1/5 of their e.s.d.

RESULTS

1,3'-Bisnaphthyl: Analysis of Observed and Calculated
Structure Amplitudes1. As function of $|F_{obs}|$

F_o	$\Sigma F_o $	$\Sigma F_c $	$\Sigma A $	N	R	\bar{R}
2	33.10	37.9	11.03	20	0.333	0.55
4	1194.63	1007.5	377.65	381	0.316	0.99
6	1150.58	1095.6	194.83	235	0.169	0.83
8	1278.04	1255.3	136.21	183	0.107	0.71
10	933.00	905.8	73.73	105	0.079	0.70
12	694.00	699.63	52.15	64	0.077	0.83
14	826.93	836.40	66.63	64	0.080	1.05
17	1004.84	956.69	54.95	65	0.053	0.81
24	1487.38	1493.25	81.81	74	0.653	1.30
33	1850.24	1800.14	102.14	66	0.056	1.30
67	2193.30	2162.35	78.63	50	0.036	1.57
110	648.36	666.89	23.71	8	0.037	2.06

As function of $\sin\theta/\lambda$

.20	845.15	819.49	50.72	27	0.060	1.03
.30	2190.96	2149.26	125.62	93	0.052	1.03
.40	2299.49	2270.33	204.60	184	0.033	1.03
.45	1705.99	1706.12	109.37	119	0.034	0.82
.50	1476.63	1520.34	131.63	149	0.032	0.77
.55	1241.24	1257.49	103.52	151	0.035	0.70
.60	1196.40	1157.24	144.68	184	0.121	0.75
.65	1061.34	1024.54	137.36	181	0.122	0.73
.70	646.91	581.60	109.53	114	0.169	0.93
.75	342.37	303.47	67.06	68	0.196	0.93
.80	146.09	98.13	53.70	34	0.368	1.51
1.00	41.87	29.47	15.36	11	0.367	1.40

As function of reciprocal lattice index

0	1642.53	1636.29	116.78	94	0.071	1.03
1	2697.20	2616.72	222.64	213	0.077	1.03
2	2683.17	2632.49	201.32	209	0.076	0.86
3	1792.01	1741.73	155.26	167	0.087	0.83
4	1674.48	1642.12	152.26	180	0.091	0.87
5	1142.88	1106.89	137.57	157	0.120	0.83
6	814.94	778.61	123.27	146	0.151	0.87
7	677.23	595.52	140.65	140	0.215	0.87

1.3. MOLECULAR DIMENSIONS AND DESCRIPTION OF THE STRUCTURE

Final atomic coordinates and temperature parameters are listed in Tables 2 and 3, respectively, with standard deviations given as units in the last place. Bond lengths and angles and their standard deviations are shown in Figure 1 and in Tables 4 and 5.

The molecular structure of 1,1'-binaphthyl viewed in projection along the g axis is shown in Figure 2. The planar naphthalene residues are linked by a bond across the diad axis, h . The length of this bond ($1.475 \pm 0.005 \text{ \AA}$) is very close to the value of 1.479 \AA predicted by Dewar and Schmeising (1959) and to the value of 1.477 \AA predicted by Pauling's method with a new order-length curve (Cruickshank and Sparks, 1960).

The equation of the best plane through the naphthalene nucleus is

$$0.0512X + 0.5596Y - 0.8272Z = -1.6754$$

where X, Y and Z are orthogonal axes defined in the following formulae:

$$X = gx + gzc \cos \beta;$$

$$Y = by;$$

$$Z = gzs \sin \beta;$$

Deviations from this plane are shown in Table 8. A χ^2 test with 7 degrees of freedom has a significance level of 0.10, indicating that the naphthalene ring systems

are essentially planar. If a mean plane is calculated through either of the benzene rings of the naphthalene system, three of the carbon atoms in the other ring show very large deviations from that plane. This does not imply that the naphthalene system is buckled; it just shows that the "best" planes through the individual rings are slightly different from the best plane through the whole system.

In Tables 6 and 7, some of the more important intra- and inter-molecular contacts are shown. Primed numbers refer to atoms whose positions are generated by the symmetry operation of the diad axis. It seems likely that the most important intra-molecular forces are carbon-carbon repulsions between C(2)...C(2)', C(8)...C(8)', and C(8)...C(9)'; whereas, the most important inter-molecular forces arise from the very large number of C...H interactions. Each naphthalene residue has 32 non-bonded C...H contacts shorter than $3.50 \overset{\circ}{\text{A}}$. The compromise between inter- and intra-molecular forces produces an angle of 68° between the planes of the two ring systems.

Badar et al. (1965) have interpreted the solid state (nujol mull) infra-red spectrum of the low melting form of 1,1'-binaphthyl in terms of a short non-bonded contact between H(17) and C(8'). Present calculations show that H(17) has two short intra-molecular contacts, and four inter-molecular contacts with the portion of the molecule

at $-x, -y, -z$ in cell (0,1,1):

H(17)...C(9)'	2.823 Å	H(17)...C(8)	2.991 Å
...C(8)'	2.991	...C(7)	3.057
		...H(17)	2.464
		...H(16)	2.555

The extra peak in the spectrum of the low melting form may be caused by C...H interactions but both inter- and intra-molecular contacts are involved. This would seem to cast some doubt on the argument that the high melting form must have a trans- configuration since it lacks this peak in its spectrum. The authors have reported that the high melting form is optically-active. This implies that it cannot crystallize in a centrosymmetric space group. Any crystal modification that reduced the number of C...H contacts would probably reduce inter-molecular repulsion forces. If this allowed the angle between the planes to expand, it would decrease the intra-molecular interactions involving H(17). It is not necessary to postulate a change in molecular configuration to explain the differences in the two spectra. However, the point will remain "not proven" until suitable crystals of the high melting polymorph become available for X-Ray analysis.

1.4. COMPARISON OF THEORY AND EXPERIMENT

Cruickshank and Sparks (1960) have shown that Pauling valence bond theory and Huckel molecular orbital theory give very good agreement with experiment for

naphthalene, anthracene and various alternant hydrocarbons.

They have recommended that a correlation of the form,

$$r = 1.477 - (1.477 - 1.337) \frac{1.333p}{0.333p + 1}$$

be used for valence-bond calculations, and have suggested that a linear variation between $m=0.40$ at $r = 1.46$ and $m = 0.85$ at $r = 1.34$ would improve the performance of simple molecular orbital theory. In these relationships, r is the bond length, p is the π -bond order calculated by the Pauling superposition method and m is the π -bond order calculated by the molecular orbital method. Trotter (1964) has recently used these order-length correlations to compare theoretical predictions with experimental results for a large number of aromatic molecules.

Table 9 shows π -bond orders for the various bonds in the asymmetric unit of 1,1'-binaphthyl. The numbering used corresponds to the normal convention for naphthalene. The H.M.O. bond orders quoted here have been taken from Coulson and Streitwieser (1965)

Valence bond theory predicts that the bond-lengths in the naphthalene nuclei of 1,1'-binaphthyl will be the same as those in naphthalene itself. For this reason, it is interesting to compare the lengths obtained in the present study with the values obtained by Cruickshank and Sparks for naphthalene and with those predicted by the two

theories. These lengths are compared in Table 10, and the root mean square differences are shown at the foot of each column.

A χ^2 test with 10 degrees of freedom shows that the dimensions of the naphthalenic residue are not significantly different from those of naphthalene. However, the differences between observed and theoretical values are highly significant. In both cases, agreement is worse for the short bonds than it is for the long ones and the major source of disagreement is the very short C(5)-C(6) bond. It is likely that the discrepancies will be reduced when the experimental values are corrected for vibrational oscillation. Some of the bonds that show worst agreement are the ones likely to be most affected by this correction.

Table 2

1,1'-Binaphthyl: Final Fractional Atomic Coordinates.
Standard deviations are given as units in the last place

C(1)	0.03550(10)	0.1057(44)	0.25208(25)
C(2)	0.06095(12)	-0.05065(44)	0.18985(29)
C(3)	0.12841(14)	-0.06107(48)	0.19519(31)
C(4)	0.17047(12)	0.08794(48)	0.26514(29)
C(5)	0.18919(11)	0.41283(47)	0.10531(29)
C(6)	0.16698(13)	0.56895(47)	0.47151(30)
C(7)	0.09985(13)	0.57249(44)	0.46830(29)
C(8)	0.05754(10)	0.43030(40)	0.39799(26)
C(9)	0.07905(10)	0.26431(38)	0.32733(25)
C(10)	0.14702(10)	0.25506(41)	0.33171(27)

Final coordinates and isotropic temperature parameters for the hydrogen atoms.

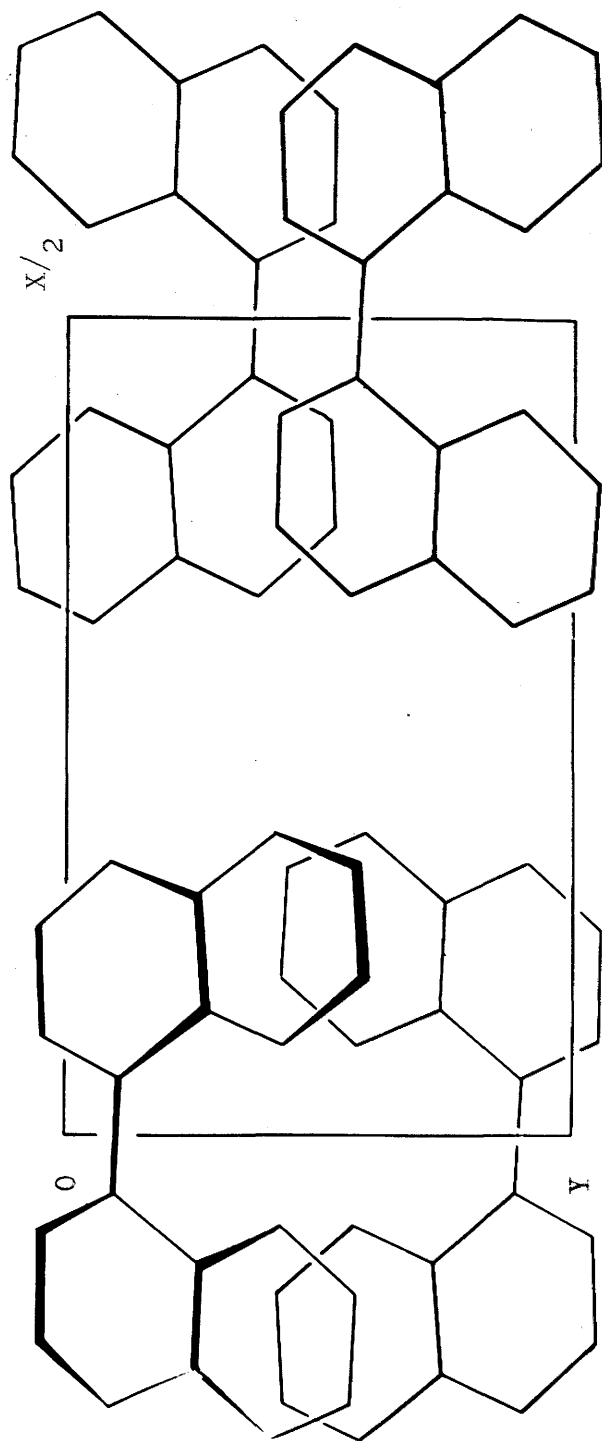
H(11)	0.0301(15)	-0.1524(49)	0.1337(30)	0.0385
H(12)	0.1443(17)	-0.1673(55)	0.1494(35)	0.0517
H(13)	0.2179(18)	0.0827(53)	0.2679(36)	0.0579
H(14)	0.2320(18)	0.4007(60)	0.4043(37)	0.0590
H(15)	0.1965(17)	0.6831(56)	0.5195(36)	0.0534
H(16)	0.0827(16)	0.6920(53)	0.5167(33)	0.0485
H(17)	0.0152(16)	0.4322(57)	0.3985(34)	0.0510

Table 3

Anisotropic temperature parameters of the carbon atoms.

	U	U	U	2U ₂₃	2U ₃₁	2U ₁₂
C(1)	0.0301	0.0360	0.0370	0.0071	0.0153	0.0051
C(2)	0.0379	0.0432	0.0501	-0.0071	0.0217	0.0114
C(3)	0.0435	0.0526	0.0542	-0.0078	0.0321	0.0325
C(4)	0.0262	0.0566	0.0492	0.0140	0.0149	-0.0049
C(6)	0.0359	0.0501	0.0529	-0.0042	0.0125	-0.0196
C(7)	0.0380	0.0461	0.0457	-0.0115	0.0186	-0.0089
C(8)	0.0270	0.0435	0.0407	-0.0062	0.0241	-0.0013
C(9)	0.0278	0.0395	0.0332	0.0093	0.0156	0.0090
C(10)	0.0280	0.0451	0.0370	0.0174	0.0.82	0.0062

Figure 2



The crystal structure of 1,1'-Binaphthyl: orthogonal projection along c^*

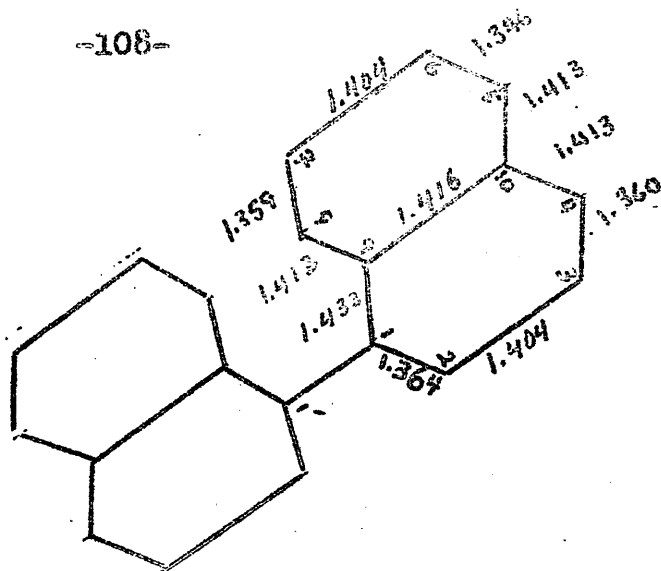


Table 4

1,1'-Binaphthyl: Bond Lengths.

Standard deviations are given as

units in the last place,

C(1)	C(1)	1.4748(46) Å
C(1)	C(2)	1.3645(37)
C(2)	C(3)	1.4036(38)
C(3)	C(4)	1.3598(41)
C(4)	C(10)	1.4127(39)
C(10)	C(5)	1.4135(37)
C(5)	C(6)	1.3450(42)
C(6)	C(7)	1.4037(38)
C(7)	C(8)	1.3590(37)
C(8)	C(9)	1.4128(35)
C(9)	C(10)	1.4162(29)
C(9)	C(1)	1.4328(32)

Bonds involving hydrogen.

C(2)	H(11)	0.983(31) Å
C(3)	H(12)	0.929(35)
C(4)	H(13)	0.989(37)
C(5)	H(14)	0.904(37)
C(6)	H(15)	0.995(36)
C(7)	H(16)	0.987(34)
C(8)	H(17)	0.889(34)

Table 5

1,1'-Binaphthyl: Bond Angles.(°)

C(1)	C(2)	C(3)	122.04	(25)
C(2)	C(3)	C(4)	119.84	(27)
C(3)	C(4)	C(10)	120.79	(24)
C(4)	C(10)	C(9)	119.33	(22)
C(4)	C(10)	C(5)	122.43	(21)
C(9)	C(10)	C(5)	118.23	(22)
C(10)	C(5)	C(6)	122.30	(22)
C(5)	C(6)	C(7)	119.81	(26)
C(6)	C(7)	C(8)	119.66	(26)
C(7)	C(8)	C(9)	122.11	(21)
C(8)	C(9)	C(10)	117.88	(21)
C(8)	C(9)	C(1)	123.07	(19)
C(10)	C(9)	C(1)	119.04	(21)
C(9)	C(1)	C(2)	118.95	(20)

Bond angles involving hydrogen.

C(1)	C(2)	H(11)	118.1	(1.8)
H(11)	C(2)	C(3)	119.7	(1.8)
C(2)	C(3)	H(12)	120.3	(2.1)
H(12)	C(3)	C(4)	119.8	(2.1)
C(3)	C(4)	H(13)	120.2	(2.2)
H(13)	C(4)	C(10)	119.0	(2.2)
C(10)	C(5)	H(14)	114.8	(2.4)
H(14)	C(5)	C(6)	122.8	(2.4)
C(5)	C(6)	H(15)	121.9	(2.1)
H(15)	C(6)	C(7)	118.2	(2.1)
C(6)	C(7)	H(16)	120.5	(1.9)
H(16)	C(7)	C(8)	119.8	(1.9)
C(7)	C(8)	H(17)	120.2	(2.3)
H(17)	C(8)	C(9)	117.7	(2.3)

Table 6

1,1'-Binaphthyl: Intra-molecular contacts < 3.50 A.

C(1)	C(2)	2.456
C(1)	C(9)	2.534
C(1)	H(11)	2.595
C(1)	H(17)	2.631
C(1)	C(8)	2.971
C(2)	H(11)	3.008
C(2)	C(2)	3.107
C(2)	H(17)	3.470
C(8)	H(17)	2.999
C(8)	C(8)	3.321
C(8)	C(9)	3.333
C(9)	H(17)	2.813
C(9)	C(9)	3.277
H(11)	H(11)	2.951
H(17)	H(17)	2.906

Table 7

Inter-molecular contacts. Equivalent position numbers refer to the following transformations.

1. x, y, z	5. $1/2+x, 1/2+y, z$
2. $-x, y, 1/2-z$	6. $1/2-x, 1/2+y, 1/2-z$
3. $-x, -y, -z$	7. $1/2-x, 1/2-y, -z$
4. $x, -y, -1/2+z$	8. $1/2+x, 1/2-y, -1/2+z$

C-H contacts < 3.15 A.

C(1)	H(16)	4	(0, 1, 0)	3.100
C(2)	H(16)	4	(0, 1, 0)	2.983
C(3)	H(16)	4	(0, 1, 0)	2.962
C(4)	H(16)	4	(0, 1, 0)	3.040
C(4)	H(15)	4	(0, 1, 0)	3.051
C(5)	H(14)	7	(0, 0, 1)	2.958
C(5)	H(13)	6	(0, 0, 0)	3.135
C(7)	H(17)	3	(0, 1, 1)	3.057
C(8)	H(17)	3	(0, 1, 1)	2.999
C(8)	H(11)	4	(0, 0, 1)	3.140
C(10)	H(16)	4	(0, 1, 0)	3.140

C-C contacts < 3.60 A.

C(2)	C(7)	1	(0, -1, 0)	3.600
C(3)	C(6)	1	(0, -1, 0)	3.580
C(5)	C(5)	7	(0, 0, 1)	3.450
C(6)	C(3)	1	(0, 1, 0)	3.580
C(7)	C(2)	1	(0, 1, 0)	3.600

H-H contacts < 2.50 A.

H(17)	H(17)	3	(0, 1, 1)	2.46
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Table 8

1,1-Binaphthyl: Mean Planes.

Deviations from the planes are shown in units of angstroms $\times 10^{-4}$.

A. Plane through naphthalene residue:

$$0.0512X + 0.5596Y - 0.8272Z = -1.6754$$

B. Plane through atoms 1,2,3,4,9,10:

$$0.0567X + 0.5544Y - 0.8303Z = -1.6774$$

C. Plane through atoms 5,6,7,8,9,10 :

$$0.0451X + 0.5653Y - 0.8237Z = -1.6536$$

	A	B	C
C(1)	99	10	3
C(2)	-6	14	-222
C(3)	-97	1	-400
C(4)	-105	-40	-371
C(5)	117	32	16
C(6)	-6	-197	12
C(7)	-127	396	-23
C(8)	-62	299	6
C(9)	77	-48	22
C(10)	110	63	-32

Root mean square deviation from planes,

90 37 20

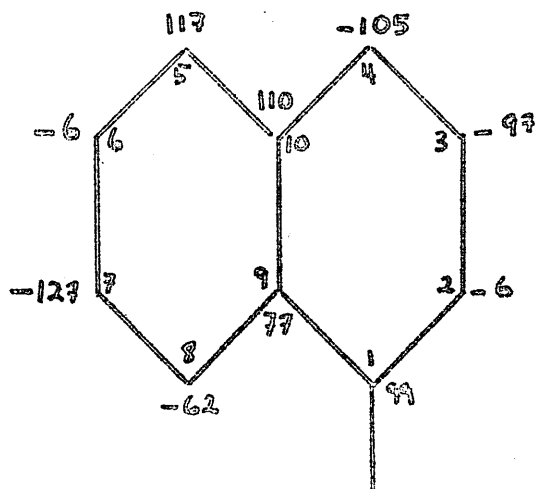


Table 9

1,1'-Binaphthyl: $\bar{\Pi}$ -Bond Orders.

Normal naphthalene numbering is used.

		V.B.	H.M.O.
C(1)	C(2)	0.667	0.659
C(2)	C(3)	0.333	0.621
C(3)	C(4)	0.667	0.712
C(4)	C(10)	0.333	0.550
C(10)	C(5)	0.333	0.554
C(5)	C(6)	0.333	0.722
C(6)	C(7)	0.333	0.607
C(7)	C(8)	0.667	0.717
C(8)	C(9)	0.333	0.569
C(9)	C(10)	0.333	0.523
C(1)	C(9)	0.333	0.507
C(1)	C(1)	0.000	0.404

Table 10

A comparison of bond lengths.

		<u>Binaphthyl</u>	<u>naphthalene</u>	<u>V.B.</u>	<u>H.M.O.</u>
C(1)	C(2)	1.364	1.364	1.375	1.391
C(2)	C(3)	1.404	1.415	1.421	1.401
C(3)	C(4)	1.360	1.364	1.375	1.377
C(4)	C(10)	1.413	1.421	1.421	1.420
C(10)	C(5)	1.413	1.421	1.421	1.419
C(5)	C(6)	1.346	1.364	1.375	1.374
C(6)	C(7)	1.404	1.415	1.421	1.405
C(7)	C(8)	1.359	1.364	1.375	1.376
C(8)	C(9)	1.413	1.421	1.421	1.415
C(9)	C(10)	1.416	1.418	1.421	1.427
C(9)	C(1)	1.433	1.421	1.421	1.432
C(1)	C(1)	1.475		1.477	1.459

Root mean square difference

0.009

0.014

0.015

1,1-Binaphthyl Structure Factor Tables Ctd.

H	K	L	Fo	Fo	H	K	L	Fo	Fo	H	K	L	Fo	Fo	H	K	L	Fo	Fo
18			10.7	-11.3	9			4.9	4.2	7			28.4	27.4	-22	4	4	2.6	2.0
2			25.5	-2.1	11			17.1	5.6	1			27.3	-26.3	-16	4	4	22.7	21.5
12			43.8	-23.4	7			12.2	-17.4	1			31.9	-29.1	-14	4	4	11.3	11.0
20			23.8	24.0	3			24.8	-25.0	1			0.9	5.3	-12	4	4	6.2	-6.0
28			26.7	24.8	3			30.4	-30.1	1			7.4	-1.3	-10	4	4	2.9	3.1
36			14.8	14.8	1			69.2	70.3	1			16.5	-15.5	-8	4	4	4.6	-3.0
44			54.4	56.4	1			7.2	-7.9	1			17.6	15.1	-6	4	4	10.5	-9.4
52			8.7	-8.5	7			11.8	-11.6	1			16.7	-16.9	-4	4	4	6.4	7.2
60			42.7	42.0	7			23.3	-25.4	1			51.9	51.8	0	4	4	10.0	11.1
68			26.7	25.7	7			5.5	-6.6	1			46.7	46.1	0	4	4	19.5	-19.6
76			28.5	-27.7	7			4.2	-6.6	1			25.4	-24.2	4	4	4	25.6	-27.1
84			22.0	21.5	7			3.7	2.8	1			10.7	-10.9	8	4	4	2.2	1.8
92			32.8	-33.1	7			2.7	-1.5	1			2.9	2.1	10	4	4	3.6	3.5
100			6.5	6.4	7			5.6	-6.0	0			3.7	-1.3	16	4	4	1.8	1.5
108			8.8	3.7	7			3.2	-5.1	0			23.3	-2.6	18	4	4	4.5	4.6
116			8.8	8.3	7			3.2	-6.2	0			14.1	14.3	20	4	4	5.8	5.5
124			4.4	-4.7	7			21.8	-22.9	0			10.2	11.3	22	4	4	3.7	3.6
132			10.1	11.0	7			8.8	8.9	0			9.9	9.3	24	4	4	3.7	3.6
140			7.1	7.0	7			8.8	3.4	0			17.7	-16.4	18	4	4	5.8	-4.7
148			4.8	5.0	7			8.8	-7.9	0			10.5	-9.2	14	4	4	3.2	3.5
156			10.7	12.3	11			2.7	-2.7	11			2.6	6.0	12	4	4	8.0	9.0
164			2.2	-2.7	11			3.1	-2.5	11			7.2	7.3	10	4	4	7.4	-8.1
172			33.2	30.9	19			13.1	-14.0	11			10.6	-10.6	8	4	4	16.0	16.0
180			22.2	21.2	19			7.7	-5.6	11			3.7	-2.2	6	4	4	8.7	8.5
188			15.5	14.5	19			2.9	-5.7	11			7.0	-5.1	4	4	4	3.0	-2.9
196			4.4	5.9	19			17.7	-18.4	10			4.1	-4.2	2	4	4	10.2	11.7
204			35.9	-33.8	19			15.2	15.9	10			6.1	-6.3	0	4	4	4.1	3.8
212			25.0	23.6	19			26.9	27.4	10			3.1	-2.4	4	4	4	27.0	26.4
220			7.5	3.7	19			9.7	-9.4	10			3.2	3.2	-6	4	4	9.5	8.7
228			40.4	38.6	19			10.0	-10.0	10			14.0	14.0	-8	4	4	22.1	21.8
236			4.4	5.1	19			37.8	-37.5	10			3.3	1.9	-12	4	4	16.0	-15.5
244			27.4	-27.3	19			5.5	-5.7	10			3.6	-4.4	-14	4	4	17.3	-17.4
252			9.9	9.4	19			10.2	10.0	10			4.3	5.4	-16	4	4	15.4	15.1
260			3.3	3.6	19			5.3	-5.2	10			7.0	7.0	-18	4	4	9.7	9.7
268			11.1	-7.8	19			2.2	1.4	10			6.1	-6.0	-20	4	4	3.4	-1.9
276			8.8	7.9	19			4.4	4.2	10			7.1	-7.6	-22	4	4	5.5	5.5
284			9.8	-7.9	19			7.7	8.4	10			24.4	-25.2	-24	4	4	4.6	-4.7
292			11.1	11.7	19			8.8	8.7	10			15.6	15.6	-18	4	4	7.0	-7.0
300			14.8	14.4	19			12.2	12.2	10			4.6	4.7	-14	4	4	28.9	-28.6
308			14.8	18.2	19			13.3	-12.3	10			3.2	-3.8	-10	4	4	3.2	-3.1
316			2.4	2.9	19			4.3	-5.5	10			6.7	5.2	-8	4	4	4.2	-4.0
324			16.6	15.9	19			5.5	-1.9	10			3.3	2.0	-6	4	4	25.2	23.0
332			15.5	15.3	19			12.2	-4.7	10			8.6	7.4	-4	4	4	11.4	10.5
340			31.9	30.0	19			13.3	10.9	10			5.5	4.8	-2	4	4	3.3	-2.9
348			9.9	9.9	19			15.5	-12.6	10			4.5	-5.1	0	4	4	11.1	10.0
356			1.1	1.5	19			7.5	-1.8	10			14.9	-16.0	8	4	4	23.6	22.6
364			17.3	18.1	19			6.6	-4.5	10			16.4	-17.1	10	4	4	2.2	-1.4
372			37.9	38.3	19			7.7	-1.9	10			3.6	3.6	12	4	4	19.0	-19.5
380			16.6	17.7	19			5.4	5.5	10			6.8	8.4	14	4	4	7.9	-8.5
388			16.6	11.1	19			4.4	4.2	10			8.9	-8.9	16	4	4	7.7	-7.7
396			26.6	23.3	19			3.3	3.5	10			5.1	4.7	18	4	4	6.7	-7.2
404			26.6	23.3	19			11.1	11.1	10			2.2	1.5	20	4	4	3.8	-5.3
412			13.5	-14.1	19			5.1	5.1	10			8.5	-8.1	22	4	4	9.1	7.4
420			16.6	16.8	19			12.2	-12.5	10			2.2	1.5	16	4	4	17.9	17.4
428			26.6	26.6	19			9.9	-9.9	10			8.5	-8.1	14	4	4	26.8	25.5
436			26.6	26.6	19			8.8	-8.9	10			3.0	3.0	12	4	4	12.0	-12.0
444			13.5	-14.1	19			9.9	-9.9	10			11.1	-11.1	10	4	4	3.6	-4.3
452			16.6	16.6	19			11.1	11.1	10			11.9	-13.2	8	4	4	36.4	-34.0
460			26.6	26.6	19			8.8	10.0	10			7.0	-5.4	6	4	4	3.7	-3.7
468			26.6	26.6	19			14.7	-14.8	10			5.1	-5.8	4	4	4	22.2	24.5
476			13.5	-13.3	19			18.3	-20.1	10			6.5	7.1	2	4	4	7.9	8.4
484			26.6	26.6	19			18.3	-21.2	10			6.8	2.6	0	4	4	1.7	-1.7
492			13.5	-13.3	19			7.1	7.3	10			7.2	4.5	-2	4	4	8.1	7.7
500			26.6	26.6	19			31.4	-29.7	10			2.7	-1.0	-4	4	4	2.5	-3.8
508			13.5	-13.3	19			18.3	18.6	10			14.0	-12.7	-6	4	4	21.9	-20.9
516			26.6	26.6	19			7.1	7.2	10			7.2	-6.3	-8	4	4	5.9	-4.6
524			13.5	-13.3	19			18.3	18.6	10			7.3	7.6	-10	4	4	32.2	31.0
532			26.6	26.6	19			67.0	-67.7	10			8.7	-10.4	-12	4	4	7.4	8.4
540			13.5	-13.3	19			24.7	-24.6	10			6.6	-8.2	-14	4	4	9.4	-9.1
548			26.6	26.6	19			15.4	15.1	10			5.4	-4.5	-16	4	4	6.7	-6.4
556			26.6	26.6	19			15.1	14.6	10			9.4	9.6	-18	4	4	3.3	1.7
564			26.6	26.6	19			17.4	16.7	10			4.1	-5.2	20	4	4	2.6	-2.5
572			26.6	26.6	19			4.5	-4.6	10			3.2	2.6	18	4	4	3.4	3.0
580			26.6	26.6	19			3.3	3.2	10			3.4	3.2	16	4	4	7.7	-7.7
588			26.6	26.6	19			2.2	2.2	10			11.1	-11.1	14	4	4	3.1	3.1
596			26.6	26.6	19			10.3	10.3	10			7.1	-7.3	12	4	4	12.1	11.5
604			26.6	26.6	19			20.3	-20.3	10			5.9	-6.5	10	4	4	8.3	-8.5
612			26.6	26.6	19			16.6	-16.5	10			6.6	7.4	8	4	4	15.3	-13.0
620			26.6	26.6	19			16.6	-16.5	10			2.1	2.4	6	4	4	32.7	-30.7
628			26.6	26.6	19			3.3	3.6	10			12.4	11.8	4	4	4	2.2	-1.7
636			26.6	26.6	19			4.7	4.1	10			30.8	31.7	2	4	4	5.1	5.2
644			26.6	26.6	19			18.3	-17.0	10			20.0	21.2	-3	4	4	4.3	4.4
652			26.6	26.6	19			18.3	-17.0	10			35.8	-36.8	-7	4	4	4.9	-4.4
660			26.6	26.6	19			2.0	-3.0	10			1.9	-1.7	-11	4	4	2.7	3.7
668			26.6	26.6	19			10.0	-9.2	10			13.7	-14.3	-15	4	4	4.5	4.6
676			26.6	26.6	19			6.4	5.4	10			12.3	13.6	-19	4	4	3.3	3.4
684			26.6	26.6	19			6.4	5.4	10			2.7	2.8	-23	4	4	3.2	1.4
692			26.6	26.6	19			18.3	-18.3	10			12.5	13.0	-19	4	4	2.8	-0.2
700			26.6	26.6	19			18.3	-18.3	10			19.3	-19.5	-17	4	4	12.1	11.2
708			26.6	26.6	19	</													

1,1-Binaphthyl Structure Factor Tables Ctd.

H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc
18			10.7	-11.3	11	9	8	4.9	4.2	7	1	1	28.4	27.4	-22	4	4	2.6	2.0
12			22.2	2.1	17	7	7	17.1	5.6	3	1	1	27.3	26.3	-16	4	4	22.7	21.5
0			29.5	-23.4	11	9	8	17.1	-17.4	3	1	1	31.9	29.1	-14	4	4	11.3	11.0
0			43.3	-40.2	12	5	5	12.5	-13.0	1	1	1	24.9	23.9	-12	4	4	6.6	-6.0
0			23.8	24.0	24	9	7	24.9	-25.1	3	1	1	30.4	30.1	-10	4	4	6.6	3.1
0			26.7	24.8	30	4	5	30.4	-30.1	5	1	1	14.8	14.8	-8	4	4	4.6	-3.0
0			54.4	56.4	69	2	2	69.2	70.3	7	1	1	54.4	56.4	-6	4	4	10.5	-9.4
0			8.7	-8.5	7	2	2	7.2	-7.9	9	1	1	42.7	42.0	-4	4	4	6.4	7.2
0			23.8	25.3	11	8	8	11.8	-11.6	-11	1	1	26.7	25.7	-2	4	4	10.0	11.1
0			26.7	-25.7	11	8	8	23.8	-25.4	-13	1	1	42.7	42.0	-2	4	4	19.5	-19.6
0			4.4	5.9	17	7	7	4.0	-6.6	-15	1	1	26.7	25.7	-2	4	4	25.6	-27.1
0			22.0	-21.5	17	7	7	22.0	-21.5	-17	1	1	4.4	5.9	-4	4	4	2.2	2.2
0			32.8	-33.6	22	3	3	32.8	-33.6	-23	1	1	10.7	10.7	-8	4	4	3.6	3.5
0			6.6	6.4	27	3	3	6.6	-6.0	-25	1	1	22.0	21.5	-10	4	4	11.0	12.2
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	6.6	6.4	-16	4	4	1.8	1.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-18	4	4	4.5	4.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-20	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-22	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-24	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-26	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-28	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-30	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-32	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-34	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-36	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-38	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-40	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-42	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-44	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-46	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-48	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-50	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-52	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-54	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-56	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-58	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-60	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-62	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-64	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-66	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-68	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-70	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-72	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-74	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-76	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-78	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-80	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-82	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-84	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-86	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-88	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-90	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-92	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-94	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-96	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-98	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-100	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-102	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-104	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-106	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-108	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-110	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-112	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-114	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-116	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-118	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-120	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-122	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-124	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-126	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-128	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-130	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-132	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-134	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-136	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-138	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-140	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-142	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-144	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-146	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-148	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-150	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6	-152	4	4	5.5	5.5
0			26.6	26.6	27	3	3	26.6	-26.6	-27	0	0	26.6	26.6					

1,1'-Binaphthyl Structure Factor Tables Ctd.

H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc
13	13	13	7.3	-7.4	25	5	1	3.1	-4.4	12	6	4	7.1	-8.1	-15	7	7	10.7	9.5
13	13	13	3.0	4.1	23	5	1	3.2	5.3	14	6	4	15.0	14.6	-25	7	7	3.2	-1.8
13	13	13	4.5	7.5	17	5	1	17.1	16.9	16	6	4	14.7	14.7	-23	7	7	3.3	3.0
13	13	13	4.9	4.9	13	5	1	26.1	-26.4	18	6	4	3.9	4.0	-21	7	7	7.7	-1.1
13	13	13	3.1	3.1	11	5	1	12.2	12.2	14	6	4	10.7	11.0	-17	7	7	3.3	3.6
13	13	13	4.4	4.4	9	5	1	6.4	6.4	12	6	4	6.2	6.9	-15	7	7	3.3	3.3
13	13	13	4.6	4.6	7	5	1	3.0	3.0	10	6	4	1.0	2.9	-13	7	7	2.6	2.9
13	13	13	4.3	4.3	5	5	1	7.0	7.0	8	6	4	2.6	2.3	-11	7	7	2.6	2.3
13	13	13	3.7	3.7	3	5	1	14.1	14.9	6	6	4	3.6	2.9	-7	7	7	6.7	5.7
13	13	13	3.9	3.9	1	5	1	5.2	-7.5	4	6	4	8.5	-8.6	-5	7	7	2.3	2.3
13	13	13	2.7	2.7	23	5	1	2.7	-3.0	2	6	4	3.1	-3.8	-3	7	7	7.7	6.6
13	13	13	1.6	1.6	21	5	1	12.4	-11.6	1	6	4	1.7	-1.7	1	7	7	4.4	4.4
13	13	13	1.5	1.5	19	5	1	5.6	-5.9	16	6	4	1.2	-1.2	1	7	7	3.3	3.3
13	13	13	1.1	1.1	17	5	1	1.7	-1.7	14	6	4	0.3	-0.3	1	7	7	3.1	3.1
13	13	13	1.1	1.1	15	5	1	3.9	3.7	12	6	4	0.6	0.6	1	7	7	2.5	2.5
13	13	13	1.1	1.1	13	5	1	4.0	3.7	10	6	4	0.9	0.9	1	7	7	1.5	1.5
13	13	13	1.1	1.1	11	5	1	3.8	3.8	8	6	4	0.8	0.8	1	7	7	0.8	0.8
13	13	13	1.1	1.1	9	5	1	6.4	6.4	6	6	4	0.7	0.7	1	7	7	0.7	0.7
13	13	13	1.1	1.1	7	5	1	7.6	7.6	4	6	4	0.9	0.9	1	7	7	0.9	0.9
13	13	13	1.1	1.1	5	5	1	6.8	6.8	2	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	3	5	1	7.7	7.7	1	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	1	5	1	3.8	3.8	1	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	25	5	1	1.8	1.8	24	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	23	5	1	7.2	7.2	22	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	21	5	1	1.8	1.8	20	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	19	5	1	7.4	7.4	18	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	17	5	1	3.8	3.8	16	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	15	5	1	6.6	6.6	14	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	13	5	1	1.8	1.8	12	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	11	5	1	7.2	7.2	10	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	9	5	1	1.8	1.8	8	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	7	5	1	7.7	7.7	6	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	5	5	1	1.5	1.5	4	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	3	5	1	7.7	7.7	2	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	1	5	1	5.2	5.2	1	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	25	5	1	5.1	4.7	24	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	23	5	1	3.2	3.3	22	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	21	5	1	3.9	3.3	20	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	19	5	1	7.5	7.5	18	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	17	5	1	6.6	6.6	16	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	15	5	1	1.8	1.8	14	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	13	5	1	7.2	7.2	12	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	11	5	1	1.8	1.8	10	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	9	5	1	7.7	7.7	8	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	7	5	1	1.5	1.5	6	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	5	5	1	7.7	7.7	4	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	3	5	1	5.2	4.7	2	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	1	5	1	5.1	4.4	1	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	25	5	1	3.2	3.3	24	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	23	5	1	3.9	3.3	22	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	21	5	1	7.5	7.5	20	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	19	5	1	6.6	6.6	18	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	17	5	1	1.8	1.8	16	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	15	5	1	7.2	7.2	14	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	13	5	1	1.8	1.8	12	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	11	5	1	7.7	7.7	10	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	9	5	1	1.5	1.5	8	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	7	5	1	7.7	7.7	6	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	5	5	1	5.2	4.7	4	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	3	5	1	5.1	4.4	2	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	1	5	1	3.2	3.3	1	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	25	5	1	3.9	3.3	24	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	23	5	1	7.5	7.5	22	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	21	5	1	6.6	6.6	20	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	19	5	1	1.8	1.8	18	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	17	5	1	7.2	7.2	16	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	15	5	1	1.8	1.8	14	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	13	5	1	7.7	7.7	12	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	11	5	1	1.5	1.5	10	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	9	5	1	7.7	7.7	8	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	7	5	1	5.2	4.7	6	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	5	5	1	5.1	4.4	4	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	3	5	1	3.2	3.3	2	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	1	5	1	3.9	3.3	1	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	25	5	1	7.5	7.5	24	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	23	5	1	6.6	6.6	22	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	21	5	1	1.8	1.8	20	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	19	5	1	7.2	7.2	18	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	17	5	1	1.8	1.8	16	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	15	5	1	7.7	7.7	14	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	13	5	1	1.5	1.5	12	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	11	5	1	7.7	7.7	10	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	9	5	1	5.2	4.7	8	6	4	1.1	1.1	1	7	7	1.1	1.1
13	13	13	1.1	1.1	7	5	1	5.1	4.4	6	6	4</							

THE CRYSTAL AND MOLECULAR STRUCTURE OF PERYLENE

In 1963, Tanaka discovered a second crystalline form of perylene which he designated as the β -form. From studies of absorption and fluorescence spectra of the two polymorphs, he concluded that the previously known α -form is a ground-state-excited-state dimer. To support his arguments, he determined the crystal structure of β -perylene in two projections. The $h0l$ projection is well resolved and was refined to an R-value of 17%. The y-coordinates were derived by assuming that the molecule is planar and has a structure similar to the α -form. Because there is serious overlap in the $0kl$ projection, the y-parameters were not refined. The R-value for this zone was 27%. In view of the current interest in molecules containing bonds with extreme π -bond orders, it seemed worthwhile to extend Tanaka's study to three dimensions.

The crystal structure of α -perylene was first established from projections along two axes by Donaldson, Robertson and White in 1953. Their measurement of $1.50 \pm 0.03 \text{ \AA}$ for the length of the peri bonds was in very good agreement with Coulson's (1951) theoretical prediction of 1.50 \AA . Camerman and Trotter (1964) have made a three-dimensional study of α -perylene using counter data. They report the value of $1.471 \pm 0.006 \text{ \AA}$ for the length of the long bonds and state that the molecule is significantly

distorted from planarity.

There are no other detailed analyses of perylene itself in the literature, but the molecule forms donor-acceptor complexes with a wide variety of acceptor molecules including fluorenil and pyromellitic dianhydride. The crystal structures of the π -molecular complexes of perylene with these two compounds have been determined (Hansen, 1963; Baeyens and Herbstreit, 1965).

The two polymorphs exhibit fluorescence when irradiated under ultra-violet light. This provides an easy method for distinguishing between them, since the fluorescence from the α -form is red and that from the β -form is green.

2.1. EXPERIMENTAL

There were a few crystals of the β -form in the sample of analytical grade perylene obtained from Koch-Light Laboratories. Since these were not suitable for single crystal studies, they were set aside to be used as seed crystals. When portions of the sample were recrystallized from benzene and from petroleum ether, the first "crop" consisted mainly of the α -form. If the supernatant liquid was decanted and seeded with crystals of the β -form, it yielded a good crop of very thin hexagonal plates that were too brittle to be cut. However, it was possible to grow very large crystals by allowing benzene solutions to evaporate slowly for several months. Crystals grown

in this manner were cut to the shape of a uniform rod and used to obtain cell dimensions and intensity data.

2.1.1. CRYSTAL DATA

Formula	$C_{20}H_{12}$	F.W. 252.3
System	monoclinic	
Lattice constants	$a = 9.777 \text{ \AA}$ $b = 5.904$ $c = 10.589$ $\beta = 96.75^\circ$	
Absent Spectra	$h0l$ if l odd $0k0$ if k odd	
Space group	$P2_1/c$	
Volume	607.0 \AA^3	
Density(obs.)	1.38 gm/cc	
Molecules/cell	2	
Density(calc.)	1.380 gm/cc	
Linear absorption coefficient μ , ($Mo K\alpha$; $\lambda = 0.7107 \text{ \AA}$)		0.920 cm^{-1}

Since there are two molecules per unit cell, the centre of the molecule must lie on a centre of inversion.

The cell dimensions listed here have been measured from precession and Weissenberg photographs and have been confirmed by checking reciprocal lattice spacings of high angle reflections on the diffractometer. The length of the c -axis is considerably shorter than Tanaka's value.

of 11.27 Å but the pattern of intensities is in agreement with the reflection data quoted in his paper.

2.1.2. REFLECTION DATA

Reflection data were obtained on a Hilger-Watts linear diffractometer using Mo K α radiation. The diffractometer is equipped with SrO-ZrO₂ balanced filters and a scintillation counter. Two crystals were used. Data from reciprocal lattice nets h0 ℓ to h4 ℓ were collected on one crystal and data from nets h5 ℓ to h7 ℓ were collected on the second. After Lorentz-polarisation corrections had been applied, the scale of the observed structure amplitudes was approximately twice the absolute scale. All data were included in the refinement. There are no "unobserved" planes because reflections with negative net counts are given an intensity of +1 by the data processing program.

2.2. REFINEMENT OF THE STRUCTURE

Although the cell dimensions determined in the present study were substantially different from those measured by Tanaka, the intensities of the reflections in the h0 ℓ zone agreed with the values quoted in Tanaka's paper. It seemed reasonable to use his coordinates as the starting point for the refinement of the structure.

When the "minimum residual" method of Rhuys and

Stanley (1963) was applied to the unique unit projection of the R-value fell from 37% to 16% in eight cycles. Refinement x- and z-coordinates from this projection were combined with Tanaka's calculated y-coordinates and used to compute structure factors for the five reciprocal lattice nets, h0l to h4l. The initial R-value was 54%, but this was reduced to 21.5% after three cycles of Fourier refinement. After termination-of-series corrections had been applied to the atomic parameters, refinement was continued by the method of least squares. Three isotropic least-squares calculations with unit weights improved the R-value to 18.9%. At this point, a weighting function of the form

$$w = 1 - \exp(-2 \sin^2 \theta / \lambda^2),$$

was applied. R was reduced to 15.0% by two further isotropic calculations but was improved by only 0.5% when the temperature factors of the carbon atoms were allowed to refine anisotropically. When hydrogen atoms were included in the structure-factor calculations, R fell to 13.6% and dropped to 12.7% after two cycles of anisotropic refinement.

Data from layers h5l to h7l were included at this point. Three full-matrix least-squares cycles were calculated to refine the scale factor between the two batches of data. In order to avoid a singularity in the matrix, the U_{22} temperature parameters were held constant.

The R-value was 16.4% for 1895 reflections.

The unweighted structure factors were analysed as a function of $\sin \theta/\lambda$, the index of the reciprocal lattice layer and the magnitude of F_{obs} as shown in Table 11. The weighting scheme devised from this examination of the data had the following form:

$$\text{if } \sin^2 \theta/\lambda^2 \leq 0.16, \text{ then } w = (\sin^2 \theta/\lambda^2)/0.16;$$

$$\text{if } |F_{\text{obs}}| \geq 7.0, \text{ then } w = 7/|F_{\text{obs}}|.$$

Planes with $|F_{\text{calc}}| \leq 1/3 |F_{\text{obs}}|$ were considered to be unreliably phased and were given a weight of zero.

A separate analysis of the weak reflections showed that R was over 200% for reflections with $|F_{\text{obs}}| < 0.5$ electrons and over 80% for reflections with $0.5 \leq |F_{\text{obs}}| < 1.0$. Most of the reflections with structure amplitudes smaller than one electron were among these that had been given an intensity of one count per minute by the data processing program because the net count was negative. It is therefore surprising to find that $|F_{\text{calc}}|$ is larger than $|F_{\text{obs}}|$ for nearly all of them. This may indicate that there is some systematic error in the measurement of the weak reflections. In order to investigate this point it will be necessary to alter the programs and to process the data again. As a temporary expedient the 468 planes with $|F_{\text{obs}}| < 1.0$ were excluded from the refinement. Of the remaining 1317 planes, 126 had been given a weight of zero. The R-value was 13.6% and $\Sigma w\Delta^2$ was 498.31. After

Table 11

β -Perylene: Analysis of Observed and Calculated
Structure Amplitudes for Reflections with $F_0 \neq 0$

As a function of $|F_{obs}|$

	$\Sigma F_o $	$\Sigma F_c $	$\Sigma A $	N	B	\bar{A}
2	930.90	823.96	196.50	933	0.533	0.53
3	643.56	547.49	172.37	262	0.268	0.66
4	488.46	461.19	80.93	141	0.166	0.57
8	1548.57	1534.66	153.33	273	0.099	0.56
16	1352.16	1297.64	97.74	123	0.072	6.79
33	1386.67	1355.39	73.08	64	0.053	1.14
111	381.81	391.05	26.56	9	0.070	2.95

As a function of $\sin \theta / \lambda$

0.20	491.51	494.32	51.01	32	0.104	1.59
0.40	2170.07	2089.89	234.40	279	0.108	0.84
0.50	1469.36	1489.22	150.42	298	0.102	0.50
0.60	1169.82	1133.33	196.38	441	0.168	0.44
0.70	967.06	887.18	262.60	490	0.272	0.54
0.80	406.60	286.39	173.66	228	0.425	0.76
1.00	57.71	31.05	32.58	37	0.564	0.88

As a function of reciprocal lattice index

0	939.05	930.39	99.06	140	0.105	0.71
1	1512.15	1489.49	163.62	260	0.108	0.63
2	1393.97	1351.83	165.52	262	0.119	0.63
3	782.22	740.37	131.03	254	0.158	0.52
4	777.81	752.31	125.58	237	0.162	0.53
5	618.68	595.58	140.14	249	0.226	0.58
6	356.25	282.97	129.42	208	0.363	0.62
7	352.00	268.44	146.14	204	0.415	0.72

three further cycles of least squares, R was 13.3%, $\sum w\Delta^2$ was 460.56 and the refinement of the carbon atoms had converged.

2.3. INTERIM RESULTS

Atomic coordinates are listed in Table 12 with their standard deviations. Temperature parameters are shown in Table 13. The hydrogen atoms were placed by a computer program and have not been refined; their temperature factors are probably slightly low.

Table 14 is a list of bond lengths and bond angles derived from the parameters of the carbon atoms. All C-H bonds are 1.09 Å long.

The polystyrene molecule is remarkably planar as is shown in Table 15. The equation of the "best" plane through the whole molecule is

$$0.2633X - 0.5793Y - 0.7714Z = 0 ,$$

where X, Y and Z are orthogonal axes defined by the equations

$$X = ax + c \cos \beta ,$$

$$Y = by ,$$

$$Z = cz \sin \beta .$$

The root mean square deviation from this plane (3.4×10^{-3} Å) is less than the average estimated standard deviation of atomic position. A more severe test of planarity is provided by the magnitude of the deviations from a plane

Table 12

β -Perylene: Final Fractional Atomic Coordinates. Standard deviations are given as units in the last place.

C(1)	0.05911(30)	-0.16889(61)	0.08899(27)
C(2)	0.11785(35)	-0.32806(75)	0.17450(35)
C(3)	0.26034(33)	-0.32329(70)	0.21807(34)
C(4)	0.34271(31)	-0.16218(67)	0.17535(29)
C(5)	0.37200(36)	0.17590(68)	0.04415(33)
C(6)	0.31745(31)	0.33318(56)	-0.03988(27)
C(7)	0.17620(30)	0.33403(57)	-0.08412(28)
C(8)	0.08948(35)	0.17241(68)	-0.04359(34)
C(9)	0.14413(37)	0.00226(78)	0.04423(37)
C(10)	0.28757(39)	0.00439(76)	0.08855(37)

Calculated positions of hydrogen atoms (not refined).

H(11)	0.0531	-0.4611	0.2082
H(12)	0.3032	-0.4486	0.2876
H(13)	0.4531	0.1654	0.2080
H(14)	0.4055	0.0082	0.0767
H(15)	0.3844	0.4609	-0.0748
H(16)	0.1347	0.4640	-0.1514

Table 13

Anisotropic Temperature Parameters of Carbon Atoms.

	U_{11}	U_{22}	U_{33}	$2U_{12}$	$2U_{23}$	$2U_{13}$
C(1)	0.0307	0.0377	0.0312	0.0001	0.0104	0.0020
C(2)	0.0353	0.0551	0.0452	0.0027	-0.0001	0.0145
C(3)	0.0287	0.0520	0.0439	-0.0123	0.0072	-0.0119
C(4)	0.0294	0.0449	0.0347	-0.0181	0.0079	0.0021
C(5)	0.0402	0.0427	0.0438	-0.0021	0.0084	-0.0237
C(6)	0.0328	0.0338	0.0310	-0.0017	0.0115	0.0006
C(7)	0.0315	0.0341	0.0333	-0.0096	0.0087	0.0098
C(8)	0.0363	0.0437	0.0480	0.0077	0.0117	0.0118
C(9)	0.0332	0.0566	0.0519	0.0066	0.0105	-0.0344
C(10)	0.0436	0.0507	0.0495	0.0271	0.0068	0.0234

Table 14

6. Perylene: Bond Lengths. (Å)

Standard deviations are given as units in the last place.

C(1)	C(2)	1.3822(54)
C(2)	C(3)	1.4154(50)
C(3)	C(4)	1.3574(54)
C(4)	C(10)	1.4093(58)
C(10)	C(5)	1.4209(59)
C(5)	C(6)	1.3516(51)
C(6)	C(7)	1.4052(44)
C(7)	C(8)	1.3781(51)
C(8)	C(9)	1.4292(60)
C(9)	C(10)	1.4256(56)
C(9)	C(1)	1.4245(56)
<u>peri-bonds</u>		1.4758(52)

C-H bonds are all 1.09 Å long.

Bond-Angles with their standard deviations.

C(1)	C(2)	C(3)	121.13(37)
C(2)	C(3)	C(4)	120.35(37)
C(3)	C(4)	C(10)	120.64(32)
C(4)	C(10)	C(5)	121.37(33)
C(4)	C(10)	C(9)	119.65(38)
C(9)	C(10)	C(5)	118.99(39)
C(10)	C(5)	C(6)	120.52(34)
C(5)	C(6)	C(7)	121.21(32)
C(6)	C(7)	C(8)	120.68(31)
C(7)	C(8)	C(9)	119.52(33)
C(8)	C(9)	C(10)	119.08(38)
C(8)	C(9)	C(1)	121.80(33)
C(10)	C(9)	C(1)	119.12(38)
C(9)	C(1)	C(2)	119.11(31)

Table 15

β -Perylene: Mean Planes.

Deviations are shown in Ångströms $\times 10^{-4}$.

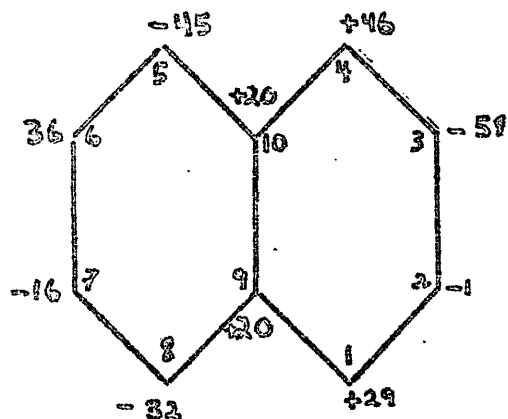
Plane A is the plane through the whole carbon skeleton.

Plane B is the plane through the central ring.

	A	B
C(1)	29	-8
C(2)	-1	-72
C(3)	-58	-141
C(4)	46	-14
C(5)	-45	-45
C(6)	36	70
C(7)	-16	31
C(8)	-32	-8
C(9)	20	8
C(10)	20	-4

Root mean square deviation:

34



through the central ring of the molecule. None of the atoms deviates significantly from this plane, which is defined by the equation.

$$0.2630X - 0.5773Y - 0.7730Z = 0.$$

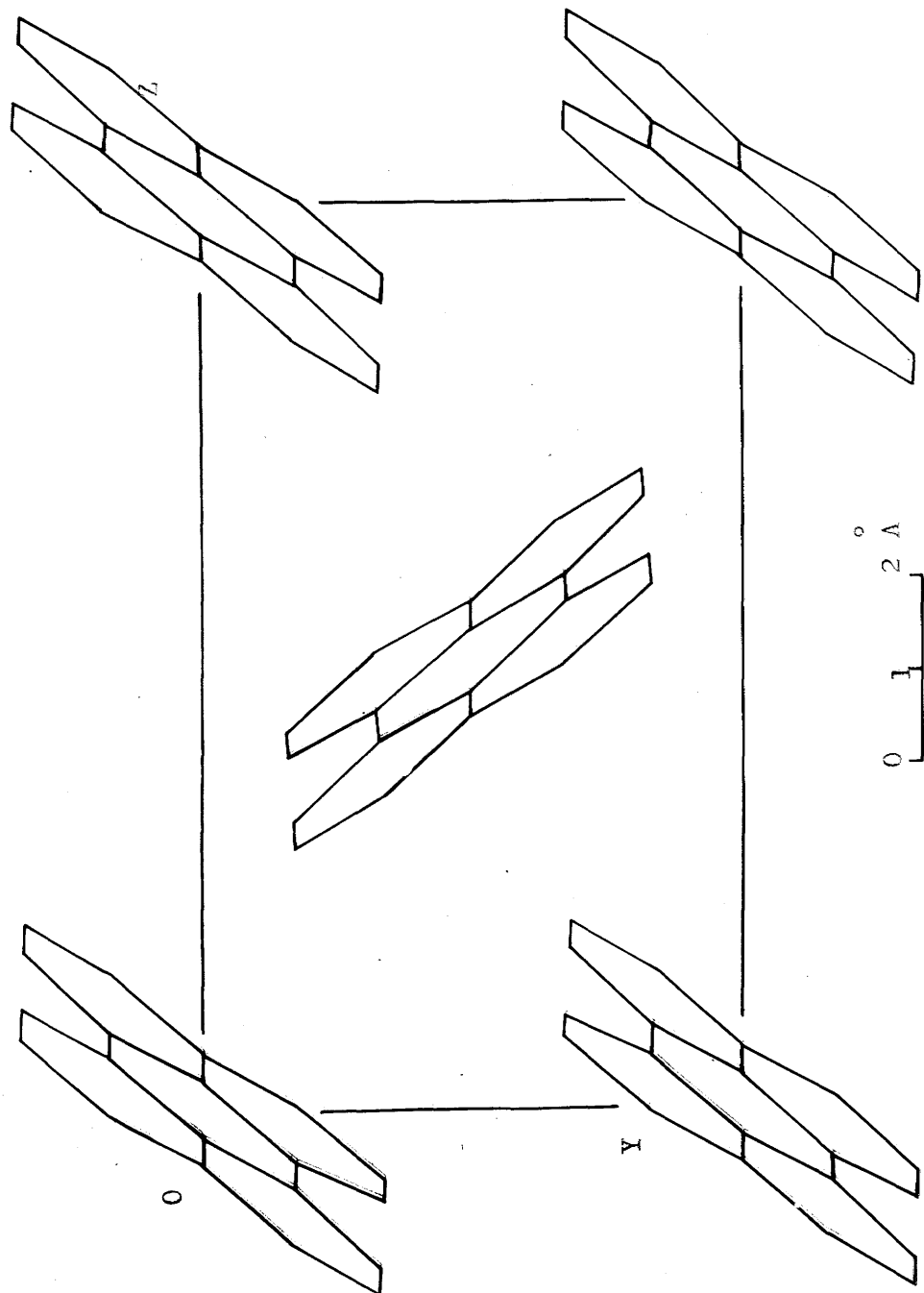
The crystal structure of β -perylene is analogous to those of naphthalene and anthracene (Abrahams, Robertson & White, 1949 ; Sinclair, Robertson & Mathieson, 1950).

The orthogonal projections shown in Figures 2 and 3 illustrate this similarity. Figure 2 is a projection of the whole unit cell on to the (100) plane. Figure 3 shows half of the projection on to (010).

It is interesting to compare the results of this analysis with theoretical predictions and with other crystallographic studies. Table 16 lists the bond lengths in β -perylene with those predicted by various theories. Table 17 summarizes the results of X-Ray studies involving β -perylene; and Table 18 compares the dimensions of β -perylene with those of quaterylene, naphthalene and 1,4-diacetylene.

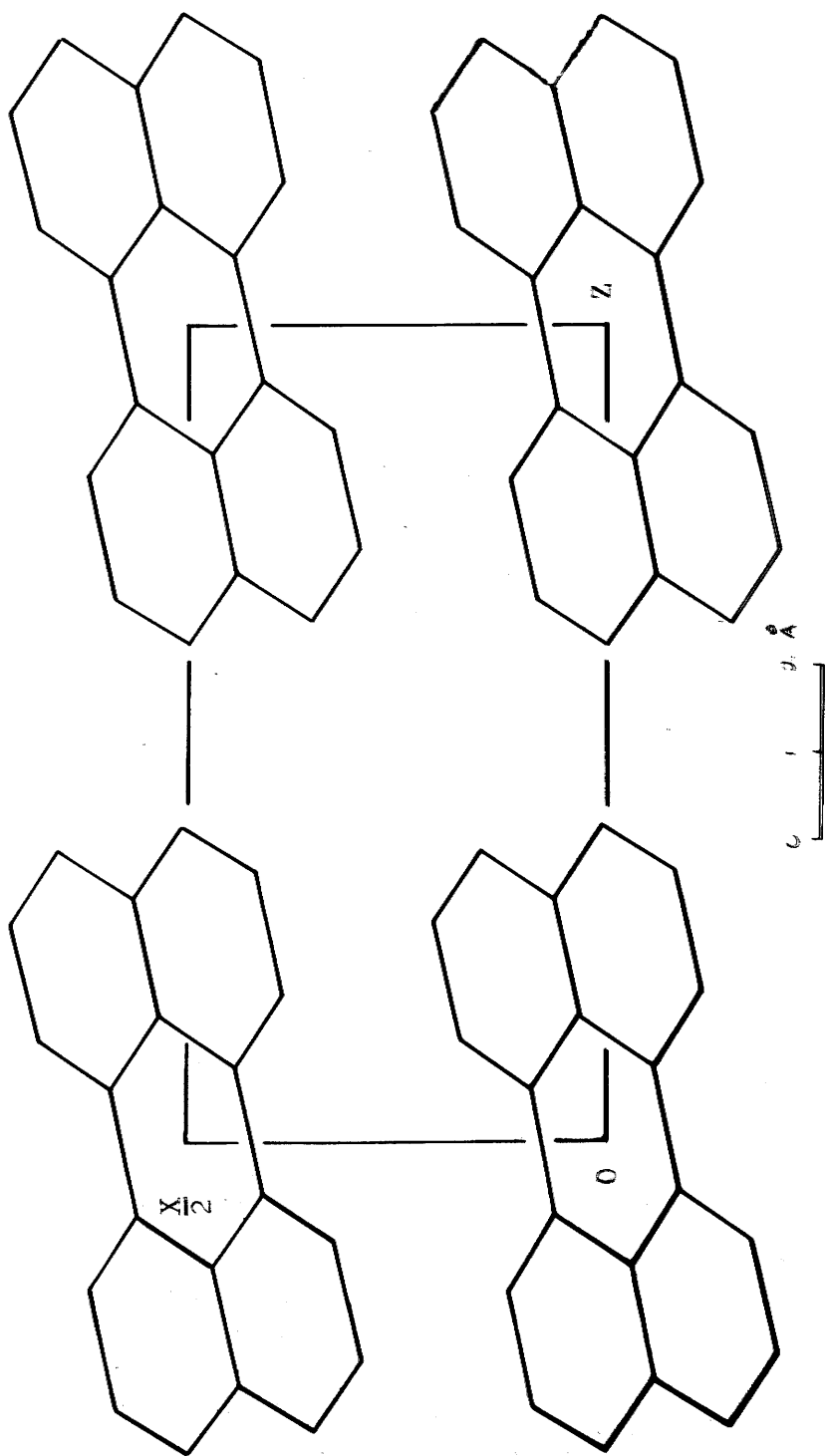
The H.M.O. bond-lengths in Table 16 have been obtained by combining the π -bond orders tabulated by Woodward and Brauman (1965) with the order length correlations suggested by Gmelchuk and Spinks (1960). Bond lengths predicted by the Revised Hückel-Mo (R.H.M.) method and the split π orbital (S.P.O.) method were taken from a paper by Bowen and Glotcher (1969). Bond lengths given

Figure 2



6-perylene: orthogonal projection on to (100)

Fig 3



β -perylene orthogonal projection on to (010)

differences between experimental and theoretical lengths are shown at the foot of each column. The goodness-of-fit becomes worse as the theory becomes more elegant. The length of the peri-bonds is the main source of disagreement, but even when this bond is not considered, the same general trend is observed. The relative π -bond orders predicted by molecular orbital theory follow the observed variations in bond length very closely.

The more elegant P.P.P. and S.P.O. calculations predict that the central bond in the naphthalenic residue should be shorter than the two outer bonds but the reverse situation is observed in the experiment. The 9-10 bond in β -perylene is longer than either the 2-3 bond or the 6-7 bond. Although these differences are not large enough to be significant, they are probably real since they are observed in 1,1'-binaphthyl and in the complexes of perylene listed in Table 17.

The dimensions of the perylene molecule obtained from various three-dimensional X-Ray studies are shown in Table 17. The average standard deviation of bond length appears at the top of each column; the root mean square difference is shown at the bottom. The perylene molecule is necessarily centrosymmetric in β -perylene, in the perylene-fluorenil complex (Hanson, 1963), and in the perylene-PMDA complex (Boeyens and Herbstein, 1963), but not in α -perylene (Gammann and Trotter, 1964). The

β -Perylene: Bond Length Comparisons

Table 16

		β -perylene	V.B.	H.M.O.	P.P.P.	S.P.O.
C(1)	C(2)	1.380	1.375	1.395	1.381	1.375
C(2)	C(3)	1.410	1.421	1.399	1.413	1.419
C(3)	C(4)	1.355	1.375	1.378	1.375	1.371
C(4)	C(10)	1.415	1.421	1.420	1.421	1.425
C(9)	C(10)	1.426	1.421	1.427	1.404	1.397
C(9)	C(1)	1.427	1.421	1.426	1.422	1.426
peri-bonds		1.476	1.477	1.457	1.456	1.462

Root mean square difference: peri-bonds included.

0.010 0.013 0.014 0.015

Root mean square difference: peri-bonds excluded.

0.010 0.011 0.012 0.013

Table 17

		β -perylene $\sigma=0.005$	perylene- fluorani $\sigma=0.005$	perylene PMDA $\sigma=0.010$	α -perylene $\sigma=0.011$
C(1)	C(2)	1.382	1.380	1.393	1.381
C(2)	C(3)	1.415	1.419	1.407	1.401
C(3)	C(4)	1.357	1.358	1.364	1.360
C(4)	C(10)	1.409	1.405	1.366	1.412
C(10)	C(5)	1.421	1.415	1.445	1.396
C(5)	C(6)	1.352	1.364	1.366	1.362
C(6)	C(7)	1.405	1.401	1.383	1.405
C(7)	C(8)	1.378	1.379	1.381	1.384
C(8)	C(9)	1.429	1.427	1.407	1.427
C(9)	C(10)	1.426	1.434	1.443	1.434
C(9)	C(1)	1.424	1.431	1.423	1.420
peri-bonds		1.476	1.473	1.493	1.473

Root mean square difference:

0.005

0.019

0.009

bond lengths listed for α -perylene are the results of Mason's (private communication) refinement of the data published by Cameron and Trotter. It should be noted that the average standard deviation reported here is approximately twice the estimate quoted in the original paper, despite a much lower R-value.

In Table 18, β -perylene is compared with quaterylene (Shrivastava and Speakman, 1960), naphthalene (Cruickshank and Sparks, 1960), and 1,1'-binaphthyl. The root mean square differences between the dimensions of β -perylene and those of the other molecules are small except in the case of quaterylene. This is likely to be a reflection of the fact that the structure of quaterylene was determined from two projections; it does not necessarily imply a real difference in molecular structure.

DISCUSSION

Although the analyses of these structures are not yet complete, the precision of the measured bond-lengths (0.003 to 0.006 Å) is sufficient to allow a certain amount of discussion.

Cruickshank (1956) has shown that when libration corrections are neglected, the apparent bond lengths are systematically shorter than their true values. In 1,1'-binaphthyl, the thermal vibrations are markedly anisotropic and libration corrections have not yet been applied. The bond that is likely to be most affected by these corrections

is the 5-6 bond. In the perylene structure, libration corrections will probably be very small since there is little anisotropy in the thermal parameters. In the present state of the analysis, the molecule appears to have its maximum vibration amplitude almost perpendicular to the plane of the ring. This unlikely situation can be explained in terms of a poor scale factor between the two batches of data.

The dimensions of 1,1'-binaphthyl and β -perylene are very similar. The root mean square difference is only 0.008 when "chemically equivalent" bonds are averaged. If the individual bond lengths are compared, the root mean square difference increases to 0.011. However, none of the differences between corresponding bond-lengths is significant.

The lengths of the sp_2-sp_2 bonds reported in this study are $1.475 \pm 0.005 \text{ \AA}$ in 1,1'-binaphthyl and $1.476 \pm 0.005 \text{ \AA}$ in β -perylene. These lengths are in good agreement with the value of $1.473 \pm 0.005 \text{ \AA}$ in the perylene-fluorenil complex and the value of $1.473 \pm 0.010 \text{ \AA}$ in α -perylene (Mason's refinement) but they are in disagreement with the values of 1.53 \AA in quaterylene, $1.493 \pm 0.010 \text{ \AA}$ in the perylene-PMDA complex, and $1.493 \pm 0.015 \text{ \AA}$ from electron diffraction measurements of perylene in the vapour phase (Trastberg, 1965).

The para-bond length quoted for quaterxylene is the average value of six independent bond lengths and would be expected to be very accurate in spite of the fact that the structure was determined from two projections. The discrepancy between this value and the 1.48 Å single bond length predicted by Devar and Schneising (1959) has been attributed to a stretching of the linking bonds caused by steric repulsions between the hydrogen atoms in the ortho-positions. However, recent studies suggest that hydrogen contacts of this type are "softer" than had been supposed. Bailey and Mason (1963) have reported H...H contacts of 1.85 Å in 1,8-bisdehydro-[14]-annulene. Hirshfeld, Rabinovich and Bregman (1965) have found that the mean separation of the inner hydrogens in [16]-annulene is 1.97 Å. There is no apparent deformation of C-C bonds in either structure. Agreement between the lengths of the linking bonds in the two molecules in the present study would seem to confirm that H...H contacts of this type do not cause significant lengthening of C-C bonds.

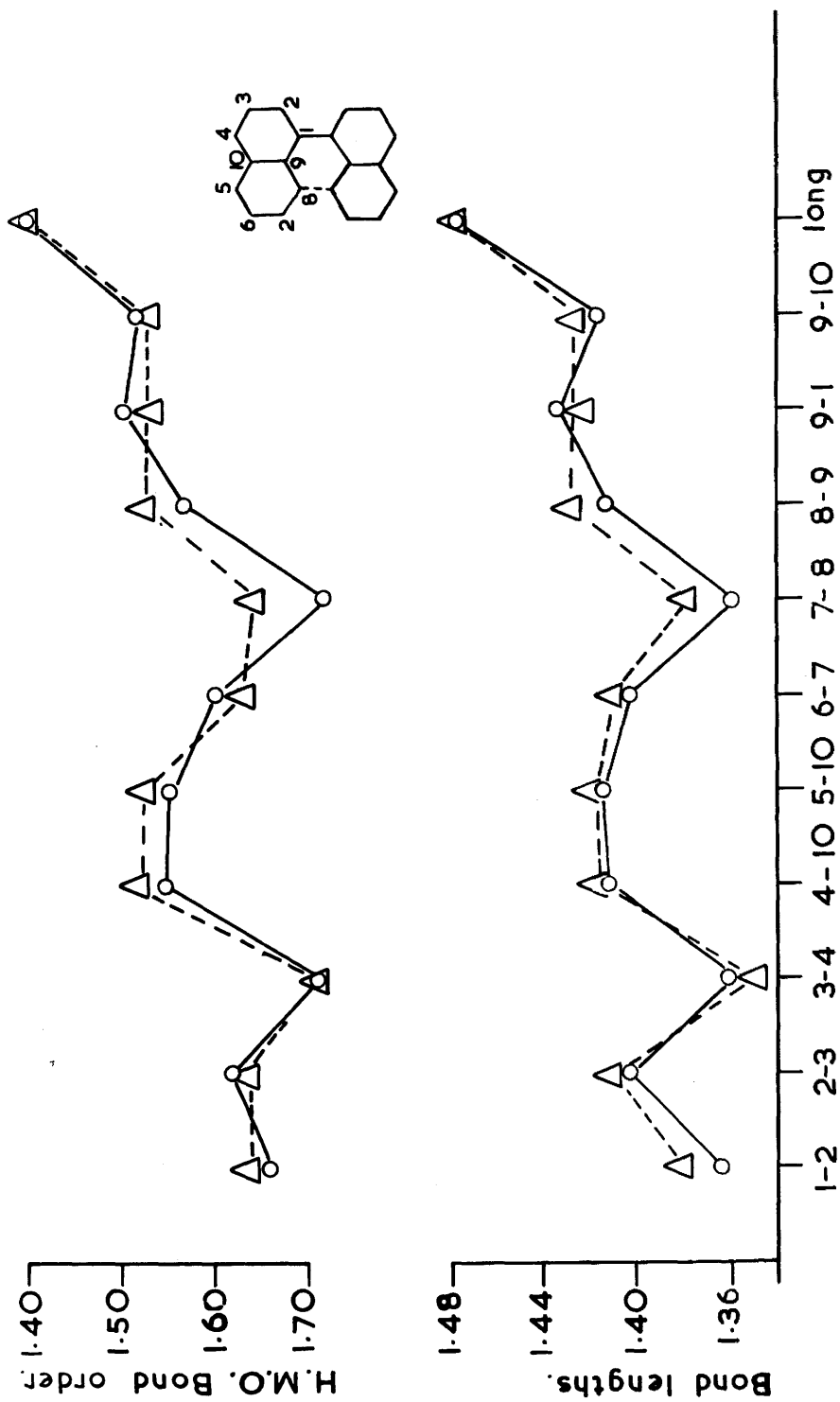
Differences between the lengths quoted in this work and those observed in the perylene-PMMA complex are, in general, quite large; but certain experimental details in the latter study should be considered. Thermal vibration amplitudes in the complex are large and anisotropic; limited computer capacity made it necessary to refine the parameters of only 6 atoms in each least-squares cycle; the refinement was terminated before convergence.

It seems likely that the standard deviations have been underestimated.

The electron diffraction study is difficult to evaluate. Perylene is a very large molecule to be investigated by this method, and certain simplifying assumptions must be made. Traetteberg has assumed that the naphthalenic residues in perylene have the same molecular dimensions as naphthalene itself and has determined the lengths of the peri-bonds by subtracting the molecular intensity function of naphthalene from the radial distribution function for perylene. Results obtained from a calculation of this type are extremely sensitive to changes in the model. On the other hand, it is possible that there are real differences in the dimensions of the molecule in the vapour phase (300°C) and in the solid.

It is interesting to note that the trend in π -bond orders predicted by the Hückel molecular orbital method is very close to the trend in the bond lengths observed in this study. (See Figure 4). This suggests that the goodness-of-fit will probably improve as more accurate bond-length determinations become available and better order-length curves are devised. The most serious discrepancy between the experimental bond lengths in β -perylene and the predictions of P.P.P. and S.F.O. calculations is the relative π -bond order of the central

-134-
Figure 4



Trend in π - bond order and observed bond length
 perylene -o- 1,1'-binaphthyl Δ

bond of the naphthalenic residue. According to these two theories, this bond should be shorter than the outer bonds, but the reverse situation is observed in the four structures involving the perylene molecule and in the structure of 1,1'-binaphthyl.

The agreement between the dimensions of the two polymorphs of perylene and between perylene and 1,1'-binaphthyl is better than one would expect from the size of the standard deviations estimated for the individual structure determinations. None of the differences between these molecules is significant at the level of precision of the analyses. The dimensions of the naphthalenic residues do not differ significantly from those of naphthalene itself.

CONCLUSIONS

The results of the present work show that the bond lengths in the naphthalenic residues of 1,1'-binaphthyl and β -perylene are the same, within the experimental error, as those of naphthalene itself. This can be interpreted as direct evidence that the aromatic character of the perylene molecule is localised in the naphthalenic residues.

The average value for the sp_2-sp_2 single bond lengths in these two structures is 1.475 ± 0.005 Å.

B-perylene Structure Factor Tables

H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc	H	K	L	Fo	Fc
11	11	11	1.6	0.0	11	11	11	1.5	0.0	6	4	5	7.2	7.3	1	1	1	5.2	5.9
11	11	10	0.9	0.0	11	11	10	2.5	0.4	1	1	2	10.5	10.5	1	1	3	11.7	11.5
11	11	9	0.6	0.0	11	11	9	0.0	0.0	1	1	4	19.0	18.8	1	1	5	19.0	18.8
11	11	8	0.3	0.0	11	11	8	0.0	0.0	1	1	6	29.9	29.2	1	1	7	29.9	29.2
11	11	7	0.0	0.0	11	11	7	0.0	0.0	1	1	8	42.8	42.8	1	1	9	42.8	42.8
11	11	6	0.0	0.0	11	11	6	0.0	0.0	1	1	10	57.7	57.7	1	1	10	57.7	57.7
11	11	5	0.0	0.0	11	11	5	0.0	0.0	1	1	11	72.6	72.6	1	1	11	72.6	72.6
11	11	4	0.0	0.0	11	11	4	0.0	0.0	1	1	12	87.5	87.5	1	1	12	87.5	87.5
11	11	3	0.0	0.0	11	11	3	0.0	0.0	1	1	13	102.4	102.4	1	1	13	102.4	102.4
11	11	2	0.0	0.0	11	11	2	0.0	0.0	1	1	14	117.3	117.3	1	1	14	117.3	117.3
11	11	1	0.0	0.0	11	11	1	0.0	0.0	1	1	15	132.2	132.2	1	1	15	132.2	132.2
11	10	11	1.6	0.0	11	10	11	1.5	0.0	1	10	11	1.5	0.0	1	10	11	1.5	0.0
11	10	10	0.9	0.0	11	10	10	2.5	0.4	1	10	10	2.5	0.4	1	10	10	2.5	0.4
11	10	9	0.6	0.0	11	10	9	0.0	0.0	1	10	9	0.0	0.0	1	10	9	0.0	0.0
11	10	8	0.3	0.0	11	10	8	0.0	0.0	1	10	8	0.0	0.0	1	10	8	0.0	0.0
11	10	7	0.0	0.0	11	10	7	0.0	0.0	1	10	7	0.0	0.0	1	10	7	0.0	0.0
11	10	6	0.0	0.0	11	10	6	0.0	0.0	1	10	6	0.0	0.0	1	10	6	0.0	0.0
11	10	5	0.0	0.0	11	10	5	0.0	0.0	1	10	5	0.0	0.0	1	10	5	0.0	0.0
11	10	4	0.0	0.0	11	10	4	0.0	0.0	1	10	4	0.0	0.0	1	10	4	0.0	0.0
11	10	3	0.0	0.0	11	10	3	0.0	0.0	1	10	3	0.0	0.0	1	10	3	0.0	0.0
11	10	2	0.0	0.0	11	10	2	0.0	0.0	1	10	2	0.0	0.0	1	10	2	0.0	0.0
11	10	1	0.0	0.0	11	10	1	0.0	0.0	1	10	1	0.0	0.0	1	10	1	0.0	0.0
11	9	11	1.6	0.0	11	9	11	1.5	0.0	1	9	11	1.5	0.0	1	9	11	1.5	0.0
11	9	10	0.9	0.0	11	9	10	2.5	0.4	1	9	10	2.5	0.4	1	9	10	2.5	0.4
11	9	9	0.6	0.0	11	9	9	0.0	0.0	1	9	9	0.0	0.0	1	9	9	0.0	0.0
11	9	8	0.3	0.0	11	9	8	0.0	0.0	1	9	8	0.0	0.0	1	9	8	0.0	0.0
11	9	7	0.0	0.0	11	9	7	0.0	0.0	1	9	7	0.0	0.0	1	9	7	0.0	0.0
11	9	6	0.0	0.0	11	9	6	0.0	0.0	1	9	6	0.0	0.0	1	9	6	0.0	0.0
11	9	5	0.0	0.0	11	9	5	0.0	0.0	1	9	5	0.0	0.0	1	9	5	0.0	0.0
11	9	4	0.0	0.0	11	9	4	0.0	0.0	1	9	4	0.0	0.0	1	9	4	0.0	0.0
11	9	3	0.0	0.0	11	9	3	0.0	0.0	1	9	3	0.0	0.0	1	9	3	0.0	0.0
11	9	2	0.0	0.0	11	9	2	0.0	0.0	1	9	2	0.0	0.0	1	9	2	0.0	0.0
11	9	1	0.0	0.0	11	9	1	0.0	0.0	1	9	1	0.0	0.0	1	9	1	0.0	0.0
11	8	11	1.6	0.0	11	8	11	1.5	0.0	1	8	11	1.5	0.0	1	8	11	1.5	0.0
11	8	10	0.9	0.0	11	8	10	2.5	0.4	1	8	10	2.5	0.4	1	8	10	2.5	0.4
11	8	9	0.6	0.0	11	8	9	0.0	0.0	1	8	9	0.0	0.0	1	8	9	0.0	0.0
11	8	8	0.3	0.0	11	8	8	0.0	0.0	1	8	8	0.0	0.0	1	8	8	0.0	0.0
11	8	7	0.0	0.0	11	8	7	0.0	0.0	1	8	7	0.0	0.0	1	8	7	0.0	0.0
11	8	6	0.0	0.0	11	8	6	0.0	0.0	1	8	6	0.0	0.0	1	8	6	0.0	0.0
11	8	5	0.0	0.0	11	8	5	0.0	0.0	1	8	5	0.0	0.0	1	8	5	0.0	0.0
11	8	4	0.0	0.0	11	8	4	0.0	0.0	1	8	4	0.0	0.0	1	8	4	0.0	0.0
11	8	3	0.0	0.0	11	8	3	0.0	0.0	1	8	3	0.0	0.0	1	8	3	0.0	0.0
11	8	2	0.0	0.0	11	8	2	0.0	0.0	1	8	2	0.0	0.0	1	8	2	0.0	0.0
11	8	1	0.0	0.0	11	8	1	0.0	0.0	1	8	1	0.0	0.0	1	8	1	0.0	0.0
11	7	11	1.6	0.0	11	7	11	1.5	0.0	1	7	11	1.5	0.0	1	7	11	1.5	0.0
11	7	10	0.9	0.0	11	7	10	2.5	0.4	1	7	10	2.5	0.4	1	7	10	2.5	0.4
11	7	9	0.6	0.0	11	7	9	0.0	0.0	1	7	9	0.0	0.0	1	7	9	0.0	0.0
11	7	8	0.3	0.0	11	7	8	0.0	0.0	1	7	8	0.0	0.0	1	7	8	0.0	0.0
11	7	7	0.0	0.0	11	7	7	0.0	0.0	1	7	7	0.0	0.0	1	7	7	0.0	0.0
11	7	6	0.0	0.0	11	7	6	0.0	0.0	1	7	6	0.0	0.0	1	7	6	0.0	0.0
11	7	5	0.0	0.0	11	7	5	0.0	0.0	1	7	5	0.0	0.0	1	7	5	0.0	0.0
11	7	4	0.0	0.0	11	7	4	0.0	0.0	1	7	4	0.0	0.0	1	7	4	0.0	0.0
11	7	3	0.0	0.0	11	7	3	0.0	0.0	1	7	3	0.0	0.0	1	7	3	0.0	0.0
11	7	2	0.0	0.0	11	7	2	0.0	0.0	1	7	2	0.0	0.0	1	7	2	0.0	0.0
11	7	1	0.0	0.0	11	7	1	0.0	0.0	1	7	1	0.0	0.0	1	7	1	0.0	0.0
11	6	11	1.6	0.0	11	6	11	1.5	0.0	1	6	11	1.5	0.0	1	6	11	1.5	0.0
11	6	10	0.9	0.0	11	6	10	2.5	0.4	1	6	10	2.5	0.4	1	6	10	2.5	0.4
11	6	9	0.6	0.0	11	6	9	0.0	0.0	1	6	9	0.0	0.0	1	6	9	0.0	0.0
11	6	8	0.3	0.0	11	6	8	0.0	0.0	1	6	8	0.0	0.0	1	6	8	0.0	0.0
11	6	7	0.0	0.0	11	6	7	0.0	0.0	1	6	7	0.0	0.0	1	6	7	0.0	0.0
11	6	6	0.0	0.0	11	6	6	0.0	0.0	1	6	6	0.0	0.0	1	6	6	0.0	0.0
11	6	5	0.0	0.0	11	6	5	0.0	0.0	1	6	5	0.0	0.0	1	6	5	0.0	0.0
11	6	4	0.0	0.0	11	6	4	0.0	0.0	1	6	4	0.0	0.0	1	6	4	0.0	0.0
11	6	3	0.0	0.0	11	6	3	0.0	0.0	1	6	3	0.0	0.0	1	6	3	0.0	0.0
11	6	2	0.0	0.0	11	6	2	0.0	0.0	1	6	2	0.0	0.0	1	6	2	0.0	0.0
11	6	1	0.0	0.0	11	6	1	0.0	0.0	1	6	1	0.0	0.0	1	6	1	0.0	0.0
11	5	11	1.6	0.0	11	5	11	1.5	0.0	1	5	11	1.5	0.0	1	5	11	1.5	0.0
11	5	10	0.9	0.0	11	5	10	2.5	0.4	1	5	10	2.5	0.4	1	5	10	2.5	0.4
11	5	9	0.6	0.0	11	5	9	0.0	0.0	1	5	9	0.0	0.0	1	5	9	0.0	0.0
11	5	8	0.3	0.0	11	5	8	0.0	0.0	1	5	8	0.0	0.0	1	5	8	0.0	0.0
11	5	7	0.0	0.0	11	5	7	0.0	0.0	1	5	7	0.0	0.0	1	5	7	0.0	0.0
11	5	6	0.0	0.0	11	5	6	0.0	0.0	1	5	6	0.0	0.0	1	5	6	0.0	0.0
11	5	5	0.0	0.0	11	5	5	0.0	0.0	1	5	5	0.0	0.0	1	5	5	0.0	0.0
11	5	4	0.0	0.0	11	5	4	0.0	0.0	1	5	4	0.0	0.0	1	5	4	0.0	0.0
11	5	3	0.0	0.0	11	5	3	0.0	0.0	1	5	3	0.0	0.0	1	5	3	0.0	0.0
11	5	2	0.0	0.0	11	5	2	0.0	0.0	1	5	2	0.0	0.0	1	5	2	0.0	0.0
11	5	1	0.0	0.0	11	5	1	0.0	0.0	1	5	1	0.0	0.0	1	5	1	0.0	0.0
11	4	11	1.6	0.0	11	4	11	1.5	0.0	1	4	11	1.5	0.0	1	4	11	1.5	0.0
11	4	10	0.9	0.0	11	4	10	2.5	0.4	1	4	10	2.5	0.4	1	4	10	2.5	0.4
11	4	9	0.6	0.0	11	4	9	0.0	0.0	1	4	9	0.0	0.0	1	4	9	0.0	0.0
11	4	8	0.3	0.0	11	4	8	0.0	0.0	1	4	8	0.0	0.0	1	4	8	0.0	0.0
11	4	7	0.0	0.0	11	4	7	0.0	0.0	1	4	7	0.0	0.0	1	4	7	0.0	0.0
11	4	6	0.0	0.0	11	4	6	0.0	0.0	1	4	6	0.0	0.0	1	4	6	0.0	0.0
11	4	5	0.0	0.0	11	4	5	0.0	0.0										

Appendix I

Comments on Choice of
Unit Cell in a Triclinic Lattice

CHOICE OF UNIT CELL IN A TRICLINIC SYSTEM

In the triclinic system, the choice of unit cell is not dictated by symmetry considerations; thus there are an infinite number of ways of choosing primitive cells in a triclinic lattice. However, each lattice is uniquely described by the Bravais "reduced cell"; the cell whose edges are the three shortest non-co-planar translations in the lattice. For purposes of compilation, it is desirable to describe the lattice in terms of this reduced cell. The two most common algorithms for deriving a reduced cell from any arbitrary primitive cell are the Delaunay reduction and the Dirichlet reduction. There are certain reasons for preferring the latter.

In any parallelepiped, sets of angles occur in enantiomorphous pairs at opposite corners. One pair of corners has the angles α, β, γ homogeneous; that is, they are all obtuse (or all acute). The other three pairs of corners are of mixed types α, β', γ' or α', β, γ' , where α', β', γ' are the acute (or obtuse) supplements of α, β, γ . If we define vectors $\underline{a}, \underline{b}, \underline{c}$ along the edges from a homogeneous corner, the scalar products $\underline{a} \cdot \underline{b}$, $\underline{b} \cdot \underline{c}$, and $\underline{a} \cdot \underline{c}$ all have the same sign. This allows a cell to be defined uniquely in terms of the homogeneous triplet with the three shortest translations. Within the limits imposed by the condition that the angles be homogeneous,

there are two possible parallelepipeds - one obtuse and the other acute. Only one of these corresponds to the Bravais-reduced cell with edges shorter than any face- or body-diagonal.

The Delaunay reduced cell is tri-obtuse and so will correspond to the Bravais-reduced cell only when the latter happens to be obtuse. When the Bravais-reduced cell is tri-acute, the Delaunay-reduced cell will have diagonals shorter than its edges. In the first edition of Crystal Data the Delaunay reduced cell was used as the standard cell. In 20% of the cases reported, the cell has angles that differ from 90° by more than 30° ; in some cases the difference is as much as 60° , and there seems to be no theoretical reason why the Delaunay method could not produce a cell with angles approaching 180° . When the Bravais-reduced cell is tri-acute, the Delaunay reduction can be applied to the reciprocal cell (Blanton, 1937), though Buerger has shown that the shortest a^*, b^*, c^* in reciprocal space do not necessarily correspond to the shortest a, b, c in direct space.

The Dirichlet reduction will always produce the cell with the three shortest translations. Given any triplet, a, b, c , it forms the reduced triplet by using multiples of each vector in turn in attempts to shorten the other two edges. The steps can be performed in any

order and they are repeated until there is no further change. The order convention adopted ($a < b < c$ or $a < c < b$) has no effect on the reduction.

In the second edition of Crystal Data the Bravais-reduced cell is used as the standard cell in all cases; but the condition imposed is that α and β be non-acute. In the case of a tri-acute parallelepiped, this corresponds to a shift of origin to the "right-handed" corner whose angles are α', β', γ .

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