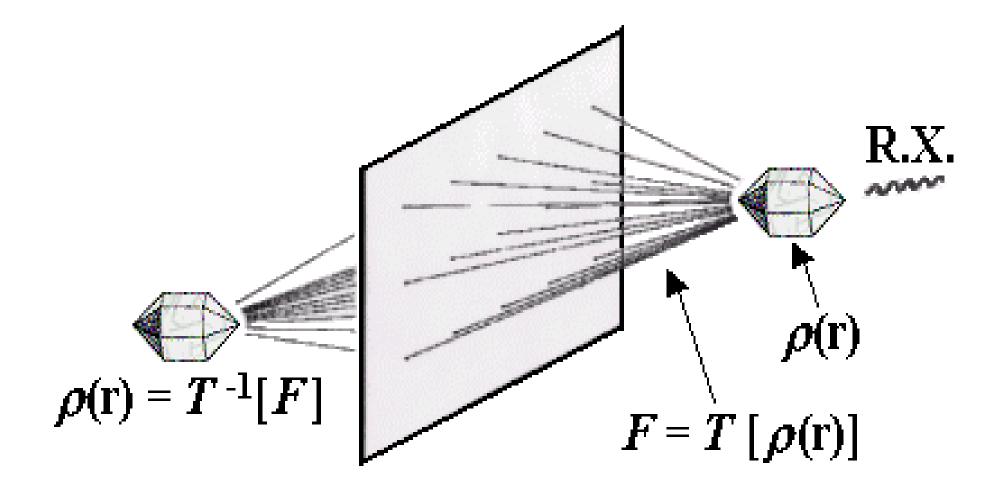
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Direct Methods

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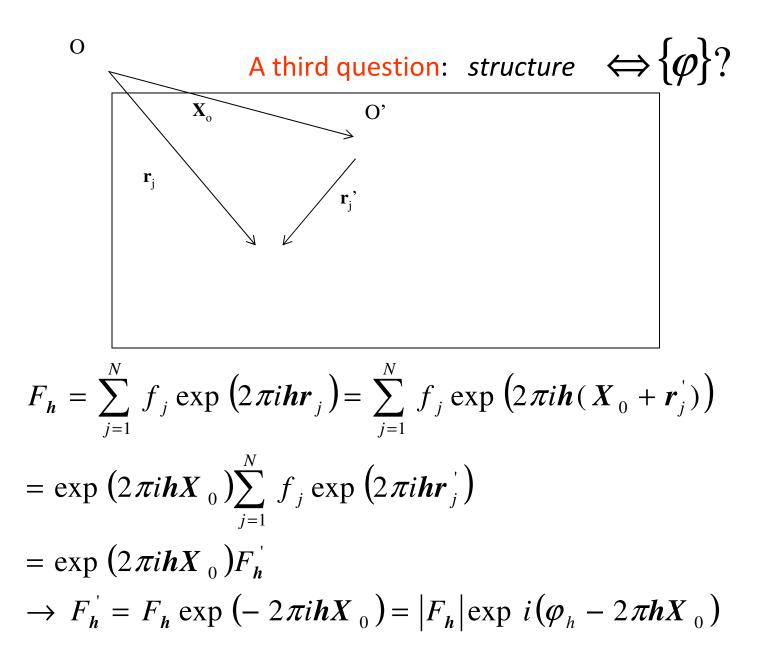


Let us answer the following questions:

crystal structure
$$\Rightarrow \{|F|^2\}$$
?
 $\{|F|^2\} \Rightarrow crystal structure ?$

$$|F_{h}|^{2} = \sum_{j=1}^{N} f_{j} \exp(2\pi i h r_{j}) \sum_{j=1}^{N} f_{j} \exp(-2\pi i h r_{j})$$
$$= \sum_{i,j=1}^{N} f_{i} f_{j} \exp[2\pi i h (r_{i} - r_{j})]$$
$$= \sum_{j=1}^{N} f_{j}^{2} + \sum_{i \neq j=1}^{N} f_{i} f_{j} \exp[2\pi i h (r_{i} - r_{j})]$$

As a consequence : $\{F|^2\} \Leftrightarrow \rho(r)$



A fourth basic question

How can we derive the phases from the diffraction moduli ? This seems contradictory: indeed

The phase values depend on the origin chosen by the user, the moduli are independent of the user.

The moduli are *structure invariants*, the phases are not structure invariants.

Evidently, from the moduli we can derive information only on those combinations of phases (if they exist) which are structure invariants.

The simplest invariant : *the triplet invariant*

Use the relation

$$F'_{h} = F_{h} \exp(-2\pi i h X_{0})$$

to check that the invariant $F_h F_k F_{-h-k}$ does not depend on the origin.

$$\begin{aligned} F_{h}F_{k}F_{-h-k} &= |F_{h}| \exp i(\phi_{h} - 2\pi hX_{0})|F_{k}| \exp i(\phi_{h} - 2\pi kX_{0}) \\ &|F_{-h-k}| \exp i[\varphi_{-h-k} + 2\pi (h+k)X_{0}] \\ &= |F_{h}||F_{k}||F_{-h-k}| \exp i(\phi_{h} + \phi_{k} + \varphi_{-h-k}) \\ &\text{The sum } (\phi_{h} + \phi_{k} + \phi_{-h-k}) \text{ is called triplet phase invariant} \end{aligned}$$

Structure invariants

Any invariant satisfies the condition that the sum of the indices is zero:

doublet invariant : $F_h F_{-h} = |F_h|^2$

triplet invariant : $F_h F_k F_{-h-k}$

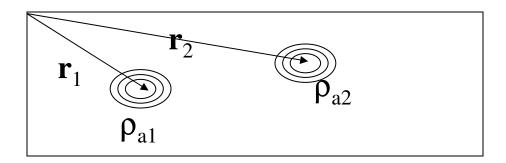
quartet invariant : $F_h F_k F_l F_{-h-k-l}$

quintet invariant : $F_h F_k F_l F_m F_{-h-k-l-m}$

.

The prior information we can use for deriving the phase estimates may be so summarised:

1) *atomicity*: the electron density is concentrated in atoms:



$$\rho(\mathbf{r}) = \sum_{j=1}^{N} \rho_{aj} \left(\mathbf{r} - \mathbf{r}_{j} \right)$$
2) **positivity** of the electron density:

$$\rho(\mathbf{r}) > 0 \implies f > 0$$

3) uniform distribution of the atoms in the unit cell.

The Wilson statistics

• Under the above conditions Wilson (1942,1949) derived the structure factor statistics. The main results where:

$$\langle |F_{\mathbf{h}}|^2 \rangle = \sum_{j=1}^N f_j^2$$
 (1)

- Eq.(1) is :
- a) resolution dependent (f_i varies with θ),
- b) temperature dependent: $f_j = f_j^0 \exp(-B_j \sin^2 \theta / \lambda^2)$
- From eq.(1) the concept of normalized structure factor arises:

$$E_{\mathbf{h}} = F_{\mathbf{h}} / (\sum_{j=1}^{N} f_j^2)^{1/2}$$

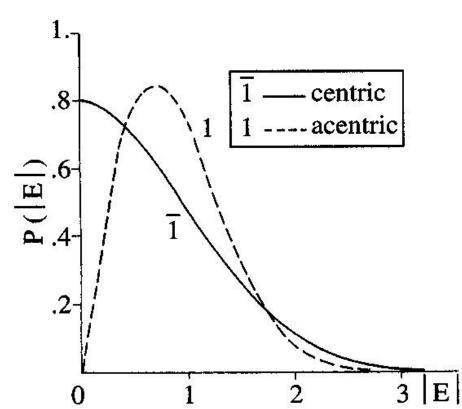
The Wilson Statistics

• |E|-distributions:

$$P_{1}(|E|) = 2|E|\exp(-|E|^{2})$$

$$P_{\overline{1}}(|E|) = \sqrt{\frac{2}{\pi}}\exp(-|E|^{2}/2)$$
and
$$|E|^{2} >= 1$$

in both the cases. The statistics may be used to evaluate the average themel factor and the absolute scale factor.

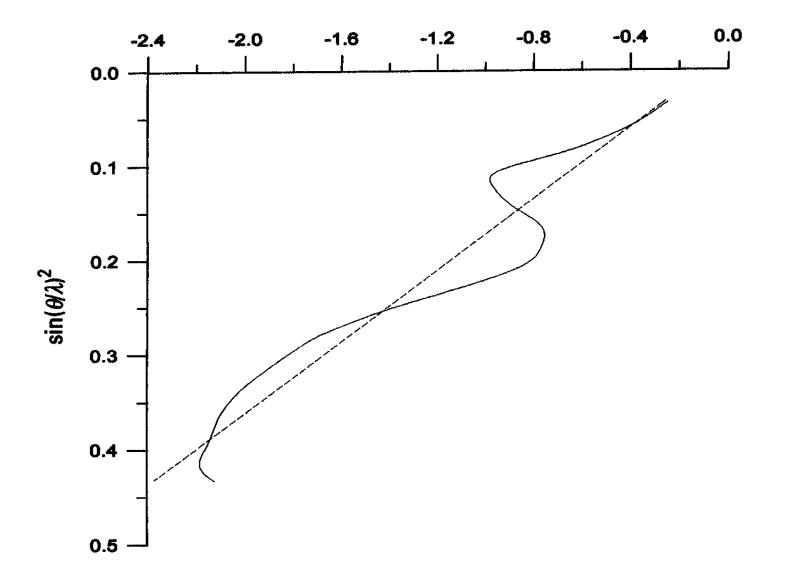


The Wilson plot $F_{\mathbf{h}} = \sum_{j=1}^{N} f_j \exp 2\pi i \mathbf{h} \mathbf{r}_j = \sum f_j^0 \exp\left(-B_j \frac{\sin^2 \theta}{\lambda^2}\right) \exp 2\pi i \mathbf{h} \mathbf{r}_j$ $\sum \exp\left(-B\frac{\sin^2\theta}{\lambda^2}\right)\sum f_j^0 \exp 2\pi i\mathbf{hr}_j$

$$|F_{h}|_{obs}^{2} = K |F_{h}|^{2} = K |F_{h}^{0}|^{2} \exp\left(-2Bs^{2}\right)$$

$$(F_{h}|_{obs}^{2} \ge K < |F_{h}^{0}|^{2} \ge \exp(-2Bs^{2}) = K\Sigma_{s}^{0} \exp(-2Bs^{2})$$

$$\ln \left(\frac{<\left|F_{h}\right|_{obs}^{2}}{\Sigma_{s}^{0}} \right) = \ln K - 2B_{s}^{2}$$



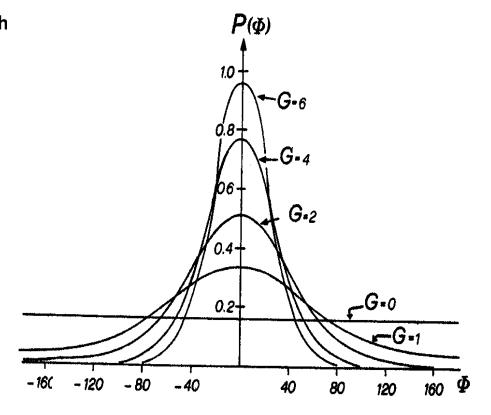
The Cochran formula

$$\Phi_{h,k} = \varphi_h + \varphi_k + \varphi_{-h-k} = \varphi_h + \varphi_k - \varphi_h$$
$$P(\Phi_{hk}) \approx [2\pi I_0]^{-1} \exp(G \cos \Phi_{hk})$$

where G = 2 | $E_h E_k E_{h+k}$ |/N^{1/2}

Accordingly:

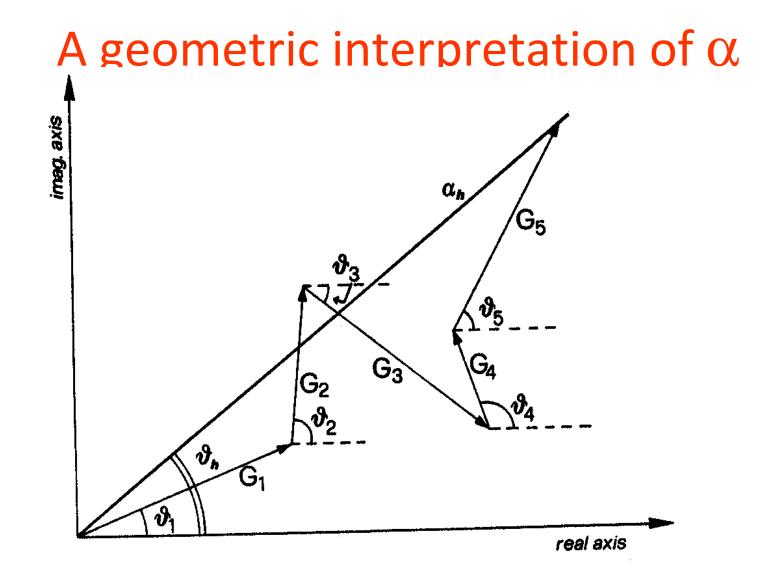
- $\phi_{h} \approx \phi_{k} \phi_{h-k}$
- $\phi_{\textbf{h}} + \phi_{\textbf{k}} \phi_{\textbf{h}+\textbf{k}} \approx 0 \quad \propto \qquad \quad G = 2 \ \mid \textbf{E}_{\textbf{h}} \ \textbf{E}_{\textbf{k}} \ \textbf{E}_{\textbf{h}+\textbf{k}} \ \mid / \textbf{N}^{1/2}$ $\varphi_h - \varphi_k - \varphi_{h-k} \approx 0 \propto G = 2 | E_h E_k E_{h-k} |/N^{1/2}$ \propto G = 2 | E_h E_k E_{h-k} |/N^{1/2}



The tangent formula

A reflection can enter into several triplets. Accordingly
$$\begin{split} \phi_h &\approx \phi_{k1} + \phi_{h-k1} = \theta_1 \quad \text{with } \mathsf{P}_1(\phi_h) \quad \propto \quad \mathsf{G}_1 = 2 \mid \mathsf{E}_h \mathsf{E}_{k1} \mathsf{E}_{h-k1} \mid /\mathsf{N}^{1/2} \\ \phi_h &\approx \phi_{k2} + \phi_{h-k2} = \theta_2 \quad \text{with } \mathsf{P}_2(\phi_h) \quad \propto \quad \mathsf{G}_2 = 2 \mid \mathsf{E}_h \mathsf{E}_{k2} \mathsf{E}_{h-k2} \mid /\mathsf{N}^{1/2} \\ \hline \\ \phi_h &\approx \phi_{kn} + \phi_{h-kn} = \theta_n \quad \text{with } \mathsf{P}_n(\phi_h) \quad \propto \quad \mathsf{G}_n = 2 \mid \mathsf{E}_h \mathsf{E}_{kn} \mathsf{E}_{h-kn} \mid /\mathsf{N}^{1/2} \\ \hline \\ \mathsf{Then} \\ \mathsf{P}(\phi_h) &\approx \quad \prod_j \mathsf{P}_j(\phi_h) \approx \mathsf{L}^{-1} \prod_j \exp\left[\mathsf{G}_j \cos\left(\phi_h - \theta_j\right)\right] \\ &= \mathsf{L}^{-1} \exp\left[\alpha \cos\left(\phi_h - \theta_h\right)\right] \\ \end{split}$$

$$\tan \theta_h = \frac{\sum G_j \sin \theta_j}{\sum G_j \cos \theta_j} = \frac{T}{B}, \qquad \alpha_h = \left(T^2 + B^2\right)^{1/2}$$



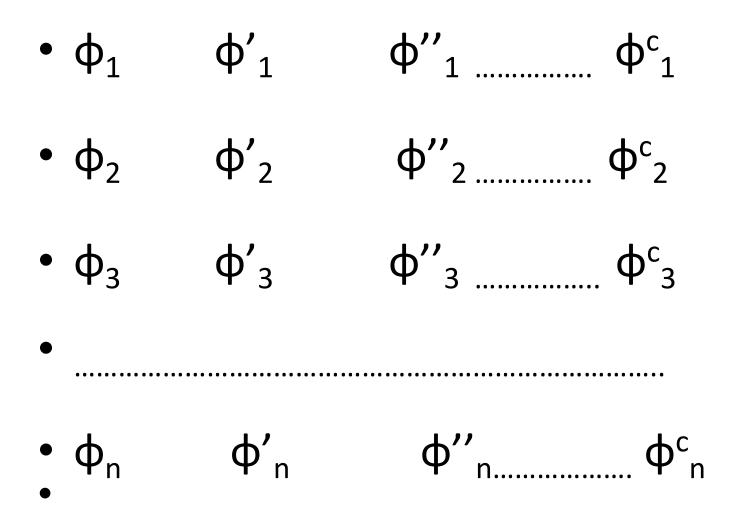
The random starting approach

To apply the tangent formula we need to know one or more pairs ($\phi_k + \phi_{h-k}$). Where to find such an information?

The most simple approach is the *random starting approach*. Random phases are associated to a *chosen* set of reflections. The tangent formula should drive these phases to the correct values. The procedure is cyclic (up to *convergence*).

How to recognize the correct solution? *Figures of merit* can or cannot be applied

Tangent cycles



• Ab initio phasing

- SIR2009 is able to solve
- -small size structures (up to 80 atoms in the a.u.);
- -medium-size structures (up to 200);
- -large size (no upper limit)
- It uses
- Patterson deconvolution techniques
- (multiple implication transformations)
- as well as

Direct methods

- to obtain a starting set of phases. They are extended and refined via
- electron density modification techniques

• Direct methods limits for proteins:

- 1) the large size (proteins range from 300 atoms in the asymmetric unit to several thousands). The G factor in the Cochran formula are very small.
- 2) data resolution

- To overcome the limits one is obliged to :
- -increase the number of direct methods trials.
 The cost to pay concerns the computing time.
- improve and extend the the poor phases available by DM by exploiting some specific features of the proteins (e.g., the solvent, etc.).

About the data resolution limit

- Atomic resolution at length was considered a necessary (and not sufficient) condition for **ab initio phasing** (Sheldrick rule), condition relaxed later on (up to 1.2 Å). If it is not satisfied:
- - the atomicity condition is violated;
- the number of reliable triplet invariants exploitable by the tangent procedure is small.
- - Patterson and EDM procedures are less effective;
- - the small ratio
- *number of observations/ number of parameters*
- make least squares unreliable.