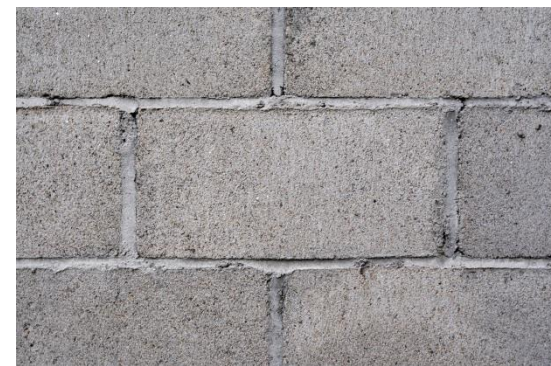


Quantitative Phase Analysis with Diffraction Methods

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Books

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- *Introduction to X-Ray Powder Diffractometry*, Jenkins & Snyder, Publisher, Wiley 1996.
- *Quantitative X-Ray Diffractometry*, Zevin & Kimmel, Springer-Verlag, New York, 1996.
- *Industrial of X-Ray Diffraction*, Chung & Smith, Marcel Dekker, New York, 2000.
- *Structure Determination from Powder Diffraction Data*, David, Shankland, McCusker & Baerlocher Editors, Oxford University Press, New York, 2002.
- *Fundamentals of Powder Diffraction and Structural Characterization of Materials*, Pecharsky & Zavalij, 2009, Springer.
- *Powder Diffraction – Theory and Practice*, Dinnebier & Billinge Editors, 2009, RSC Pub (QPA: Ch.11).
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- B.H. O'Connor & M.D. Raven, *Application of the Rietveld refinement procedure in assaying powdered mixtures*, Powder Diffr. 1988. **3**, 2-6
- D.L. Bish & S.A. Howard, *Quantitative phase analysis using the Rietveld method*, J. Appl. Cryst. 1988. **21**, 86-91.
- R.J. Hill & C. J. Howard, *Quantitative phase analysis from neutron powder diffraction data using the Rietveld method*, J. Appl. Cryst. 1987. **20**, 467-474.
- N.V.Y. Scarlett & I. C. Madsen, *Quantification of phases with Partial Or No Known Crystal Structure*, Powder Diffr. 2006. **21**, 278-284.
- C. Giannini, A. Guagliardi & Millini *Quantitative phase analysis by combining the Rietveld and the whole-pattern decomposition methods*, J.Appl.Cryst (2002). **35**, 481

Outlook

- I. Defining the QPA problem
- II. Mathematical background
- III. Single-peak (or single-line) QPA methods
- IV. Whole patterns QPA methods → Rietveld and Rietveld-like methods

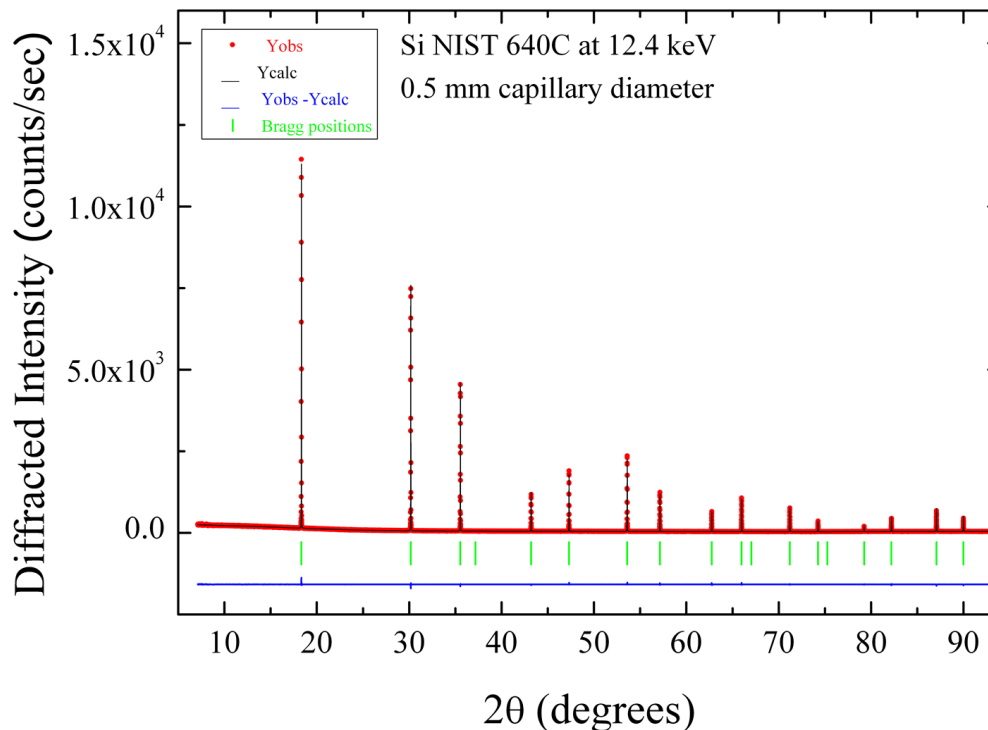
Qualitative *versus* Quantitative phase analysis

Principal use of powder diffraction technique is the identification of crystalline or disordered structures (or phases)

Why?

A powder diffraction pattern is DIRECTLY produced by the structure of the component phases and can, therefore, fingerprint them

What do we learn from a powder pattern of a crystalline structure?

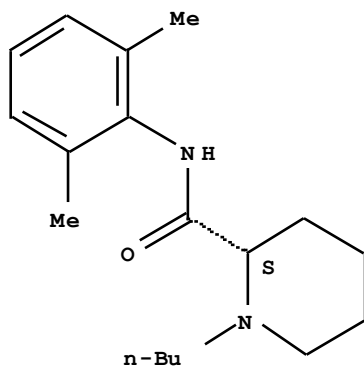


- Position of the diffracted peaks → size and dimension of the unit cell
- Intensity ratios of the diffracted peaks → type and location of atoms in the unit cell
- Full Width at Half Maximum (FWHM) of the diffracted peaks → intrinsic properties of the materials (i.e. microstructural analysis)

Polymorphism of drugs

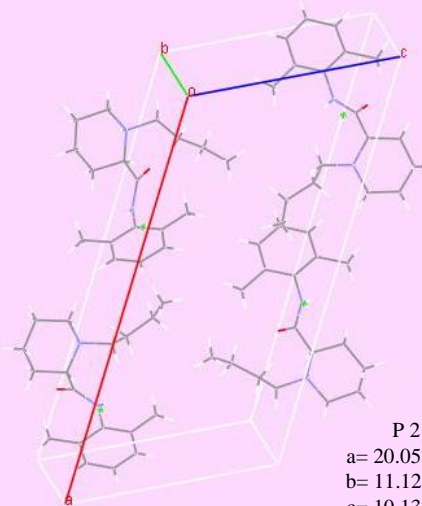
Polymorphism is the ability of substances with **identical chemical composition** to crystallize in solid state phases according to different arrangements or conformations of the basic molecule(s) in the crystal lattice

Example of
Bupivacaine Hydrochloride



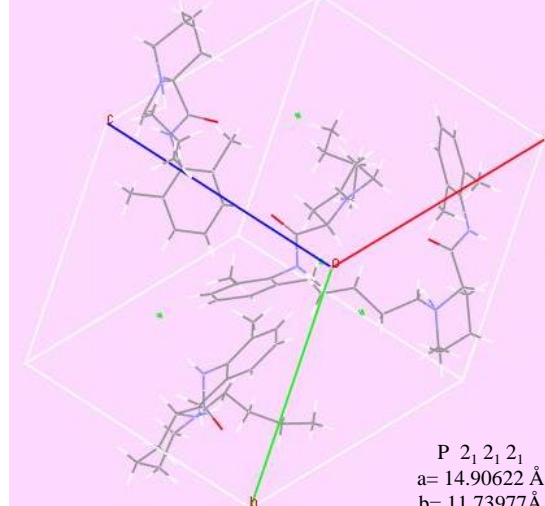
● HCl

Form B at 112 °C, monoclinic



P 2₁
a= 20.05795 Å
b= 11.12509 Å
c= 10.13290 Å
β= 116.18377 °

Form D at 20 °C, orthorhombic



P 2₁ 2₁ 2₁
a= 14.90622 Å
b= 11.73977 Å
c= 11.08386 Å

Gozzo, Masciocchi, Griesser, Niederwanger, 2010

Forms B and D share the same chemical composition, but have different solid forms
→ They are different polymorphs!

Quantitative Phase Analysis (QPA)

QPA refers to the ability of quantitatively state the abundance of the different phases that constitute a mixture.

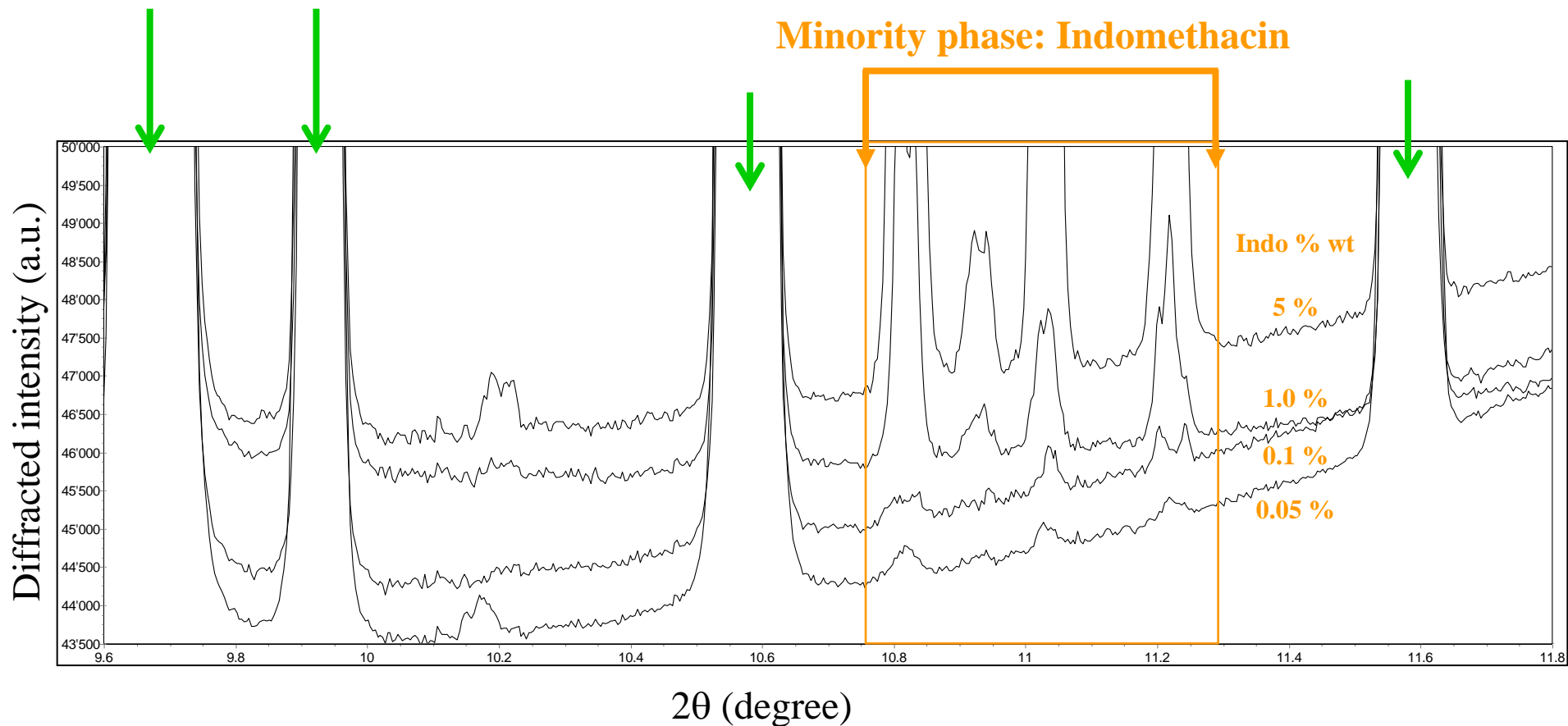
Why is this relevant?

- ❑ **Polymorphic purity**: detect and quantify unwanted polymorphic forms in both drug substance and drug product
 - Level of Detection (LoD)
 - Level of Quantitation (LoQ)
- ❑ Assess the **polymorphic composition** in drug substance and product
- ❑ In formulated materials, the **API/excipients relative proportion** is important and needs to be kept under control
- ❑ **Degree of Crystallinity** in amorphous/crystalline mixtures

QPA of a binary API physical mixtures with fast SR-XRPD

Majority phase (intensity up to 1.5 M counts): Haloperidol

Minority phase: Indomethacin




QPA analytical methods

Several are the analytical methods used to obtain quantitative phase related information:

- Based on chemical composition (so-called *normative calculation*)
- Based on properties specific to the phases of interest (e.g. magnetism, selective dissolution, density)
- Spectroscopic methods (e.g. Raman and Infrared spectroscopy, Mass spectroscopy, Nuclear Magnetic Resonance spectroscopy)
- Thermal Methods (e.g. Differential Scanning Calorimetry, ThermoGravimetric Analysis)
- Diffraction Methods → XRPD

Direct method

Diffraction information is directly produced by the crystal structure of the component phases in the mixture

QPA  determining the contribution (typically in % weight) of each component phase in a mixture

DIFFRACTION-BASED QPA METHODS

Single-peak methods:

- Intensity ratio $I_{\text{unknown}}/I_{\text{standard}}$ of one or more reflections
- No need of structural information but prone to systematic errors (e.g. caused by preferential orientation and peak overlapping)
- Need *ad-hoc* mixtures for calibration curves

Whole-pattern methods:

- Use of full diffraction patterns
- Minimization of systematic errors (e.g. due to peak overlap)
- Preferential Orientation (PO) can be modeled
- Accuracy close to X-Ray fluorescence elemental analysis, with the advantage of being sensitive to structural differences → direct QPA of polymorphs

DIFFRACTION-BASED QPA METHODS

Single-peak methods:

(adapted from Cullity, *Elements of X-Ray Diffraction*)

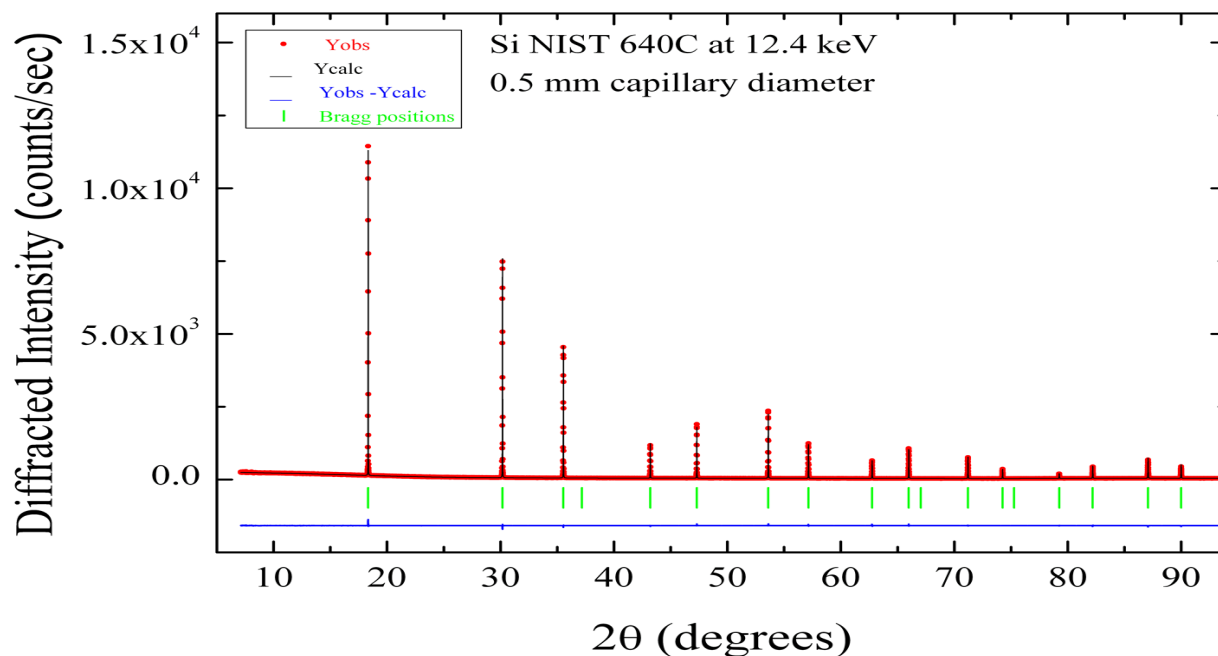
- Intensity ratio $I_{\text{unknown}}/I_{\text{standard}}$ of one or more reflections
- No need of structural information but prone to systematic errors (e.g. caused by preferential orientation and peak overlapping)
- Need *ad-hoc* mixtures for calibration curves

Rietveld-based methods:

(adapted from Madsen & Scarlett in *Powder Diffraction-Theory and Practice*)

- Use of full diffraction patterns
- Minimization of systematic errors (e.g. due to peak overlap)
- Preferential Orientation (PO) can be modeled
- Accuracy close to X-Ray fluorescence elemental analysis, with the advantage of being sensitive to structural differences → direct QPA of polymorphs

QPA with diffraction methods: math background



The diffracted intensity distribution is defined by:

Structural factors

→ crystal structure

Specimen factors

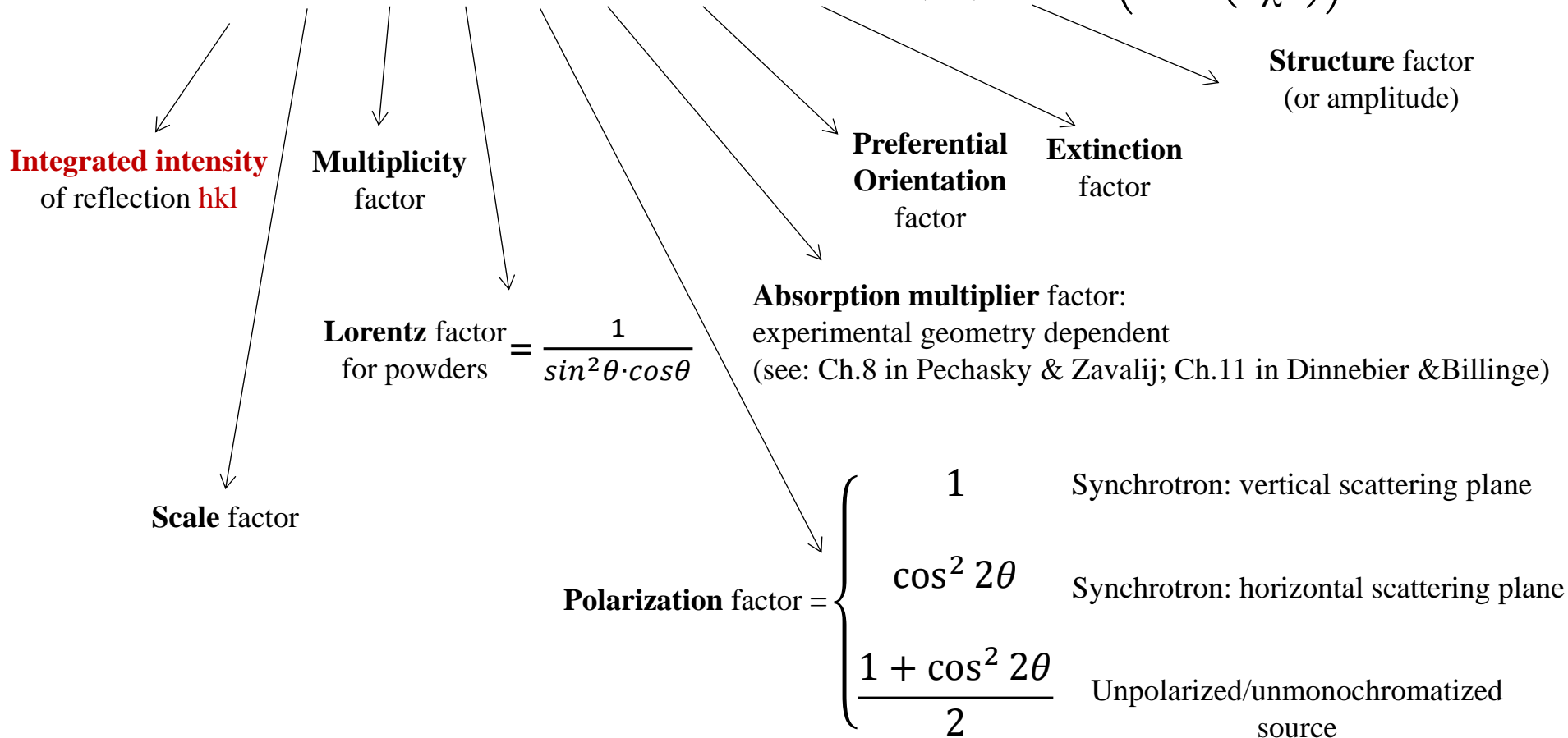
→ preferential orientation, grain size, shape and distribution, microstructure

Instrumental factors

→ properties of radiation, optics geometry, properties of detectors, slits and/or monochromator

For a powder diffraction pattern of a pure phase we can write:

$$I_{hkl} = K \cdot M_{hkl} \cdot L_{\theta} \cdot P_{\theta} \cdot A_{\theta} \cdot PO_{hkl} \cdot E_{hkl} \cdot |F_{(hkl)}|^2 \cdot \exp\left(-2B \left(\frac{\sin \theta}{\lambda}\right)\right)$$



For a powder diffraction pattern of a pure phase we can write:

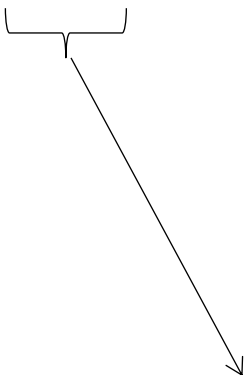
$$I_{hkl} = K \cdot M_{hkl} \cdot L_{\theta} \cdot P_{\theta} \cdot A_{\theta} \cdot PO_{hkl} \cdot E_{hkl} \cdot |F_{(hkl)}|^2 \cdot \exp\left(-2B \left(\frac{\sin \theta}{\lambda}\right)\right)$$

Lorentz factor
for powders = $\frac{1}{\sin^2 \theta \cdot \cos \theta}$

Polarization factor = $\begin{cases} 1 & \text{Synchrotron: vertical scattering plane} \\ \cos^2 2\theta & \text{Synchrotron: horizontal scattering plane} \\ \frac{1 + \cos^2 2\theta}{2} & \text{Unpolarized/unmonochromatized source} \end{cases}$

For a powder diffraction pattern of a pure phase we can write:

$$I_{hkl} = K \cdot M_{hkl} \cdot \underbrace{L_{\theta} \cdot P_{\theta}} \cdot A_{\theta} \cdot PO_{hkl} \cdot E_{hkl} \cdot |F_{(hkl)}|^2 \cdot \exp\left(-2B \left(\frac{\sin \theta}{\lambda}\right)\right)$$



**Lorentz-
Polarization factor** = $\frac{1 + \cos^2 2\theta \cos^2 2\theta_M}{\sin^2 \theta \cdot \cos \theta}$
 for powders

Lab diffractometer:

$2\theta_M$ is the Bragg angle of the reflection from a monochromator, $2\theta_M=0$ for unpolarized unmonochromatized source, $2\theta_M \neq 0$ (e.g. 26.5° graphite mono + CuK α radiation)

Synchrotron radiation:

$2\theta_M$ is the angle between the scattering direction (where we place our detector!) and the direction of acceleration of the electron (e.g. the direction of the electric field of the synchrotron e.m. radiation). In the vertical plane this is always 90° .

For a powder diffraction pattern of a **pure phase** we can write:

$$I_{hkl} = K \cdot M_{hkl} \cdot L_{\theta} \cdot P_{\theta} \cdot A_{\theta} \cdot PO_{hkl} \cdot E_{hkl} \cdot |F_{(hkl)}|^2 \cdot \exp\left(-2B \left(\frac{\sin \theta}{\lambda}\right)\right)$$



negligible

Under the hypothesis of **A**, **PO** and **E negligible**, transmission (Debye-Scherrer) geometry, synchrotron radiation with vertical diffraction plane (as at the SLS-MS-PD), with powders loaded in capillaries and a 1D display detector (e.g. Mythen II):

$$I_{hkl} = \left[\frac{I_0 \lambda^3}{32 \pi r} \cdot \frac{\sigma e^4}{m_e^2 c^4} \right] \cdot \left[\frac{M_{hkl}}{V^2} \cdot |F_{(hkl)}|^2 \cdot \left(\frac{1}{\sin^2 \theta \cdot \cos \theta} \right) \right] \cdot \exp\left(-2B \left(\frac{\sin \theta}{\lambda}\right)\right) \cdot \left[\frac{1}{\mu} \right]$$

I_0	incident beam intensity	$\mu = \rho \mu^*$	linear absorption (attenuation) coefficient of the pure phase
λ	photon beam wavelength		
σ	cross sectional area of incident beam	with ρ and μ^*	density of the pure phase and the mass absorption coefficient
e	charge of the electron		
m_e	mass of the electron		
c	speed of light	$\exp(-2B \left(\frac{\sin \theta}{\lambda}\right))$	thermal factor, B is the mean atomic displacement parameter (ADP)
V	unit cell volume		

For a powder diffraction pattern of a **mixture** (e.g. binary $\alpha+\beta$ mixture), for a hkl intensity line of phase α , we can write:

$$I_{(hkl)\alpha} = \left[\frac{I_0 \lambda^3}{32 \pi r} \cdot \frac{\sigma e^4}{m_e^2 c^4} \right] \cdot \left[\frac{M_{hkl}}{V_\alpha^2} \cdot |F_{hkl}|_\alpha^2 \left(\frac{1}{\sin^2 \theta \cdot \cos \theta} \right) \right] \cdot \exp \left(-2B_\alpha \left(\frac{\sin \theta}{\lambda} \right) \right) \cdot \left[\frac{c_\alpha}{\mu_m} \right]$$

I_0	incident beam intensity	c_α	volume fraction of phase α
λ	photon beam wavelength	$\mu_m = \rho_m \cdot \mu_m^*$	linear absorption (attenuation) coefficient of the mixture
σ	cross sectional area of incident beam		
e	charge of the electron		
m_e	mass of the electron	with ρ_m and μ_m^*	density of the mixture and the mass absorption coefficient
c	speed of light		
r	distance scattering electron-detector	$\exp(-2B \left(\frac{\sin \theta}{\lambda} \right))$	thermal factor, with B mean Atomic Displacement Parameter (ADP)
V_α	unit cell volume of phase α		

And similarly, for a h'k'l' line of phase β , we can write:

$$I_{(h'k'l')\beta} = \left[\frac{I_0 \lambda^3}{32 \pi r} \cdot \frac{\sigma e^4}{m_e^2 c^4} \right] \cdot \left[\frac{M_{hkl}}{V_\beta^2} \cdot |F_{(h'k'l')}|_\beta^2 \cdot \left(\frac{1}{\sin^2 \theta \cdot \cos \theta} \right) \right] \cdot \exp \left(-2B_\beta \left(\frac{\sin \theta}{\lambda} \right) \right) \cdot \left[\frac{c_\beta}{\mu_m} \right]$$

For a powder diffraction pattern of a mixture (e.g. binary $\alpha+\beta$ mixture), for a hkl intensity line of phase α , we can write:

$$I_{(hkl)\alpha} = \left[\frac{I_0 \lambda^3}{32 \pi r} \cdot \frac{\sigma e^4}{m_e^2 c^4} \right] \cdot \left[\frac{M_{hkl}}{V_\alpha^2} \cdot |F_{hkl}|_\alpha^2 \left(\frac{1}{\sin^2 \theta \cdot \cos \theta} \right) \right] \cdot \exp \left(-2B_\alpha \left(\frac{\sin \theta}{\lambda} \right) \right) \cdot \left[\frac{c_\alpha}{\mu_m} \right]$$

All factors are constant and independent of the concentration of the α phase with the exception of c_α and μ_m



$$I_{(hkl)\alpha} = \frac{K_{1,\alpha} \cdot c_\alpha}{\mu_m}$$

and equivalently:

$$I_{(hkl)\beta} = \frac{K_{1,\beta} \cdot c_\beta}{\mu_m}$$

The simplified expression: $I_{(hkl)\alpha} = \frac{K_{1,\alpha} \cdot c_{\alpha}}{\mu_m}$ can also be written in terms of weight

fractions w_{α} (and w_{β}): $I_{(hkl)\alpha} = \frac{K_{1,\alpha} \cdot w_{\alpha} \cdot \rho_m}{\mu_m \cdot \rho_{\alpha}} = \frac{K_{1,\alpha} \cdot w_{\alpha}}{\rho_{\alpha} \frac{\mu_m}{\rho_m}} = \frac{K_{1,\alpha} \cdot w_{\alpha}}{\rho_{\alpha} \mu_m^*}$ with ρ_{α} the density

of phase α and μ_m^* the mass absorption coefficient ($= \frac{\mu_m}{\rho_m}$).

DEMO:

Assuming a unit volume of mixture ($V_m=1$): ρ_m =density of mixture=weight of mixture $\rightarrow w_{\alpha}\rho_m$ and $w_{\beta}\rho_m$ represent the weights of the α and β contents in our binary mixture with w_{α} and w_{β} the weight fractions $\rightarrow \rho_{\alpha} = \frac{w_{\alpha}\rho_m}{c_{\alpha}} \rightarrow c_{\alpha} = \frac{w_{\alpha}\rho_m}{\rho_{\alpha}}$ and equivalently $c_{\beta} = \frac{w_{\beta}\rho_m}{\rho_{\beta}} \rightarrow \frac{c_{\alpha}}{\mu_m} = \frac{w_{\alpha}\rho_m}{\rho_{\alpha}\mu_m} = \frac{w_{\alpha}}{\rho_{\alpha}} \cdot \frac{1}{\mu_m^*}$ with μ_m linear absorption (or attenuation) coefficient and μ_m^* mass absorption coefficient.

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$\frac{w_{\beta}\rho_m}{\rho_{\beta}} \rightarrow \frac{c_{\alpha}}{\mu_m} = \frac{w_{\alpha}\rho_m}{\rho_{\alpha}\mu_m} = \frac{w_{\alpha}}{\rho_{\alpha}} \cdot \frac{1}{\mu_m^*}$ with μ_m linear absorption (or attenuation) coefficient and μ_m^* mass absorption coefficient.

The simplified expression: $I_{(hkl)\alpha} = \frac{K_{1,\alpha} \cdot c_\alpha}{\mu_m}$ can also be written in terms of weight

fractions w_α (and w_β): $I_{(hkl)\alpha} = \frac{K_{1,\alpha} \cdot w_\alpha \cdot \rho_m}{\mu_m \cdot \rho_\alpha} = \frac{K_{1,\alpha} \cdot w_\alpha}{\rho_\alpha \frac{\mu_m}{\rho_m}} = \frac{K_{1,\alpha} \cdot w_\alpha}{\rho_\alpha \mu_m^*}$ with ρ_α the density

of phase α and μ_m^* the mass absorption coefficient ($= \frac{\mu_m}{\rho_m}$).

If we can access a powder sample of pure phase α : $I_{(hkl)\alpha,pure} = \frac{K_{1,\alpha}}{\rho_\alpha \mu_\alpha^*} = \frac{K_{1,\alpha}}{\mu_\alpha}$

and we can write: $\frac{I_{(hkl)\alpha}}{I_{(hkl)\alpha,pure}} = \frac{K_{1,\alpha} \cdot w_\alpha}{\rho_\alpha \frac{\mu_m}{\rho_m}} \cdot \frac{\mu_\alpha}{K_{1,\alpha}} = \frac{w_\alpha (\mu_\alpha / \rho_\alpha)}{(\mu_m / \rho_m)} = \frac{w_\alpha (\mu_\alpha / \rho_\alpha)}{w_\alpha (\mu_\alpha / \rho_\alpha) + \mu_\beta / \rho_\beta}$

[demo p.389-390 Cullity]

$$\frac{I_{(hkl)\alpha}}{I_{(hkl)\alpha,pure}} = \frac{w_{\alpha}(\mu_{\alpha}/\rho_{\alpha})}{w_{\alpha}(\mu_{\alpha}/\rho_{\alpha}) + \mu_{\beta}/\rho_{\beta}}$$

QPA of a binary mixture can, therefore be performed provided that we can access:

- the mass absorption coefficients of the two phases (if not, a calibration curve can be prepared using mixtures of known composition)
- one pure phase (or a mixture with a known amount of that phase)
- No need of structural information, $K_{1,\alpha}$ cancels out

What can we observe?

$$\frac{I_{(hkl)\alpha}}{I_{(hkl)\alpha,pure}} = \frac{w_{\alpha}(\mu_{\alpha}/\rho_{\alpha})}{w_{\alpha}(\mu_{\alpha}/\rho_{\alpha} - \mu_{\beta}/\rho_{\beta}) + \mu_{\beta}/\rho_{\beta}}$$

- The intensity of a particular diffraction line depends on the mass absorption coefficient of the other phase
- For binary mixtures of phases with the same mass absorption coefficient: $\frac{I_{(hkl)\alpha}}{I_{(hkl)\alpha,pure}} = w_{\alpha}$

The binary mixture case that we have worked out together is an example of the so-called **single-line** or **single-peak methods of QPA**, for which *the measurement of the weight fraction of phase in a mixture depends on the measurement of the ratio of the intensity of a diffraction line from that phase to the intensity of some standard reference line!*

In the case discussed, the reference standard is the pure phase α !

DIRECT COMPARISON METHOD

INTERNAL STANDARD METHOD

DIRECT COMPARISON METHOD

Let us again consider a $\alpha+\beta$ binary mixture:

$$I_{(hkl)\alpha} = \frac{K_{1,\alpha} \cdot c_\alpha}{\mu_m} = \frac{K_{1,\alpha} \cdot w_\alpha}{\rho_\alpha \mu_m^*} \qquad I_{(h'k'l')\beta} = \frac{K_{1,\beta} \cdot c_\beta}{\mu_m} = \frac{K_{1,\beta} \cdot w_\beta}{\rho_\beta \mu_m^*}$$

Let us separate in $K_{1,\alpha}$ the phase-dependent from the phase-independent part:

$$K_{1,\alpha} = K_2 \cdot R_{1,\alpha}$$

K_2 is a constant independent of the kind and amount of diffracting substance; R depends on θ , hkl, kind of substance

$$I_{(hkl)\alpha} = \frac{K_2 \cdot R_\alpha \cdot c_\alpha}{\mu_m} \qquad \longrightarrow \qquad \frac{I_{(hkl)\alpha}}{I_{(h'k'l')\beta}} = \frac{R_\alpha \cdot c_\alpha}{R_\beta \cdot c_\beta} \qquad \longrightarrow \qquad \frac{c_\alpha}{c_\beta} = \frac{I_{(hkl)\beta}}{I_{(h'k'l')\alpha}} \cdot \frac{R_\beta}{R_\alpha}$$

$$I_{(h'k'l')\beta} = \frac{K_2 \cdot R_\beta \cdot c_\beta}{\mu_m}$$

Details here: $K_2 = \frac{I_0 \lambda^3}{32 \pi r} \cdot \frac{\sigma e^4}{m_e^2 c^4}$ and $R_\alpha = \left[\frac{M_{hkl}}{V_\alpha^2} \cdot |F_{(hkl)}|_\alpha^2 \cdot \left(\frac{1 + \cos^2 2\theta}{2} \right) \right] \cdot \exp \left(-2B_\alpha \left(\frac{\sin \theta}{\lambda} \right) \right)$

DIRECT COMPARISON METHOD

Let us again consider a $\alpha+\beta$ binary mixture:

$$I_{(hkl)\alpha} = \frac{K_{1,\alpha} \cdot c_\alpha}{\mu_m} = \frac{K_{1,\alpha} \cdot w_\alpha}{\rho_\alpha \mu_m^*} \qquad I_{(h'k'l')\beta} = \frac{K_{1,\beta} \cdot c_\beta}{\mu_m} = \frac{K_{1,\beta} \cdot w_\beta}{\rho_\beta \mu_m^*}$$

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K_2 is a constant independent of the kind and amount of diffracting substance; R depends on θ , hkl, kind of substance

$$I_{(hkl)\alpha} = \frac{K_2 \cdot R_\alpha \cdot c_\alpha}{\mu_m} \qquad \longrightarrow \qquad \frac{I_{(hkl)\alpha}}{I_{(h'k'l')\beta}} = \frac{R_\alpha \cdot c_\alpha}{R_\beta \cdot c_\beta} \qquad \longrightarrow \qquad \frac{w_\alpha}{w_\beta} = \frac{I_{(hkl)\beta}}{I_{(h'k'l')\alpha}} \cdot \frac{\rho_\alpha \cdot R_\beta}{\rho_\beta \cdot R_\alpha}$$

$$I_{(h'k'l')\beta} = \frac{K_2 \cdot R_\beta \cdot c_\beta}{\mu_m}$$

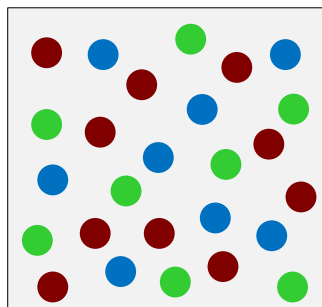
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INTERNAL STANDARD METHOD

A diffraction line from the phase that we need to quantify in a given mixture (e.g. $I_{hkl,\alpha}$) is compared with a line from a standard (e.g. $I_{h'k'l',S}$) mixed with our original mixture in known proportion \rightarrow the **ISM method is only applicable to powders!**

Let us consider a mixture M of n phases $\alpha, \beta, \gamma \dots$

Mixture M
(e.g. 3 phases)

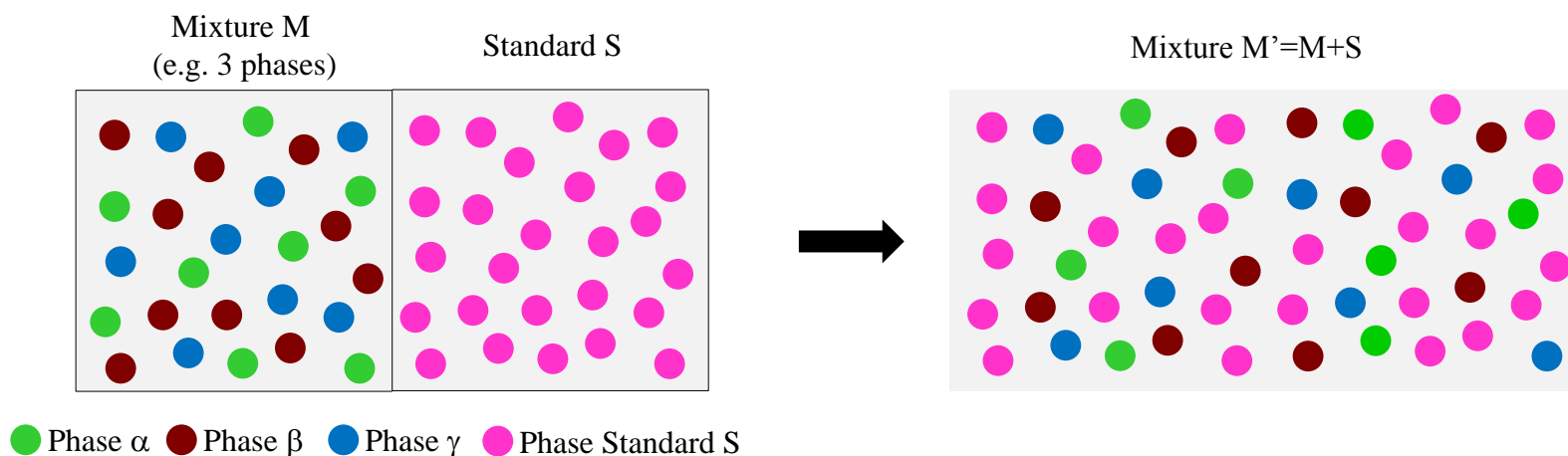


● Phase α ● Phase β ● Phase γ

We need to quantify the **amount of phase α** in a series of mixtures of type M in which the relative proportion of the other phases $\beta, \gamma \dots$ might change from mixture to mixture.

INTERNAL STANDARD METHOD: how does it work?

- i. We mix a known amount of original mixture **M** with a known amount of a known standard **S** and form a new mixture **M' = M + S** (e.g. 50% M + 50% S)



- ii. Let c_α and c'_α the volume fractions of phase α in M and M' (both unknown!) and c_S the volume fraction of the standard S (that we know!)

INTERNAL STANDARD METHOD: how does it work?

- i. We mix a known amount of original mixture M with a known amount of a known standard S and form a new mixture M'=M+S (e.g. 50% M + 50% S)
- ii. Let c_α and c'_α the volume fractions of phase α in M and M' (both unknown!) and c_S the volume fraction of the standard S (that we know!)
- iii. From a powder diffraction pattern recorded on the new mixture M', we obtain:

$$I_{(hkl)\alpha} = I_\alpha = \frac{K_{1,\alpha} \cdot c_{\alpha'}}{\mu_m} \quad \text{and similarly:} \quad I_{(h'k'l')S} = I_S = \frac{K_{1,S} \cdot c_S}{\mu_m}$$

Note that μ_m **cancels out!**

$$\longrightarrow \frac{I_\alpha}{I_S} = \frac{K_{1,\alpha} \cdot c_{\alpha'}}{K_{1,S} \cdot c_S}$$

It physically means that the variation of absorption due to the variation of the relative amounts of the other phases present in the original mixture (β, γ, \dots) does not affect the I_α/I_S ratio since such variations equivalently affects I_α and I_S !

INTERNAL STANDARD METHOD: how does it work?

- i. We mix a known amount of original mixture M with a known amount of a known standard S and form a new mixture M'=M+S (e.g. 50% M + 50% S)
- ii. Let c_α and c'_α the volume fractions of phase α in M and M' (both unknown!) and c_S the volume fraction of the standard S (that we know!)
- iii. From a powder diffraction pattern recorded on the new mixture M', we obtain:

$$\frac{I_\alpha}{I_S} = \frac{K_{1,\alpha} \cdot c'_\alpha}{K_{1,S} \cdot c_S} \quad \longrightarrow \quad \text{with} \quad \frac{c'_\alpha}{c_S} = \frac{w'_\alpha \cdot \rho_S}{\rho_\alpha \cdot w_S} \quad \longrightarrow \quad \frac{I_\alpha}{I_S} = \frac{K_{1,\alpha}}{K_{1,S}} \cdot \frac{w'_\alpha \cdot \rho_S}{\rho_\alpha \cdot w_S}$$

If w_S is kept constant in all mixtures of type M', then

$$\frac{K_{1,\alpha}}{K_{1,S}} \cdot \frac{\rho_S}{\rho_\alpha \cdot w_S} = K_3$$

$$\longrightarrow \frac{I_\alpha}{I_S} = K_3 \cdot w'_\alpha \quad \longrightarrow \quad \frac{I_\alpha}{I_S} = K_4 \cdot w_\alpha \quad \text{being:} \quad w'_\alpha = w_\alpha (1 - w_S)$$

INTERNAL STANDARD METHOD: how does it work?

- i. We mix a known amount of original mixture M with a known amount of a known standard S and form a new mixture $M'=M+S$ (e.g. 50% M + 50% S)
- ii. Let c_α and c'_α the volume fractions of phase α in M and M' (both unknown!) and c_S the volume fraction of the standard S (that we know!)
- iii. From a powder diffraction pattern recorded on the new mixture M', we obtain:

$$\frac{I_\alpha}{I_S} = K_4 \cdot w_\alpha$$

↑
Slope of the
straight line

- The intensity ratio $\frac{I_\alpha}{I_S}$ is therefore a **linear function** of the weight fraction w_α of phase α .
- A calibration curve can be prepared from XRPD measurements on a set of *ad-hoc* synthetic samples containing **known** concentrations of phase α and a **constant** concentration w_S of a suitable standard
- The concentration of α in an unknown mixture is obtained by measuring the ratio $\frac{I_\alpha}{I_S}$ in a mixture of type M' (so-called *spiked* sample) containing the unknown original mixture and the standard in the same proportion as used to build up the calibration curve.

Generalization of the Internal Standard Method → → The **Reference Intensity Ratio (RIR)**

$$\frac{I_{\alpha}}{I_S} = K_4 \cdot w_{\alpha}$$

The calibration constant K_4 depends on: $\alpha, S, (hkl)_{\alpha}, (h'k'l')_S$ and w_S

(remember: $I_{\alpha} = I_{(hkl)_{\alpha}}$ and $I_S = I_{(h'k'l')_S}$)

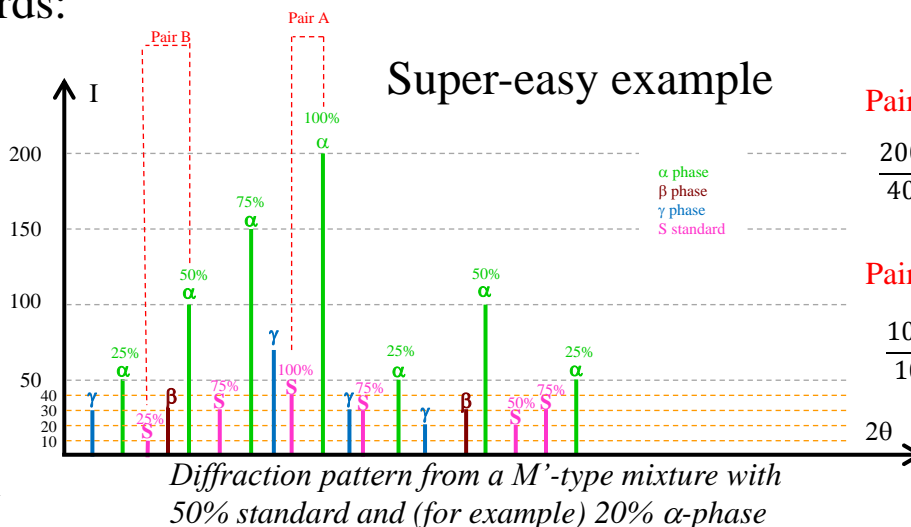
For a more reliable quantification, the use of several analyte-line/internal standard-line pair is preferable → each pair requires a calibration constant!

The use of **relative intensities I^{rel}** and so-called **Relative Intensity Ratio (RIR)** allows us to use multiple analyte/standard reflection pairs without the need of multiple reference standards:

Hubbard & Snyder, (1988)
Powder Diffraction 3, 74-78.

$$\frac{I_{i\alpha}}{I_{jS}} \cdot \frac{I_{jS}^{rel}}{I_{i\alpha}^{rel}} \cdot \frac{w_S}{w_{\alpha}} = K = RIR_{\alpha,S}$$

i denotes one hkl reflection of α phase
j denotes one h'k'l' reflection of the internal standard



Pair A:

$$\frac{200}{40} \cdot \frac{100}{100} \cdot \frac{0.5}{0.2} = 12.5 = K = RIR_{\alpha,S}$$

Pair B:

$$\frac{100}{10} \cdot \frac{25}{50} \cdot \frac{0.5}{0.2} = 12.5 = K = RIR_{\alpha,S}$$

The Reference Intensity Ratio (RIR)

$$K = RIR_{\alpha,S} = \frac{I_{i\alpha}}{I_{jS}} \cdot \frac{I_{jS}^{rel}}{I_{i\alpha}^{rel}} \cdot \frac{w_S}{w_\alpha}$$

Universal calibration constant relating the scattering power of phase α to that of the standard S

→ So-called **Reference Intensity Ratio** or **RIR**

Standard S → **NIST Corundum** → **RIR=I/I_C**

The Powder Diffraction File (PDF) contains I/I_C ratios for more than 2500 phases!

Quantitative Phase Analysis with I/I_C (RIR or Chung Method)

$$w_\alpha = \frac{I_{i\alpha}}{I_{jC}} \cdot \frac{I_{jC}^{rel}}{I_{i\alpha}^{rel}} \cdot \frac{w_S}{RIR_{\alpha,C}} = \frac{I_{i\alpha}}{I_{jC}} \cdot \frac{I_{jC}^{rel}}{I_{i\alpha}^{rel}} \cdot \frac{w_S}{(I/I_C)_\alpha}$$

With the addition of a known %wt of corundum, no calibration curves needed!

Or **standardless QPA**, if all phases in the mixture are crystalline, identified, and the RIR values known for each of them!

DIFFRACTION-BASED METHODS

Single-peak methods:

(adapted from Cullity, *Elements of X-Ray Diffraction*)

- Intensity ratio $I_{\text{unknown}}/I_{\text{standard}}$ of one or more reflections
- No need of structural information but prone to systematic errors caused by PO and peak overlapping
- Need ad-hoc mixtures for calibration curves

Rietveld-based methods:

(adapted from Madsen & Scarlett in *Powder Diffraction-Theory and Practice*)

- Use of full diffraction patterns
- Minimization of systematic errors (e.g. due to peak overlap)
- Preferential Orientation (PO) can be modeled
- Accuracy close to X-Ray fluorescence elemental analysis, with the advantage of being sensitive to structural differences → direct QPA of polymorphs

Rietveld QPA METHOD

*“The Rietveld method uses a model to calculate a diffraction pattern which is then compared with observed data. The difference between the two patterns is then reduced through least square minimization. **The refinable parameters used in the models provide the analyst with information regarding the crystal structure of the component phases, the crystalline size and strain and, importantly, their relative proportions.** The **Rietveld scale factor S**, which is a multiplier for each components’ contribution to the pattern, is related to the relative abundance of that phase and can be used in the quantification of phases.”*

(textual citation from *Quantitative Phase Analysis using the Rietveld Method*, Madsen, Scarlett, Riley & Raven, Ch.10 in *Modern Diffraction Methods*, Mittemeijer & Welzel Edts, 2013)

Rietveld QPA METHOD

$$I_{(hkl)\alpha} = \left[\frac{I_0 \lambda^3}{32 \pi r} \cdot \frac{\sigma e^4}{m_e^2 c^4} \right] \cdot \left[\frac{M_{hkl}}{V_\alpha^2} \cdot |F_{hkl}|_\alpha^2 \left(\frac{1}{\sin^2 \theta \cdot \cos \theta} \right) \right] \cdot \exp \left(-2B_\alpha \left(\frac{\sin \theta}{\lambda} \right) \right) \cdot \left[\frac{w_\alpha}{\rho_\alpha \mu_m^*} \right]$$

D-S geometry with capillary, assuming absorption, PO and extinction negligible

$$I_{(hkl)\alpha} = \frac{K_{1,\alpha} \cdot w_\alpha}{\rho_\alpha \cdot \mu_m^*} \longrightarrow w_\alpha = \frac{I_\alpha \cdot \rho_\alpha \cdot \mu_m^*}{K_{1,\alpha}} = \frac{S_\alpha \cdot (ZMV)_\alpha \cdot \mu_m^*}{K}$$

☺ $K_{1,\alpha} \propto \frac{1}{V_\alpha^2}$

☺ $I_\alpha \propto S_\alpha$

R. J. Hill, Powder Diffr. 1991, **6**, 74-77

☺ $\rho_\alpha = 1.6604 \cdot \frac{ZM_\alpha}{V_\alpha}$

Z=number of formula units

M=molecular mass of the formula unit

Rietveld QPA METHOD

$$I_{(hkl)\alpha} = \left[\frac{I_0 \lambda^3}{32 \pi r} \cdot \frac{\sigma e^4}{m_e^2 c^4} \right] \cdot \left[\frac{M_{hkl}}{V_\alpha^2} \cdot |F_{hkl}|_\alpha^2 \left(\frac{1}{\sin^2 \theta \cdot \cos \theta} \right) \right] \cdot \exp \left(-2B_\alpha \left(\frac{\sin \theta}{\lambda} \right) \right) \cdot \left[\frac{w_\alpha}{\rho_\alpha \mu_m^*} \right]$$

D-S geometry with capillary, assuming absorption, PO and extinction negligible

$$w_\alpha = \frac{S_\alpha \cdot (ZMV)_\alpha \cdot \mu_m^*}{K}$$



IMPORTANT RELATION

In 1988, O'Connor & Raven demonstrated that the scaling factor K is independent of individual phases and overall sample-related parameters (Powder Diffr. **3**, 2-6) .



This implies that K only needs to be estimated once for a given instrumental configuration and using an appropriate standard

→ **EXTERNAL STANDARD METHOD** for absolute QPA analysis

Warning: the term “*instrumental configuration*” really refers to all experimental details →
→ in DS geometry with powders in capillaries it would imply also the same powder packing!

Rietveld QPA METHOD

$$I_{(hkl)\alpha} = \left[\frac{I_0 \lambda^3}{32 \pi r} \cdot \frac{\sigma e^4}{m_e^2 c^4} \right] \cdot \left[\frac{M_{hkl}}{V_\alpha^2} \cdot |F_{hkl}|_\alpha^2 \left(\frac{1}{\sin^2 \theta \cdot \cos \theta} \right) \right] \cdot \exp \left(-2B_\alpha \left(\frac{\sin \theta}{\lambda} \right) \right) \cdot \left[\frac{w_\alpha}{\rho_\alpha \mu_m^*} \right]$$

D-S geometry with capillary, assuming absorption, PO and extinction negligible

$$w_\alpha = \frac{S_\alpha \cdot (ZMV)_\alpha \cdot \mu_m^*}{K}$$



IMPORTANT RELATION

$(ZMV)_\alpha$ is the so-called *calibration constant* for phase α that can be calculated from the structural model (either from crystallographic database or the refinement of the pure α phase)

Absolute QPA analysis can, then, be obtained provided we correctly estimate K and μ_m^* for all our mixtures and calibration standards (the latter for K determination)

Rietveld QPA METHOD: Questions & Answers

$$w_{\alpha} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha} \cdot \mu_m^*}{K}$$



Q₁: Can we still perform **absolute QPA** analyses without estimating **K** and μ_m^* ?

A₁: Yes, we can, if we “spike” our unknown mixture with a known amount w_S of an appropriate reference standard S of well known crystallographic structure (**INTERNAL STANDARD METHOD**)

$$w'_{\alpha} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha} \cdot \mu_m^*}{K}$$



$$w'_{\alpha} = w_S \cdot \frac{S_{\alpha} \cdot (ZMV)_{\alpha}}{S_S \cdot (ZMV)_S}$$



$$w_S = \frac{S_S \cdot (ZMV)_S \cdot \mu_m^*}{K}$$

$$w'_{\alpha} = w_{\alpha}(1 - w_S)$$

Remark: note that there is no need of a calibration curve as for the Internal Standard Method as implemented in the single-line diffraction method in virtue of performing here a Rietveld refinement!

Rietveld QPA METHOD: Q&A

$$w_{\alpha} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha} \cdot \mu_m^*}{K} \quad \star$$

$$w'_{\alpha} = w_S \cdot \frac{S_{\alpha} \cdot (ZMV)_{\alpha}}{S_S \cdot (ZMV)_S} \quad \star \star$$

$$w'_{\alpha} = w_{\alpha}(1 - w_S)$$

Q₂: What if we **do NOT** dispose of the whole composition of the unknown mixture?

A₂: We have 2 possibilities:

> if we can reasonably estimate K and μ_m^* , we apply \star , that is the External Standard Method.

> we spike the sample with a known amount of a known standard and apply the $\star \star$ (Internal Standard Method)

Rietveld QPA METHOD: Q&A

$$w_{\alpha} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha} \cdot \mu_m^*}{K} \quad \star$$

$$w'_{\alpha} = w_S \cdot \frac{S_{\alpha} \cdot (ZMV)_{\alpha}}{S_S \cdot (ZMV)_S} \quad \star \star$$

$$w'_{\alpha} = w_{\alpha}(1 - w_S)$$

Q₂: What if we dispose of the whole composition of the unknown mixture?

A₂: For a mixture of n crystalline phases $\alpha, \beta, \gamma, \delta \dots$ we can write:

$$w_{\alpha} + w_{\beta} + w_{\gamma} + \dots = \sum_{i=1}^n w_i = 1$$



$$w_{\alpha} = \frac{w_{\alpha}}{\sum_{i=1}^n w_i} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha}}{\sum_{i=1}^n S_i \cdot (ZMV)_i}$$

and equivalent expressions for $w_{\beta}, w_{\gamma} \dots$

Hill & Howard (JAC (1987). 20, 467-474) modification of the Rietveld QPA method, known as the *ZMV approach* inspired by the *Matrix Flushing Method* of **Chung** (JAC, 1974, 7, 519-525 and 526-531)

Rietveld QPA METHOD: Q&A

$$w_{\alpha} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha} \cdot \mu_m^*}{K} \quad \star$$

External Standard Method

$$w'_{\alpha} = w_S \cdot \frac{S_{\alpha} \cdot (ZMV)_{\alpha}}{S_S \cdot (ZMV)_S} \quad \star \star$$

$$w'_{\alpha} = w_{\alpha}(1 - w_S)$$

Internal Standard Method

$$w_{\alpha} + w_{\beta} + w_{\gamma} + \dots = \sum_{i=1}^n w_i = 1$$

$$w_{\alpha} = \frac{w_{\alpha}}{\sum_{i=1}^n w_i} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha}}{\sum_{i=1}^n S_i \cdot (ZMV)_i}$$

Hill & Howard (ZMV) Approach

- Warning:** the (ZMV) approach assumes that:
- All phases in the mixtures are crystalline!
 - We have identified them all!



If our mixture has **unknown crystalline phases** or **amorphous components**, a QPA analysis via (ZMV) approach inevitably **overestimates** the w_i weight fractions

Rietveld QPA METHOD: Q&A

$$w_{\alpha} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha} \cdot \mu_m^*}{K} \quad \star$$

External Standard Method

$$w'_{\alpha} = w_S \cdot \frac{S_{\alpha} \cdot (ZMV)_{\alpha}}{S_S \cdot (ZMV)_S} \quad \star \star$$

$$w'_{\alpha} = w_{\alpha} (1 - w_S)$$

Internal Standard Method

$$w_{\alpha} + w_{\beta} + w_{\gamma} + \dots = \sum_{i=1}^n w_i = 1$$

$$w_{\alpha} = \frac{w_{\alpha}}{\sum_{i=1}^n w_i} = \frac{S_{\alpha} \cdot (ZMV)_{\alpha}}{\sum_{i=1}^n S_i \cdot (ZMV)_i}$$



What do we do in such cases?

Hill & Howard (ZMV) Approach

We apply the Internal Standard Method:

➤ We add a known amount of an appropriate standard S

➤ We write $\star \star$ for all identified crystalline phases i and for S

➤ For each phase i we can write: $Corrected(W_i) = W_{i,Rietveld} \frac{W_{S,true}}{W_{S,Rietveld}}$

➤ $W_{unknown} = 1.0 - \sum_{k=1}^n Corrected(W_k)$

What if a **structural model is NOT available** or if it does
not work well with our experimental data?

➤ Partial Or No Known Crystal Structure (**PONKCS**) Method

Madsen & Scarlett, Powder Diffr. 21(4), 2006, 278-284;

Madsen, Scarlett & Kern, Z. Kristallogr.226 (2011) 944-955 → implemented in Topas

- *If **partial structure available** (i.e. unit cell and SG) → real structure factors substituted with empirical values derived from a Pawley or LeBail refinement performed on pure phases → an **hkl_Is phase** in Topas*
- *If **partial structure NOT available** → real structure factor substituted by fictitious phases consisting each of a series of related peaks with **FIXED** relative intensities and **GROUP-SCALED** as a single entity during the QPA analysis → a **x0_Is phase** or **peak-phase** in Topas*
- *Compute empirical **ZM** or **ZMV calibration constants** from the refinement of ad-hoc mixtures of pure phases with a known amount of a known standard (e.g. via spiking) → so-called **PONKCS phases**.*

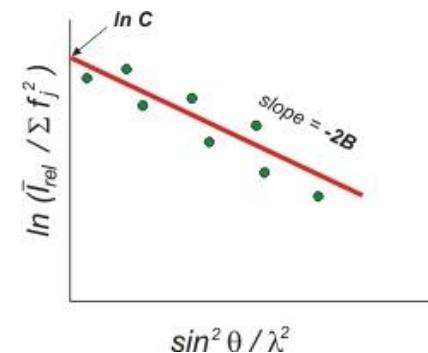
Requirements:

- Pure phases (or pure phases with known impurities) must be available
- *Ad-hoc* mixtures of pure phases with an appropriate standard in known % wt must be available to build the so-called PONKCS phase

Benefits:

- No need of a valid structural model
- PONKCS phases can be “re-used” provided the QPA analyses are conducted at the same photon energy!
→ Careful with SR-XRPD data!
- PONKCS works for **crystalline** as well as **amorphous phases**

Whole-patterns QPA Methods



➤ QUANTO+

Giannini, Guagliardi & Mililli, JAC (2002). 35, 481-490

- If *partial structure available* (i.e. unit cell and SG) → for each phase in the mixture a reflection intensity file is built via whole pattern decomposition (e.g. Le Bail refinement) performed on pure phases
- This «external» file is used instead of the calculated structural factors from the model ($|F_c|^2$) for the absolute scaling of the diffracted intensity via a Wilson plot

Requirements:

- Pure phases must be available
- Partial structure (unit cell and SG) must be known
- Crystalline phases

Benefits:

- No need to have a valid structural model
- No need to prepare *ad-hoc* mixtures with an appropriate standard!

→ implemented in Quanto (CNR-IC)

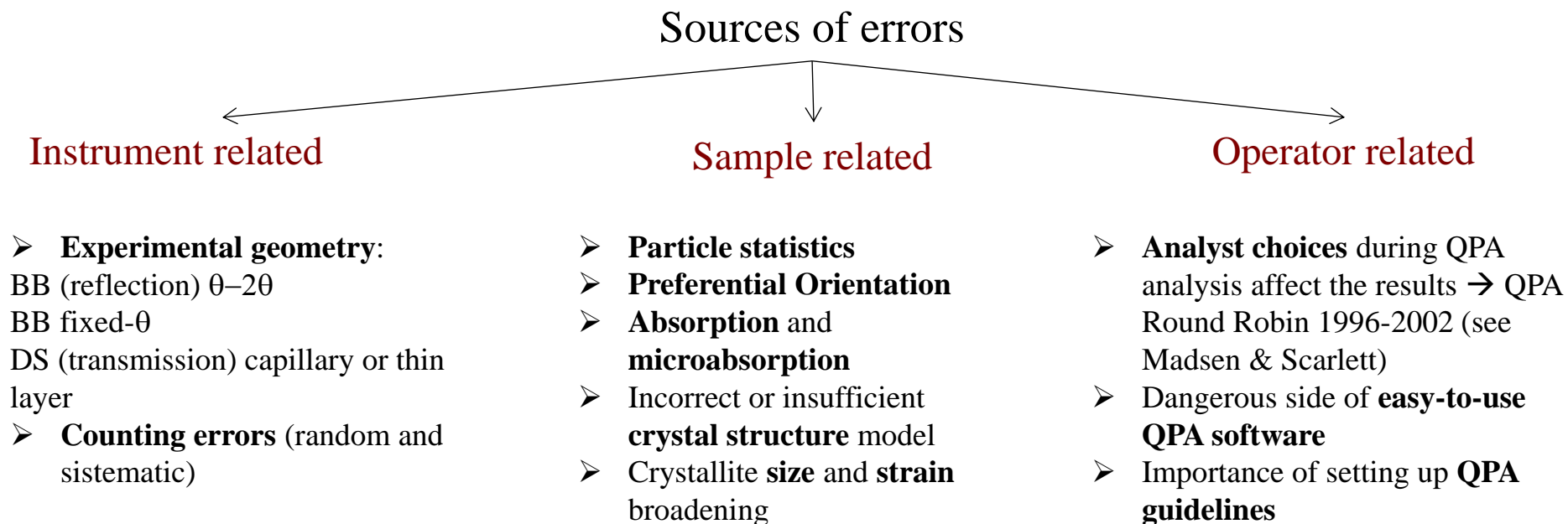
Additional topics not covered during the
lecture plus a few examples of application

Factors influencing QPA

For a detailed and exhaustive discussion and references citation see:

- ❖ Madsen & Scarlett (*Ch. 11, Powder Diffraction – Theory and Practice*), **2009**
- ❖ Madsen, Scarlett, Riley & Raven (*Ch.10, Modern Diffraction Methods*), **2013**

Accurate QPA  Accurate integrated intensity
(ca. 1-2% relative)



Accuracy & Precision associated to QPA

What sets QPA accuracy and precision?

What does a Rietveld (or Rietveld-like) refinement return?

Sources of errors

Instrument related

- **Experimental geometry:**
BB (reflection) $\theta-2\theta$
BB fixed- θ
DS (transmission) capillary or thin layer
- **Counting errors** (random and systematic)

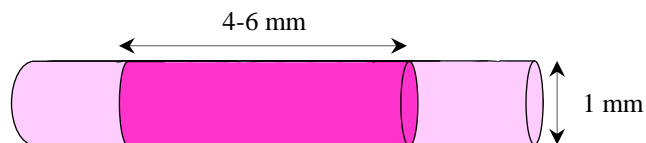
Sample related

- **Particle statistics**
- **Preferential Orientation**
- **Absorption and microabsorption**
- Incorrect or insufficient **crystal structure** model
- Crystallite **size** and **strain** broadening

Operator related

- **Analyst choices** during QPA analysis affect the results \rightarrow QPA Round Robin 1996-2002 (see Madsen & Scarlett)
- Dangerous side of **easy-to-use QPA software**
- Important of setting up **QPA guidelines**

The accuracy in the I_{hkl} estimate is strongly influenced by N_{diff} in a powder sample



Glass capillary

$$V \sim 3 - 5 \text{ mm}^3 \rightarrow N_{diff} \sim 3 \rightarrow \sigma_{PS} \sim 0.6$$

Table 11.1 Relationship between crystallite diameter and the number diffracting (after Smith²⁴).

Crystallite diameter (μm)	40	10	1
Crystallites (20 mm^3)	5.97×10^5	3.82×10^7	3.82×10^{10}
Number diffracting N_{diff}	12	760	38 000
σ_{PS}	0.289	0.036	0.005

From Madsen & Scarlett, *Powder Diffraction-Theory and Practice*, p. 309 reported from original work by D.K. Smith, *Adv. X-Ray Anal.* 1992, 35, 1-15; Elton & Salt, *Powder Diffr.*, 1996, 11, 218-229.

Sources of errors

Instrument related

- **Experimental geometry:**
BB (reflection) $\theta-2\theta$
BB fixed- θ
DS (transmission) capillary or thin layer
- **Counting errors** (random and systematic)

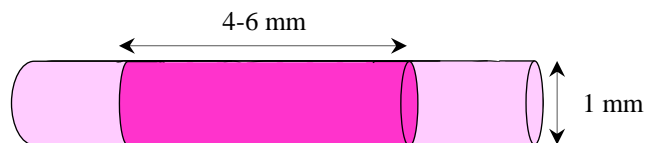
Sample related

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The accuracy in the I_{hkl} estimate is strongly influenced by N_{diff} in a powder sample



Glass capillary

$$V \sim 3 - 5 \text{ mm}^3 \rightarrow N_{diff} \sim 3 \rightarrow \sigma_{PS} \sim 0.6$$

Remedies:

- Increase instrument beam divergence
- Sample spinning or oscillation
- Increase the powder volume analyzed or powder repacking
- Mechanical comminution (grinding, milling)

Sources of errors

Instrument related

- **Experimental geometry:**
BB (reflection) $\theta-2\theta$
BB fixed- θ
DS (transmission) capillary or thin layer
- **Counting errors** (random and systematic)

Sample related

- **Particle statistics**
- **Preferential Orientation**
- **Absorption and microabsorption**
- Incorrect or insufficient **crystal structure** model
- Crystallite **size** and **strain** broadening

Operator related

- **Analyst choices** during QPA analysis affect the results → QPA Round Robin 1996-2002 (see Madsen & Scarlett)
- Dangerous side of **easy-to-use QPA software**
- Important of setting up **QPA guidelines**

DS geometry + capillary + 1D position sensitive detectors:

- intensity modulation due to inhomogeneous capillary packing
- inhomogeneous photon beam distribution

Remedies:

- Partial photon beam focusing
- Analysis of a large number of powder volumes

Sources of errors

Instrument related

- **Experimental geometry:**
BB (reflection) $\theta-2\theta$
BB fixed- θ
DS (transmission) capillary or thin layer
- **Counting errors** (random and systematic)

Sample related

- **Particle statistics**
- **Preferential Orientation**
- **Absorption and microabsorption**
- Incorrect or insufficient **crystal structure** model
- Crystallite **size** and **strain** broadening

Operator related

- **Analyst choices** during QPA analysis affect the results → QPA Round Robin 1996-2002 (see Madsen & Scarlett)
- Dangerous side of **easy-to-use QPA software**
- Important of setting up **QPA guidelines**

- The need of absorption corrections should be minimized and corrections appropriate for the given experimental geometry applied
- **Microabsorption** (i.e. absorption contrast) occurs when the phases in a mixture are characterized by different mass absorption μ^* coefficients and/or different crystallite size → the %wt of the phase with higher μ^* and/or larger crystallite size would be underestimated

Remedies:

- Choose an appropriate λ
- Reduce crystallite size

Sources of errors

Instrument related

- **Experimental geometry:**
BB (reflection) $\theta-2\theta$
BB fixed- θ
DS (transmission) capillary or thin layer
- **Counting errors** (random and systematic)

Sample related

- **Particle statistics**
- **Preferential Orientation**
- **Absorption and microabsorption**
- **Incorrect or insufficient crystal structure model**
- Crystallite **size and strain** broadening

Operator related

- **Analyst choices** during QPA analysis affect the results → QPA Round Robin 1996-2002 (see Madsen & Scarlett)
- Dangerous side of **easy-to-use QPA software**
- Important of setting up **QPA guidelines**

- Strong correlation between the **Rietveld scale factor** and the **Atomic Displacement Parameters** (ADP , B_{iso} , U_{iso}).

WARNING: very often .cif files in the crystallographic database DO NOT report such parameters → commercial programs use in such cases default values (i.e. 1 \AA^2)

Be careful: most programs require as input B_{iso} (e.g. Topas, FullProf), often you find in literature and database $U_{iso} \rightarrow B_{iso} = 8\pi^2 U_{iso}$

EXAMPLES

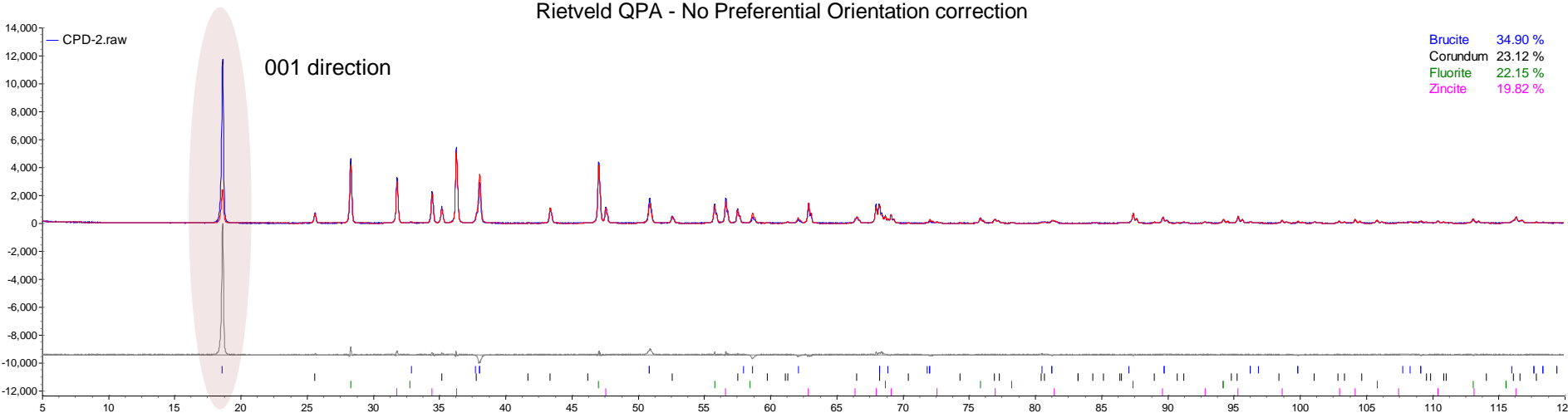
Example 1: Quantification of wt% of Corundum ($\alpha\text{-Al}_2\text{O}_3$), Fluorite (CaF_2) and Zincite (ZnO) using *ad-hoc* synthetic mixtures: **CPD-2.RAW**

Source:

IUCr CPD Round Robin on Quantitative Phase Analysis
Data available from: <http://www.iucr.org/iucr-top/comm/cpd/QARR/index.html>
Madsen, Scarlett, Cranswick and Lwin (2001). J.Appl. Cryst, 34, 409-426.

Expected %wt:
Brucite: 36.26%
Corundum: 21.27%
Fluorite: 22.53%
Zincite: 19.94%

Rietveld QPA - No Preferential Orientation correction



Corresponding to the (001) direction of brucite, the calculated diffracted intensity clearly poorly matches the experimental data

→ PO effects presumably decreases the accuracy of quantification!

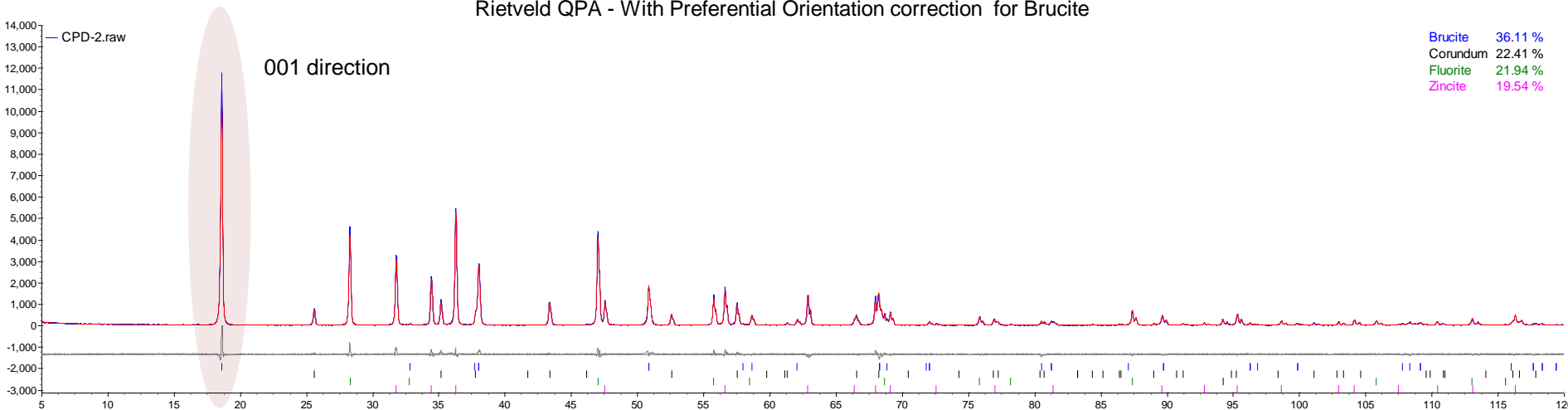
Example 1: Quantification of wt% of Corundum ($\alpha\text{-Al}_2\text{O}_3$), Fluorite (CaF_2) and Zincite (ZnO) using *ad-hoc* synthetic mixtures: **CPD-2.RAW**

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Madsen, Scarlett, Cranswick and Lwin (2001). J. Appl. Cryst, 34, 409-426.

Expected %wt:
Brucite: 36.26%
Corundum: 21.27%
Fluorite: 22.53%
Zincite: 19.94%

Rietveld QPA - With Preferential Orientation correction for Brucite

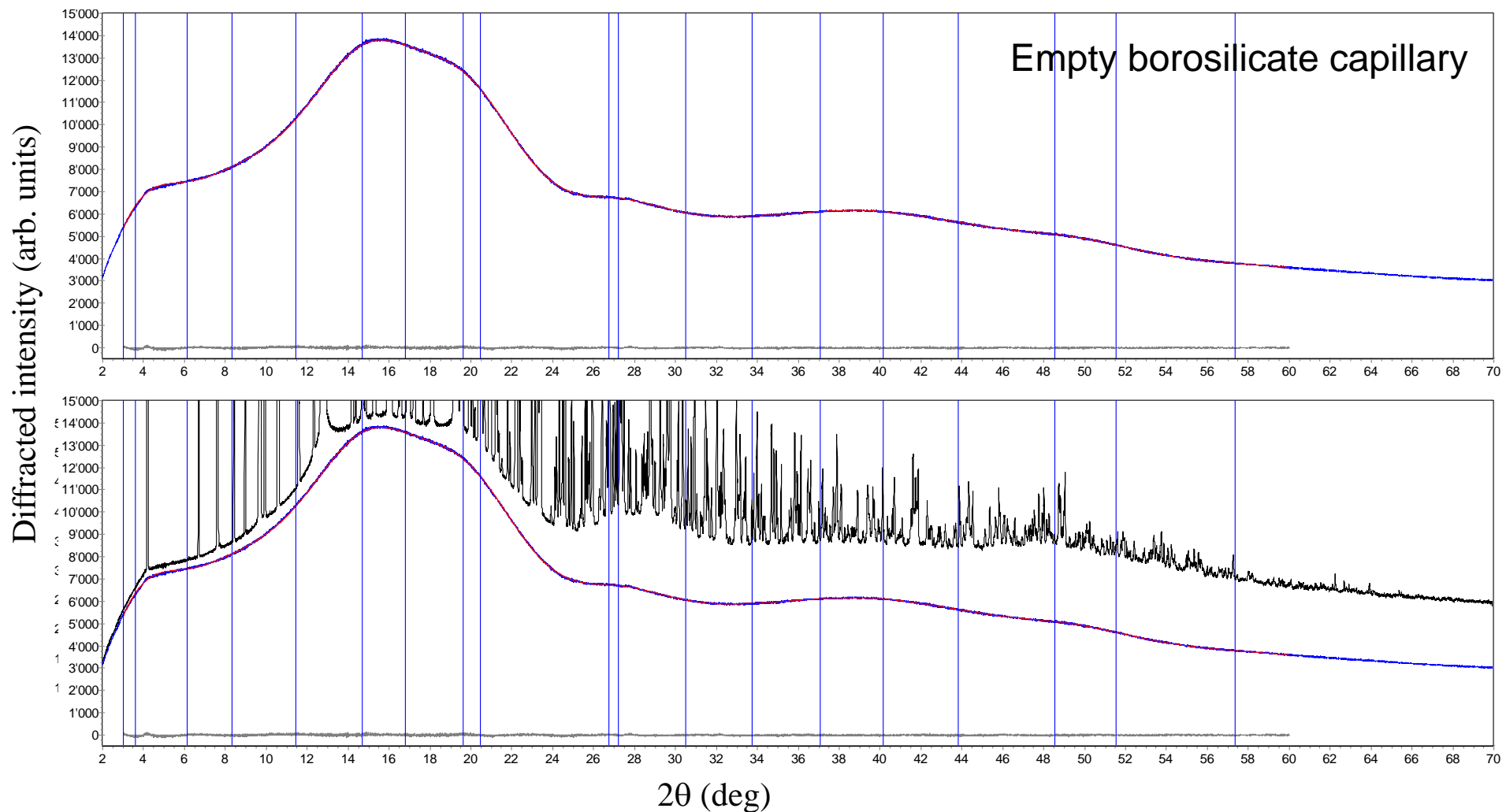


The modeling of the PO for (001) direction of brucite improves the refinement and the QPA accuracy.

The accuracy of the quantitative phase analysis strongly depends on:

- The quality of the refinement models (e.g. background, instrumental contribution, crystal structure, microstructure)
- How close to the correct values of all these refinement parameters we start the quantitative analysis

Example 2: A good description of the extrinsic background with a limited number of parameters



Example 3: Quantification of wt% of an Amorphous phase in a mixture of Fluorite (CaF₂) and Zincite (ZnO) using Corundum (α-Al₂O₃) as an internal standard (spiking method)

Expected values: 28.41% Fluorite+27.45% Zincite + 44.14% Glass

Source:

IUCr CPD Round Robin on Quantitative Phase Analysis

Data available from: <http://www.iucr.org/iucr-top/comm/cpd/QARR/index.html>

Madsen, Scarlett, Cranswick and Lwin (2001). J. Appl. Cryst, 34, 409-426.

Help:

$$w'_\alpha = w_s \cdot \frac{S_\alpha \cdot (ZMV)_\alpha}{S_s \cdot (ZMV)_s}$$



$$w_\alpha + w_\beta + w_\gamma + \dots = \sum_{i=1}^n w_i = 1$$



$$Corr(W_i) = W_i \frac{w_{s,true}}{w_{s,Rietveld}}$$

$$w'_\alpha = w_\alpha(1 - w_s)$$

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Step I: We spike the original mixture M of Fluorite and Zincite with a known amount of a known internal standard (30.79% Corundum) → Mixture of type M'=30.79% IS+69.21% M

Example 3: Quantification of wt% of an Amorphous phase in a mixture of Fluorite (CaF₂) and Zincite (ZnO) using Corundum (α-Al₂O₃) as an internal standard (spiking method)

Expected values: 28.41% Fluorite+27.45% Zincite + 44.14% Glass

Source:

IUCr CPD Round Robin on Quantitative Phase Analysis

Data available from: <http://www.iucr.org/iucr-top/comm/cpd/QARR/index.html>

Madsen, Scarlett, Cranswick and Lwin (2001). J. Appl. Cryst, 34, 409-426.

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$$w'_\alpha = w_S \cdot \frac{S_\alpha \cdot (ZMV)_\alpha}{S_S \cdot (ZMV)_S}$$



$$w_\alpha + w_\beta + w_\gamma + \dots = \sum_{i=1}^n w_i = 1$$

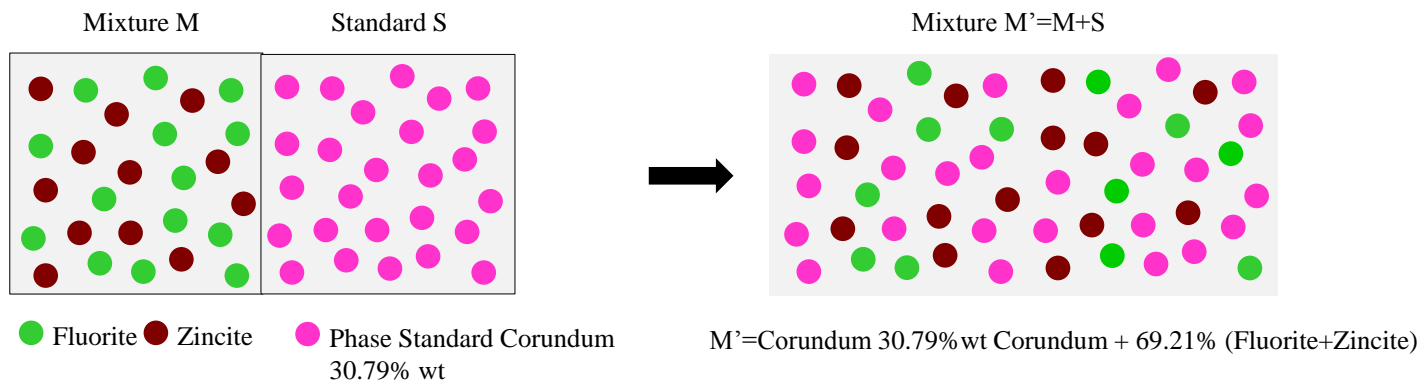


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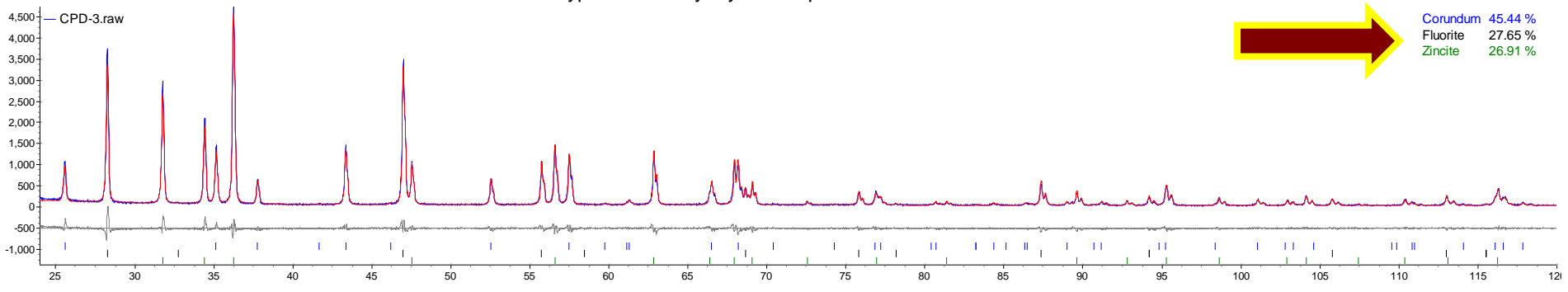
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Hypothesis: only crystalline phases in the mixture



Example 3: Quantification of wt% of an Amorphous phase in a mixture of Fluorite (CaF₂) and Zincite (ZnO) using Corundum (α-Al₂O₃) as an internal standard (spiking method)

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Step II: We perform a Rietveld QPA using the structural information of Fluorite, Zincite and Corundum → 45.44% Corundum+27.65% Fluorite+26.91% Zincite

The overestimation of Corundum of our Rietveld QPA clearly tells us that we DO have an unknown in our original mixture M!

Example 3: Quantification of wt% of an Amorphous phase in a mixture of Fluorite (CaF₂) and Zincite (ZnO) using Corundum (α-Al₂O₃) as an internal standard (spiking method)

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Step III: We calculate the correction factor $\frac{w_{S,true}}{w_{S,Rietveld}} = \frac{w_{Corundum,true}}{w_{Corundum,Rietveld}} \approx \frac{30.79\%}{45.44\%} \approx 0.68$

Step IV: We calculate the corrected % wt:

$$Corr(W'_{Fluorite}) = W'_{Fluorite} \frac{w_{S,true}}{w_{S,Rietveld}} \approx 27.65 \cdot 0.68 \approx 18.80\%$$

$$Corr(W'_{Zincite}) = W'_{Zincite} \frac{w_{S,true}}{w_{S,Rietveld}} \approx 26.91 \cdot 0.68 \approx 18.30\%$$

$$\rightarrow W'_{unknown} = 1.0 - \sum_{k=1}^n Corr(W_k) \approx 32.11\%$$

$$\rightarrow \text{In original mixture: } w_{unknown} = W'_{unknown} / (1 - w_S) \approx 46.40\%$$

Example 3: Quantification of wt% of an Amorphous phase in a mixture of Fluorite (CaF₂) and Zincite (ZnO) using Corundum (α-Al₂O₃) as an internal standard (spiking method)

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$$Corr(W'_{FL.}) = W'_{FL.} \frac{w_{S,true}}{w_{S,Rietveld}} \approx 27.65 \cdot 0.68 \approx 18.80\% \rightarrow w_{Fluorite} \approx 27.16\% \text{ in original mixture } M$$

$$Corr(W'_{Zin.}) = W'_{Zin.} \frac{w_{S,true}}{w_{S,Rietveld}} \approx 26.91 \cdot 0.68 \approx 18.30\% \rightarrow w_{Zincite} \approx 26.44\% \text{ in original mixture } M$$

$$\rightarrow W'_{unknown} = 1.0 - \sum_{k=1}^n Corr(W'_k) \approx 32.11\%$$

$$\rightarrow \text{In original mixture: } w_{unknown} = w'_{unknown} / (1 - w_S) \approx 46.40\%$$

Similarly:

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Quantification of amorphous phase with an internal standard

