Lesson 8 Publishing XRD Results



Nicola Döbelin('s humble opinions...) RMS Foundation, Bettlach, Switzerland

...totally irrelevant to the rest of the world



March 01 – 02, 2016, Freiberg, Germany

••••• Testing • Research • Consulting

Relevant Topics

- Quality of Rietveld refinement results
- Graphs
- «Experimental» section





Quality: Accurate + Precise



High Precision, High Accuracy



Low Precision, High Accuracy



High Precision, Low Accuracy



Low Precision, Low Accuracy https://phidgets.wordpress.com



••••• Testing • Research • Consulting

Quality of Rietveld Refined Data

XRD / Rietveld refinements are very difficult to validate:

- Often no complementary analytical technique available
- Hardly any reference materials available
- Refinement: Highly operator dependent

One option: Participate in round robins

 \rightarrow Compare your results with other labs

Round robin on CaP phase quantification organized by RMS Foundation / Nicola Döbelin in 2012/2013



Accuracy of Rietveld Refined Data

Round Robin Reference Sample:

- Simple 2 phase system: HA + β -TCP
- Very homogeneous distribution
- No texture
- No micro-absorption
- Highly crystalline
- Mean cryst size ~200 nm

→ Nearly «Best Case» Scenario





Round Robin

12 Labs with 26 different instruments / configurations analyzed the same powder

n=5

Results returned from one lab/instrument:

Sample	β-TCP [wt-%]	HA [wt-%]
1	28.01	71.99
2	28.22	71.78
3	28.49	71.51
4	28.29	71.71
5	28.20	71.80
Mean (Std. Dev)	28.24 (0.17)	71.76 (0.17)



Accuracy of Rietveld Refined Data

* p < 0.05 ** p < 0.01



Döbelin, N. "Interlaboratory study on the quantification of calcium phosphate phases by Rietveld refinement." Powder Diffraction, 2015 30(3): 231-241.

••••••• Testing • Research • Consulting

Uncertainty of Measurement

• Random Errors \rightarrow Precision Determine σ from *N* measurements:

$$\sigma = \sqrt{\frac{1}{N} \sum_{i=1}^{N} (x_i - \overline{x})^2}$$

◆ Systematic Errors → Accuracy
Determine ∆ with reference samples

$$\Delta = \left| \bar{x} - x_{reference} \right|$$

$$u(c) = \sqrt{\sigma^2 + \Delta^2}$$

$$U_{k=2} = 2 \cdot u(c)$$

Validation of β-TCP + HAp Quantification at RMS



Validation of β-TCP + HAp Quantification at RMS



Validation of β-TCP + HAp Quantification at RMS





Raynaud, S., Champion, E., Bernache-Assolant, D., and Laval, J.-P. (2001). "Determination of calcium/phosphorus atomic ratio of calcium phosphate apatites using X-ray diffractometry," J. Am. Ceram. Soc. 84, 359–366.

Validated uncertainties of measurement compliant with ISO 17025 accreditation



Quality of Rietveld Refined Data



RMS

Possible Definition of Quantification Limit (LOQ):

Phase Quantity $< 4 \cdot \sigma$

Possible Definition of Detection Limit (LOD):

Phase Quantity $< 2 \cdot \sigma$





LOD / LOQ Warning in Profex





Guidelines for Reporting Phase Quantities

For error bars use 3 · ESD or 2.77 · ESD

 $3 \cdot \text{ESD} = 99.7\%$ repeatability limit, normal distribution 2.77 $\cdot \text{ESD} = 95\%$ repeatability limit, t distribution [1]

Report Values < LOQ as «< 4 - ESD» Report 0.76 ± 0.23 as «< 0.92»</p>

Report Values < LOD as «not detected» Report 0.21 ± 0.11 as «not detected»

[1] ASTM E177-13: «Standard Practice for Use of the Terms Precision and Bias in ASTM Test Methods».







• • • Testing • Research • Consulting

. 50

60

. 40

30

. 30

Angle [°2theta]

Angle [°2theta]

40

50

60

17

RMS

Recommendation:

- Show all your raw data (documents your data quality)
- Show 1 full refinement (documents your refinement quality)

- All the information is in the raw data
- Only additional information in refinement plots (Icalc, Idiff, Bkgr):

«The refinement was done right»





RMS

Combine your raw patterns





Match the level of detail to the relevance of XRD for the study

Used for phase identification

- Basic intrument parameters:
 - Instrument / Manufacturer
 - Scan range (start, end, step size)
- References to PDF / ICDD / COD phase entries







Used for phase quantification

- Detailed intrument parameters:
 - Instrument / manufacturer
 - Scan range (start, end, step size [°2θ])
 - Radiation, filter / monochromator, divergence slit
- References to sources of crystal structures
- Rietveld software (program name, version, reference)





••••••• Testing • Research • Consulting



Used for advanced refinements

(structural parameters, amorphous fractions, crystallite size analysis, texture analysis etc.)

- Detailed intrument parameters:
 - Instrument / manufacturer
 - Scan range (start, end, step size [°2θ])
 - Radiation, filter / monochromator, divergence slit
 - ... (detector, masks, ASS, Soller slits, generator settings...)
- References to sources of crystal structures
- Rietveld software (program name, version, reference)
- Refinement strategy
- Any non-standard calculations, sample preparations, measurement conditions, etc...



ournal

J. Am. Ceram. Soc., 92 [7] 1592–1595 (2009) DOI: 10.1111/j.1551-2916.2009.03093.x © 2009 The American Ceramic Society

Synthesis and Structure Refinement of Zinc-Doped β-Tricalcium Phosphate Powders

Sanjeevi Kannan,[‡] Friedlinde Goetz-Neunhoeffer,[§] Jürgen Neubauer,[§] and Jose M.F. Ferreira^{†,‡}

(2) Structure Refinement

For refinement studies, all the powders were heat treated in a Thermolab furnace (Pt30%Rh/Pt6%Rh-thermocouple) with a heating rate of 5°/min to achieve a predetermined temperature of 1000°C, followed by a dwelling time of 2 h, and then cooled to room temperature at the rate of 5°/min. Although low crystalline β -Ca₃(PO₄)₂ forms at about 800°C, ^{21,22} a higher heat treatment temperature of 1000°C was required to obtain the desired well-crystallized phase for Rietveld refinement. Collection of Xray powder diffraction (XRPD) pattern data was performed with a Siemens D5000 (Karlsruhe, Germany) equipped with a diffracted beam graphite monochromator. The software TO-PAS 3 (Bruker AXS, Karlsruhe, Germany) with the fundamental parameters approach was used for Rietveld refinements. All powders were prepared and measured three times independently. Rietveld refinement was performed using the structural model of ICSD card number of $\# 6191^{23}$ for β -TCP. Refined parameters were scale factor, specimen displacement, background as Chebyshev polynomial of fifth grade, crystallite size, microstrain, lattice parameters, and occupancy factors. As the scattering factor for P^{5+} is not implemented in the TO-PAS 3 Rietveld software, the P sites were refined with a combination of Si⁴⁺ and P for pure β -Ca₃(PO₄)₂ phase. This leads to an improved refinement of Ca^{2+} occupancy and of the zinc substitution at Ca²⁺ sites. Zinc content in the heat-treated powders was analyzed by the ICP technique.



•••••• Testing • Research • Consulting

J. Am. Ceram. Soc., 93 [10] 3455-3463 (2010) DOI: 10.1111/j.1551-2916.2010.03856.x © 2010 The American Ceramic Society

Thermal Treatment of Flame-Synthesized Amorphous Tricalcium Phosphate Nanoparticles

Nicola Döbelin,^{†,‡} Tobias J. Brunner,^{§,††} Wendelin J. Stark,[§] Martin Fisch,[¶] Egle Conforto,[¶] and Marc Bohner[‡]

In situ XRD data sets were collected using a powder diffractometer (X'Pert Pro MPD, Panalytical, Almelo, the Netherlands) in θ/θ configuration using Ni-filtered CuK α radiation and a fixed divergence slit of 1/8° opening. A heating chamber (HTK-1200, Anton Paar, Ostfildern, Germany) was attached to the goniometer. The samples were prepared in a ceramic sample holder spinning at approximately 1 rps. Data sets were collected from 9°-65° 20 with a step size of 0.016° 20 and a counting time of 1 s per step at the following temperatures: 25°, 500°, 525°, 550°, 575°, 600°, 650°, 700°, 750°, 800°, 850°, 900°, 950°, 1000°C, and again at 25°C. The heating rate was set to 1°C/min for all heating sequences, which was slow enough to prevent overshooting within the accuracy of the furnace control unit (±1°C). During data collection, the temperature was held constant, and for the last data acquisition the sample was cooled from 1000° to 25°C at 60°C/min.

XRD patterns were analyzed by Rietveld refinement using the software FullProf.2k.³⁶ Starting models for the crystal struc-tures were taken from Mathew *et al.*³⁷ for α -TCP, Dickens *et al.*³⁸ for β -TCP, and Sudarsanan and Young³⁹ for hydroxyapatite (HA). No additional crystalline phases were observed in the diffraction patterns. The refinements were performed in three steps, starting with vertical sample displacement, scale factors for all phases, and sampled background points with linear interpolation until convergence was reached. In the second step, the cell parameters of all phases were released for refinement, and in the last step isotropic peak broadening of all phases and one common overall isotropic displacement parameter (Boverall) were optimized. The instrument resolution function and peak asymmetry were determined with an "NIST SRM 660a-Line Profile LaB6" standard before the refinements of the sample data sets. Peak shapes were modeled with a pseudo-Voigt function accounting for peak asymmetry due to axial divergence.4 Average crystallite sizes were calculated from isotropic Gaussian peak broadening using the Scherrer equation.

The most common method to determine X-ray amorphous fractions in a specimen is to add a known quantity of a crystalline internal standard. However, at temperatures up to 1000°C, solid-state reactions of the specimen with the standard powder are to be expected. Relative weight fractions W_P of the crystalline phase P, on the other hand, can be calculated from refined parameters and chemical information of all phase as follows⁴¹:

$$W_P = \frac{S_P(ZMV)_P}{\sum_i S_i(ZMV)_i} \tag{1}$$

where S is the Rietveld scale factor, Z the number of formula units per unit cell, M the molecular mass of the formula unit, V the unit cell volume, and i iterates over all crystalline phases. The absolute mass m_P of each crystalline component in the specimen is proportional to

$$m_P \propto S_P(ZMV)_P$$

By making the reasonable assumptions that (i) the intensity of the primary beam and the device configuration were identical for all data sets, (ii) the total amount of powder in the specimen remained constant during the experiment, and (iii) all phases involved, including the amorphous powder, have similar particle sizes and absorption coefficients, the dependency of S_P could be reduced to the phase abundance and the overall atomic displacement parameter. The latter, as well as the unit cell volume, are temperature dependent. The value m_P calculated from Eq. (2) for a constant amount of a phase P will thus show a slight linear decrease with the increasing temperature.

In order to determine the amount of X-ray amorphous material in the specimen, a 100% crystalline reference was needed. As the sum of m_P values of all crystalline phases remained constant between 800° and 1000°C, except for a steady decrease caused by the temperature-dependent parameters, it was assumed that all amorphous material was crystallized below 800°C. A linear regression through the sums of m_P values between 800° and 1000°C thus served as references for 100% crystallinity for the calculation of absolute phase quantities.



Fig. 1. In situ X-ray diffraction patterns and theoretical peak positions for α -tricalcium phosphate (TCP), β -TCP, and hydroxyapatite (HA). Just after crystallization, the main phase was α -TCP, followed by a gradual transformation to β -TCP. The peak shift of the top pattern was caused by the thermal contraction of the unit cell when cooled from 1000°C to room temperature.

Special calculations

Iourna

(2)

Summary: DOs and DON'Ts in Manuscripts

Do:

- Show your raw data (stacked plots)
- Show one full refinement
- Give details according to the relevance of XRD data for the manuscript

Don't:

- Spam the manuscript with refinement plots
- Publish results from non-standard samples
- Publish results from poor refinements!!!



JECT