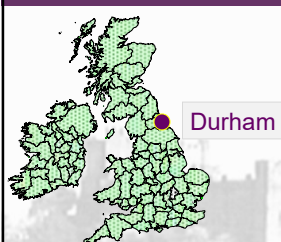
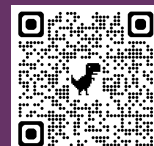


Structural Rietveld Refinements

Prof. John S.O. Evans, Durham Chemistry

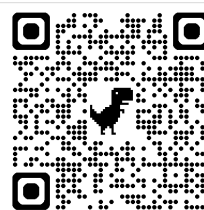
Stuttgart School, 2025



Outline of session

- Fundamentals of structural Rietveld refinement
- Short recap of Robert's content we'll use in common refinements
- Intro to INP files (launch mode)
- Exercises (>30 minutes?)

Stuttgart page



<https://topas.webspace.durham.ac.uk/stuttgart-school-2025/>

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The notes – Whole Powder Pattern Fitting (WPPF)

1. Intro/overview
2. Least squares maths
3. Constraints and restraints
4. Quantities refined
5. Factors affecting intensities
6. Agreement factors
7. Multiple data sets
8. Refinement strategy

[Listen this morning, read at your leisure]

WPPF concise notes

Whole Powder Pattern Fitting

Since $\frac{\partial \chi^2}{\partial p_k} = 0$ at the minimum, it follows that:

$$\sum_{i=1}^N w_i \left(\frac{\partial y_{calc,i}(p_k)}{\partial p_k} - \frac{\partial y_{obs,i}}{\partial p_k} \right) = 0 \quad \text{Eq. 7}$$

Changing the numerators on the left side leads to:

$$\sum_{i=1}^N w_i \left(\frac{\partial y_{calc,i}(p_k)}{\partial p_k} - y_{obs,i} \right) = 0 \quad \text{Eq. 8}$$

This equation is linear in the parameter shifts Δp_k . There will be one equation for each p_k and the set of equations can be concisely expressed in matrix notation as:

$$A \Delta p = Y \quad \text{Eq. 9}$$

with the components of the $P \times P$ matrix A (each k corresponds to a matrix row and each j corresponds to a column) given by:

$$A_{kj} = \sum_{i=1}^N w_i \left(\frac{\partial y_{calc,i}(p_k)}{\partial p_k} - y_{obs,i} \right) \frac{\partial y_{calc,i}(p_j)}{\partial p_j} \quad \text{Eq. 10}$$

and the P components of the vector Y by:

$$Y_j = \sum_{i=1}^N w_i (y_{obs,i} - y_{calc,i}(p_k)) \frac{\partial y_{calc,i}(p_j)}{\partial p_j} \quad \text{Eq. 11}$$

The equations in Eq. 9 are called the normal equations of least squares and can be solved by pre-multiplying each side of the equation by A^{-1} to give the parameter shifts Δp which should be applied to the model.

$$A^{-1} A \Delta p = A^{-1} Y \quad \text{Eq. 12}$$

In practice the approximations made in the Taylor series of Eq. 4 mean that this recipe is not perfect and won't immediately give the best fit model. The process is therefore iterated multiple times, ideally with an improvement between $y_{obs,i}$ and $y_{calc,i}$ on each iteration. The process is stopped when no further improvement is observed in the fit (i.e. there is no significant reduction in χ^2), or the parameter shifts become comparable to their uncertainties. These uncertainties, and the covariance between different parameters, can be extracted from the inverse matrix A^{-1} normalised by the reduced χ^2 (Eq. 26 later). Critical analysis of this variance-covariance matrix to assess model reliability is a crucial part of all WPPF methods.

Much of the work of WPPF software goes into the manipulation of the matrices in Eq. 9 and solution of the normal equations. Modern software packages incorporate a number of mathematical tools to ensure that the fit improves on each cycle and that the refinement doesn't diverge due to mathematical instabilities. Some programs issue warnings about the determinant of matrices being small. These are flagging that the least squares process is unstable. A small determinant of the A matrix means large values in A^{-1} and therefore large uncertainties in the refined parameters.

3. Constraints and restraints

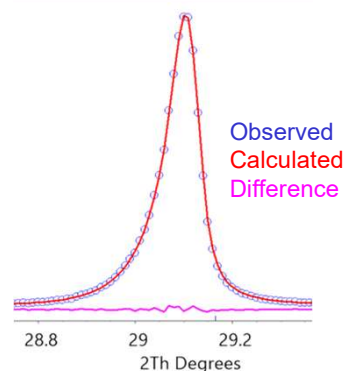
The useful information content of a powder diffraction pattern is often low due to peak overlap or a limited 2θ range. In these cases it may be helpful to include extra information in the refinement in the form of constraints or restraints (sometimes confusingly called "soft constraints").

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What is Rietveld refinement vs WPPF?

- Minimise the difference between an observed and a calculated powder diffraction pattern based on a (structural) model
- Model must contain all the relevant information
- Done by iterative non-linear least squares by calculating parameter shifts required to improve the fit of the starting model
- Maths in the notes
- Key difference to Pawley/Le Bail is that intensities are determined by a crystallographic model – so you can extract structural information



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Minimisation, R-factors, etc

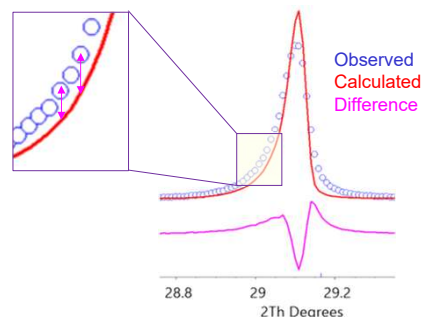
Minimise:
$$\sum_{i=1}^N w_i (y_{obs,i} - y_{calc,i})^2$$

Monitor:
$$R_{wp} = \sqrt{\frac{\sum_{i=1}^N w_i (y_{obs,i} - y_{calc,i})^2}{\sum_{i=1}^N w_i y_{obs,i}^2}}$$

Monitor:
$$GoF = \sqrt{\chi^2} = \frac{R_{wp}}{R_{exp}} = \sqrt{\frac{\sum_{i=1}^N w_i (y_{obs,i} - y_{calc,i})^2}{N - P}}$$

N is the number of data points and P the number of parameters. Least-squares uses a $1/\sigma(y_{obs,i})^2$ weighting, where $\sigma(y_{obs,i})$ is the experimental uncertainty in $y_{obs,i}$

Other R-factors (e.g. background subtracted) in notes



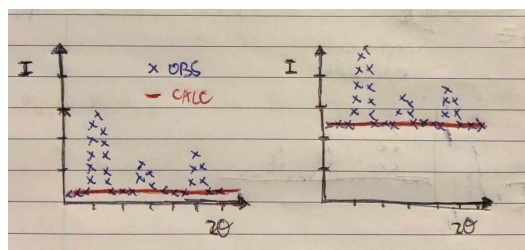
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Target R-factor for a good refinement?

- TOPAS forum question ($\times 100$): how low should my R-factor be for a good refinement?

High R_{wp}
High R_{Bragg}



Low(er) R_{wp}
High R_{Bragg}

- Answer: useful summary in Brian Toby paper cited in the notes

R factors in Rietveld analysis: How good is good enough?

Brian H. Toby
BESSRC/XOR, Advanced Photon Source, Argonne National Laboratory, Argonne, Illinois
(Received 19 December 2005; accepted 27 January 2006)

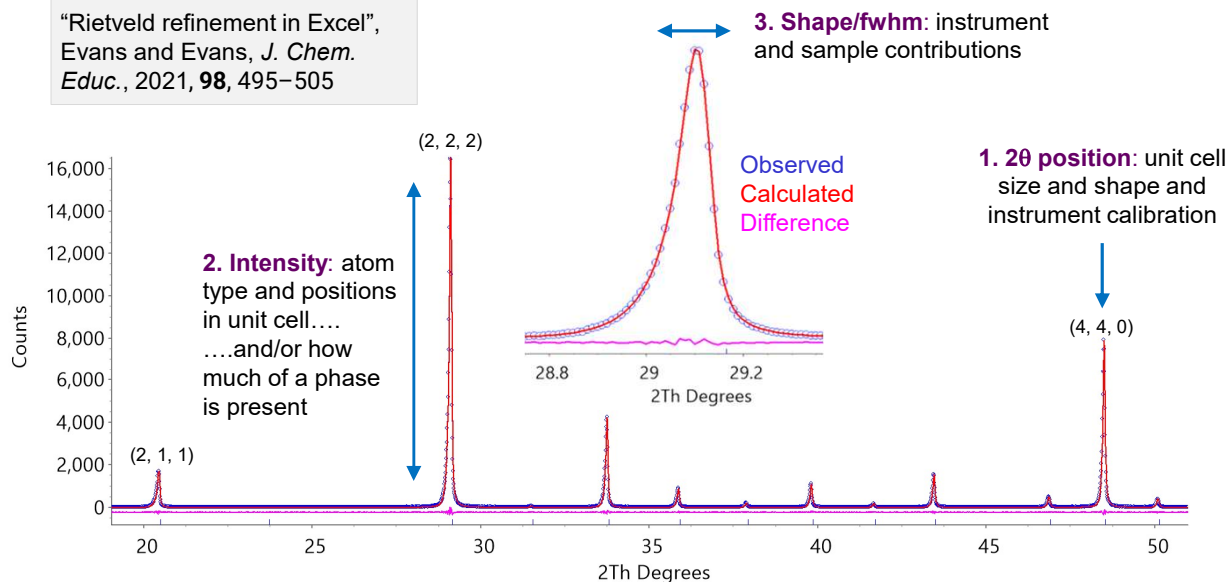
The definitions for important Rietveld error indices are defined and discussed. It is shown that while smaller error index values indicate a better fit of a model to the data, wrong models with poor quality data may exhibit smaller error index values than some superb models with very high quality data. © 2006 International Centre for Diffraction Data. [DOI: 10.1154/1.2179804]

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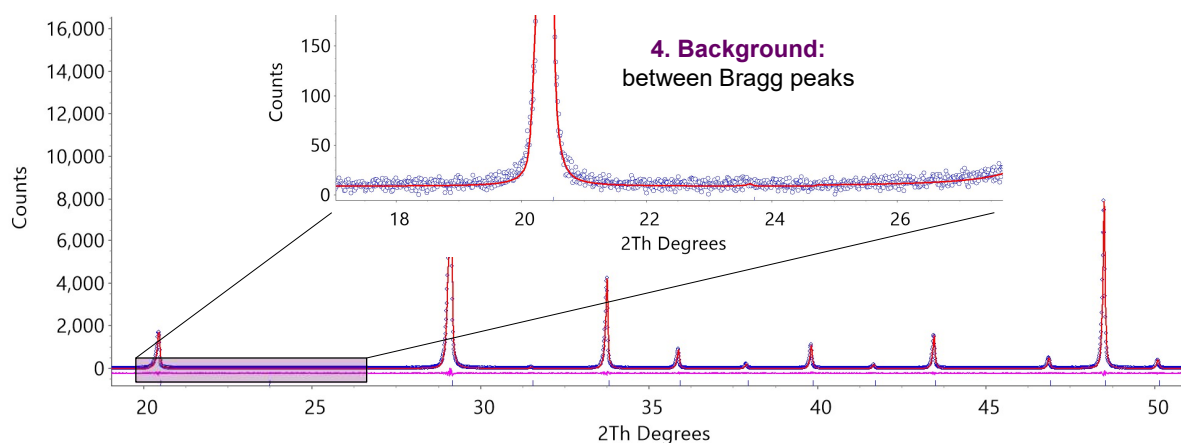
What do we have to think about in the Rietveld model?

"Rietveld refinement in Excel",
Evans and Evans, *J. Chem.
Educ.*, 2021, **98**, 495–505



What do we have to think about in the Rietveld model?

"Rietveld refinement in Excel",
Evans and Evans, *J. Chem.
Educ.*, 2021, **98**, 495–505





1. Peak position 2θ - Bragg's law

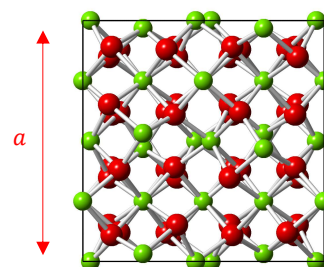
Positions:

$$\lambda = 2d_{hkl} \sin \theta_{calc} \quad \frac{1}{d_{hkl}^2} = \frac{h^2 + k^2 + l^2}{a^2}$$

$$2\theta_{obs} = 2\theta_{calc} + \Delta 2\theta$$

Depends on:

- zero point alignment error
- **sample displacement**
- sample transparency
- beam divergence
- peak asymmetry
- non-curved sample
- data reduction problems
- etc, etc



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2. Intensity equations

Intensities:

$$I_{hkl} = scale \times m \times LP \times |F_{hkl}|^2 \times other\ corrections$$

$$LP = \frac{1 + \cos^2 2\theta}{1 + \cos^2 2\theta}$$

Relates experimental intensities to scattering from unit cell

Lorentz-polarisation instrument geometry at (x_j, y_j, z_j)

$$F_{hkl} = \sum_j t_j \times occ_j \times f_j \times \exp[2\pi i(hx_j + ky_j + lz_j)]$$

$$t_j = \exp\left(\frac{-B_j \sin^2 \theta}{\lambda}\right)$$

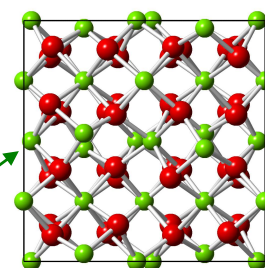
$$B_j = 8\pi^2 u^2$$

rms displacement in Å

occupancy and form factor

other corrections?

- Preferred orientation
- Texture
- Absorption
- Extinction
- Beam decay or changes
- etc



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3. Peak shapes

- Peak shapes contain fascinating information – we'll learn about this tomorrow!
- We'll (mainly) take a simple empirical approach today
- If you're doing purely structural work (depends on relative intensities) minor misfits in peak shapes might not matter
- [Exercise tip: peak shapes can be anisotropic – different for different hkl reflections (or families of reflections)]

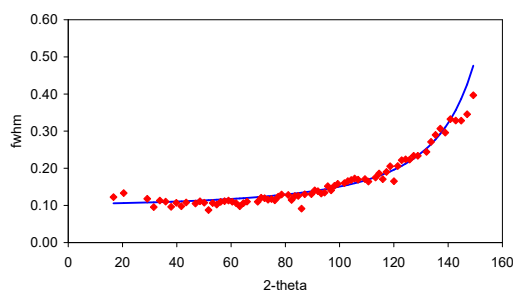
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3. Empirical peak shapes in powder diffraction

Gaussian:
$$fwhm = \left(U \tan^2 \theta + V \tan \theta + W + \frac{Z}{\cos^2 \theta} \right)^{1/2}$$

Lorentzian:
$$fwhm = X \tan \theta + \frac{Y}{\cos \theta}$$



TCHZ_Peak_Type(pku, 0.05°, pkv, -0.004°, pkw, 0.008°,
pkz, 0.03°, pkx, 0.136°, pky, 0.04°)

U, V, W, X, Y (and Z)
refineable parameters

Window Help			
Structure Microstructure Peak Type hkl			
Additional Convolutions GUI Text Rpt/Text			
Peak Type	Value	Code	Error
PV_TCHZ			
U	0.01	Refine	0
V	0.01	Refine	0
W	0.01	Refine	0
Z	0	Fix	0
X	0.01	Refine	0
Y	0	Fix	0

Peaks can be anisotropic
shape depends on hkl

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4. Background description

- Smooth function in 2-theta (polynomial, Fourier series, etc)
- Perhaps broad peaks to describe “humps”
- Perhaps a function to describe e.g. capillary or short range order effects
- Perhaps fit an experimentally measured background during the Rietveld
- [Exercise tip: don't forget point 2 on this slide!]

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How much information is in a powder pattern, what can I refine?

- Rarely enough!
- See notes for a more mathematical answer
- Bring in **restraints** and **constraints** to help refinement as “extra information”
- [Tomorrow we will use symmetry mode ideas as a different flavour of “extra information”]

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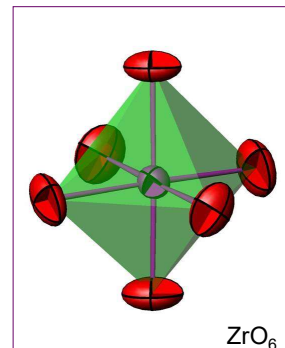
Restraints in Rietveld refinement

- You might know that a Zr-O bond in a ZrO_6 octahedron should be $\sim 2.075 \text{ \AA}$
- “extra information”
- Apply a “penalty” or “restraint” if it’s different in the model

$$\text{penalty} = (\text{model's } \text{ZrO}_{\text{dist}} - 2.075)^2$$

$$\chi^2_{\text{restraints}} = f(\text{all penalties})$$

$$\chi^2_{\text{total}} = \chi^2_{\text{data}} + \text{weight} \times \chi^2_{\text{restraints}}$$



- [What about standard uncertainties or esd’s if you do this? What do they mean?]

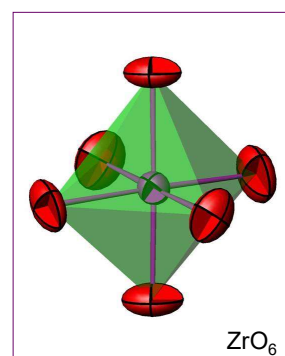
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Constraints in Rietveld refinement – e.g. rigid bodies

- You might want to force a ZrO_6 group to be a perfect octahedron
- Constraints change the least squares equations
- 21 xyz parameters become 3 positions and 3 rotations?

```
macro Octahedron(Zr1, o1, o2, o3, o4, o5, o6, 2.075)
{
  Point_for_site(s0, 0, 0, 0)
  Point_for_site(s1, r, 0, 0)
  Point_for_site(s2, -r, 0, 0)
  Point_for_site(s3, 0, r, 0)
  Point_for_site(s4, 0, -r, 0)
  Point_for_site(s5, 0, 0, r)
  Point_for_site(s6, 0, 0, -r)
}
```



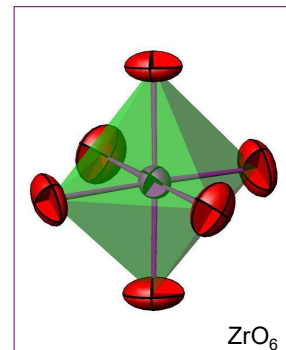
- [More detail in Sebastian’s lecture tomorrow]

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Constraints in Rietveld refinement – e.g. TLS matrices

- In a rigid group atoms ought to vibrate in related ways
- 42 u_{ij} parameters become 2 TLS parameters?

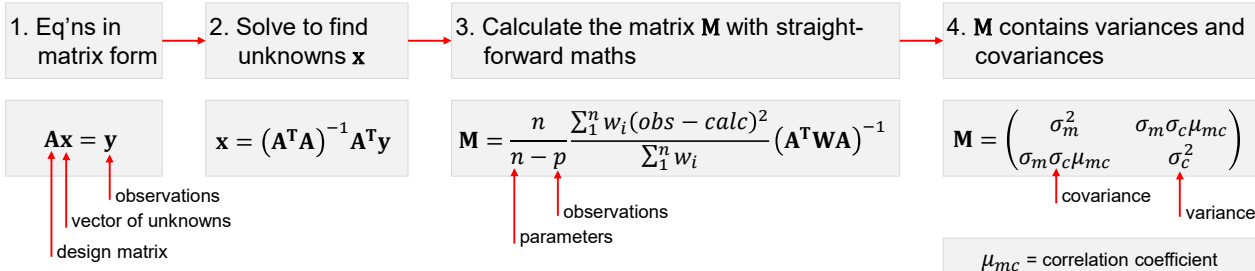


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How reliable is the model: variance covariance matrix

e.g. fit a straight line $y = mx + c$ to a set of n y_{obs} values to determine the gradient m and intercept c



Correlation matrix from software →

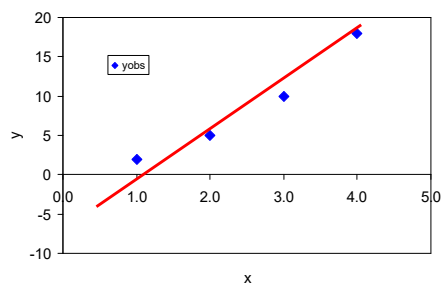
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22		
1. bkg1F871651788	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
2. bkg1F871651830	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
3. bkg1F871651808	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
4. bkg1F871651980	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
5. bkg1F871651A28	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
6. bkg1F871651A40	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
7. axial	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
8. zero	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
9. scale1F87150BAE8	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
10. lpa	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
11. bkg1F87150B0B8	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
12. x1F87150B648	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
13. bkg1F87150C208	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
14. x1F87150C118	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00		
15. y1F87150C1A8	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00	0.00	0.00		
16. x1F87150C238	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00	0.00		
17. bkg1F87150C358	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00	0.00		
18. pku	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00	0.00		
19. pku	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	0.00		
20. pku	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100		
21. pku	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100	
22. pku	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	100

There's an example in the on-line notes of how to do this by hand – tedious but good to do once in your life!

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Correlated parameters e.g. online by-hand problem



$$y = mx + c$$

$$m = 5.3(8)$$

$$c = -4.5(20)$$

$$\sigma_m = 0.79$$

$$\sigma_c = 2.17$$

$$\mu_{mc} = -0.91$$

gradient and intercept
anti-correlated

Rietveld refinement correlations

- Zero point, sample displacement, unit cell
- Wavelength, unit cell
- Occupancy, scale factor, temperature factor
- Peak shape coefficients, size, strain
- Phase fractions, temperature factors, occupancies

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Topas simplest launch mode INP file

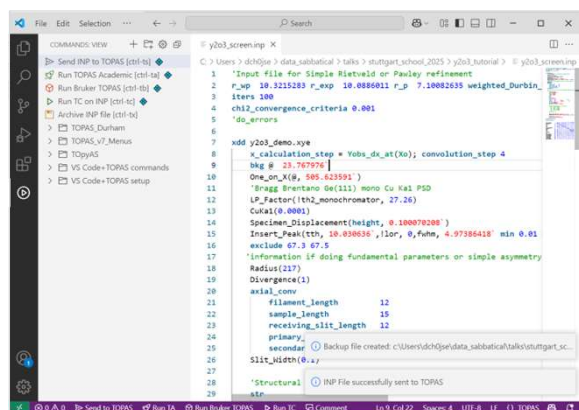
```
'Input file for Simple Rietveld
r_wp 10.3215283 r_exp 10.0886011 r_p 7.10082635 weighted_Durbin_Watson 1.98433763
gof 1.02308815
iters 100
chi2_convergence_criteria 0.001

xdd y2o3_demo.xye
x_calculation_step = Yobs_dx_at(Xo); convolution_step 4
bkg @ 0 0 0
LP_Factor(!th2_monochromator, 27.26)
CuKa1(0.0001)
str
  space_group "Ia-3"
  site Y1 x 0.25 y 0.25 z 0.25 occ Y+3 1 beq 0.25
  site Y2 x 0.97 y 0.00 z 0.25 occ Y+3 1 beq 0.25
  site O1 x 0.39 y 0.15 z 0.38 occ O-2 1 beq 0.25
  TCHZ_Peak_Type(pk_u, 0.039, pk_v, -0.02, pk_w, -0.001, !pk_z, 0.00, pk_x, 0.009, pk_y, 0.001)
```

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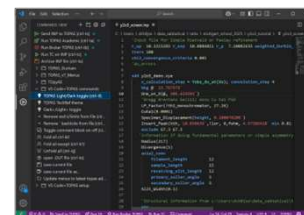


New for Stuttgart 2025: TOPAS + VS Code topas-editor



- Easier to install
- Menus simplified
- Light and dark schemes
- Easier to customise
- New commands/ways of working
- Ctrl-t keys: ta, tc, ts
- Pop up help menus
- Will have bugs!

Topas archive inp file	Ctrl+T Ctrl+X
Topas import CIF	Ctrl+T Ctrl+I
Topas insert structure	
Topas remove esds/limits from INP file	Ctrl+T Ctrl+E
Topas save and set as INP file	Ctrl+T Ctrl+S
Topas toggle comment block on or off	Ctrl+T Ctrl+/
Align-columns	
Change All Occurrences	Ctrl+F2
Refactor...	Ctrl+Shift+R
Source Action...	



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Controlling parameters in INP files

'parameters are turned on and off with a ! Symbol or using @

```
a 10.156 'fixed cell parameter with no name
a @ 10.156 'refined cell parameter with automatic name

a !lpa 10.156 'fixed cell parameter a with a name
a lpa 10.156` 'refined cell parameter a with a name

prm !prm_name 10.156 'a fixed parameter with a name
prm prm_name 10.156` 'a refined parameter with a name
```

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Quiz: TOPAS crimes/pitfalls

str

```
space_group "Ia-3"
a @ 10.598107`
b @ 10.498107`
c @ 10.698107`
a1 @ 90
be @ 90
ga @ 90
volume 1190.378`
```

str

```
space_group "Ia-3"
a lpa 10.598107`
b lpa 10.598107`
c lpa 10.598107`
a1 !alpha 90
be !beta 90
ga !gamma 90
volume 1190.378`
```

str

```
space_group "Ia-3"
a lpa 10.598107`
b = Get(a);
c = Get(a);
```

str

```
space_group "Ia-3"
Cubic(@ 10.598107`)
```

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Quiz: TOPAS crimes/pitfalls

str

```
space_group "P-3m1"
phase_name "SnS2"
a 3.6999
b 3.6999
c 6.9779
a1 90
be 90
ga 120
site Sn0      x 0.0000 y 0.0000 z 0.0000 occ Sn 1.0
site S1       x 0.3333 y 0.6666 z 0.2117 occ S 1.0

site Sn0 num_posns 1 x 0.0000 y 0.0000 z 0.0000 occ Sn 1.0
site S1  num_posns 2 x =1/3; y =2/3; z 0.2117 occ S 1.0
```

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Quiz: TOPAS crimes/pitfalls

str

```
a 4.594
b 4.594
c 2.959
al 90.
be 90.
ga 90.
space_group "P42/mnm"
site Ti1 x 0.000 y 0.000 z 0.000 occ Ti 1.0 beq 0.70000
site O1 x @ 0.306 y @ 0.306 z 0.000 occ O 1.0 beq 0.70000

site Ti1 x 0.000 y 0.000 z 0.000 occ Ti 1.0 beq 0.70000
site O1 x o1x 0.306 y o1x 0.306 z 0.000 occ O 1.0 beq 0.70000

site Ti1 x 0.000 y 0.000 z 0.000 occ Ti 1.0 beq 0.70000
site O1 x o1x 0.306 y =Get(x);:0 z 0.000 occ O 1.0 beq 0.70000
```

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Quiz: TOPAS crimes/pitfalls

```
TCHZ_Peak_Type(pku, 0.05`,pkv, -0.004`,pkw, 0.0008`,pkz, 0.003`,pkx, 0.136`,pky, 0.04`)
```

```
TCHZ_Peak_Type(pku, 0.05`,pkv, -0.004`,pkw, 0.0008`,!pkz, 0.00, pkx, 0.136`,pky, 0.04`)
```

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Quiz: TOPAS crimes/pitfalls

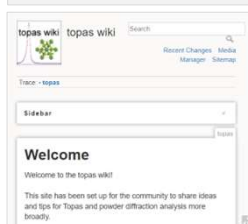
```
xdd y2o3_demo.xye
x_calculation_step = Yobs_dx_at(Xo); convolution_step 4
bkg @ 0 0 0
'Bragg Brentano Ge(111) mono Cu Ka1 PSD
LP_Factor(!th2_monochromator, 27.26)
CuKa1(0.0001)
Specimen_Displacement(height, 0.068`)
Zero_Error(@, -0.017)
...
...
```

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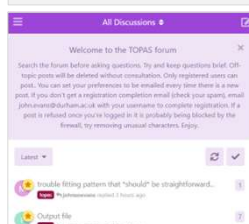


Resources?

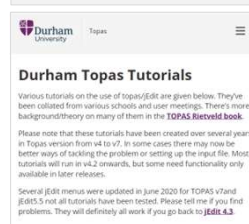
wiki site



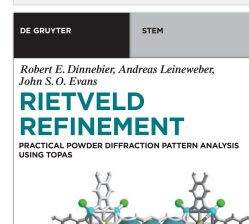
forum



web tutorials



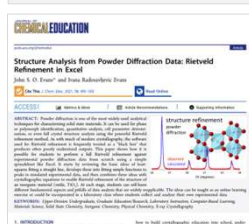
book



YouTube



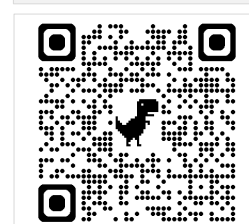
Rietveld in excel



School breaks



School page



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Exercises: individual or as a group?

- If you've never done a jEdit/VS Code TOPAS refinement before:
 - https://topas.webspace.durham.ac.uk/vscode_riet_pawley/
 - https://topas.webspace.durham.ac.uk/tutorial_tio2riet/
- If you've some experience with jEdit/VS Code:
 - https://topas.webspace.durham.ac.uk/tutorial_y2o3riet/
- If you don't have jEdit/VS Code template INP files online
- If you finish these try anything that interests you from:
 - https://topas.webspace.durham.ac.uk/topas_user_menu/
- Or try the Y_2O_3 tutorial in GSAS if you're really brave
- We will come round and help you

Stuttgart page

