

实用中子衍射晶体学
Practical Neutron Diffraction Crystallography
(II)

峰型剖面方法精修和分析结构

黄清镇

HUANG QINGZHEN

中国散裂中子源

China Spallation Neutron Source

X射线和中子粉末衍射峰的位置，强度和结构因子

Powder diffraction positions, intensity, and structural factors for XRPD & NPD

Peaks Position: $2d_{hkl} \sin \theta_{hkl} = n\lambda$

where λ is the incident beam wavelength, d and θ are the distance between successive hkl planes and Bragg angles of reflections, respectively.

Diffraction Intensity: $I_{hkl} = C|F_{hkl}|^2$

where $|F_{hkl}|^2$ is the hkl reflection amplitude of the diffracted X-ray, or neutron, or magnetic, and C is all others.

X-ray: $|F_{hkl}|^2 = |\sum f_j \exp(2\pi i (hx + ky + lz))|^2 e^{-2W}$

where f_j is the X-ray atomic scattering factor of atom j for X-ray.

Neutron: $|F_{hkl}|^2 = |\sum b_j \exp(2\pi i (hx + ky + lz))|^2 e^{-2W}$

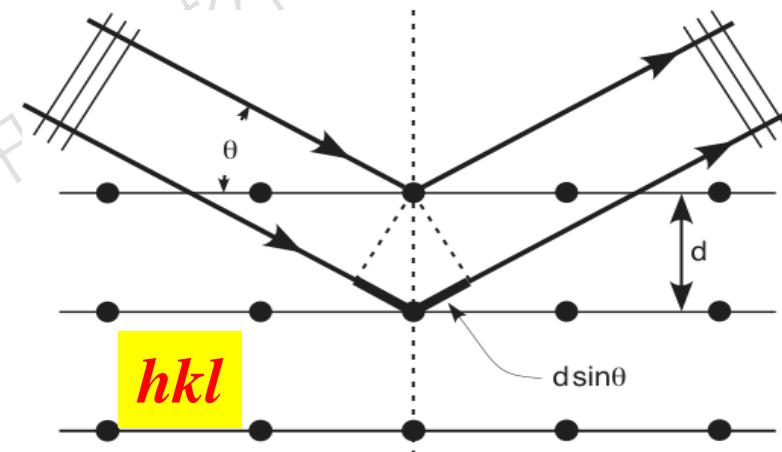
where b_j is the neutron scattering length for atom j .

Magnetic: $|F_{hkl}|^2 = |\sum q_j f_{Mj} \exp(2\pi i (hx + ky + lz))|^2 e^{-2W}$

where q_j and f_{Mj} are the magnetic interaction vector and the magnetic form factor for atom j , respectively.

X-射线衍射与中子衍射具有许多共同点。峰的位置都遵循Bragg方程：

$$2d_{hkl} \sin \theta_{hkl} = n\lambda$$



衍射峰的强度：

$$I_{hkl} = C|F_{hkl}|^2$$

式中 C 是与晶体结构无关的参数， F_{hkl} 是结构因子。

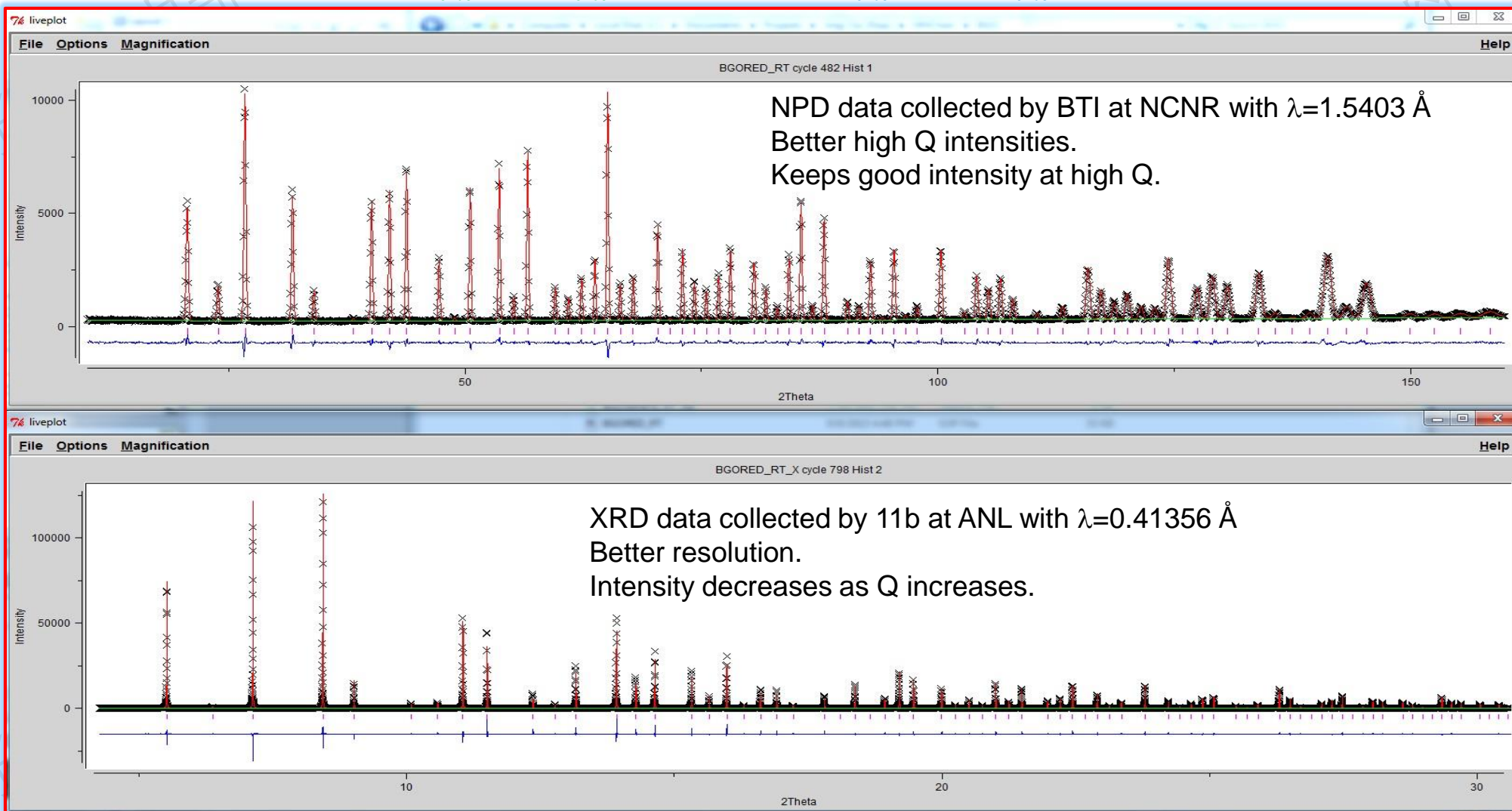
(1) 在结构因子中，除了由于元素的散射因子差异使得衍射峰强度不同外，其它与结构有关的参数都一样。

(2) 在中子衍射中，磁有序的衍射谱图是一套独立于晶体结构的衍射图，因此磁相可作为一个独立的相精化得到磁结构信息。

XRPD & NPD 是结构分析的有力的工具

$$2d_{hkl} \sin \theta_{hkl} = n\lambda$$

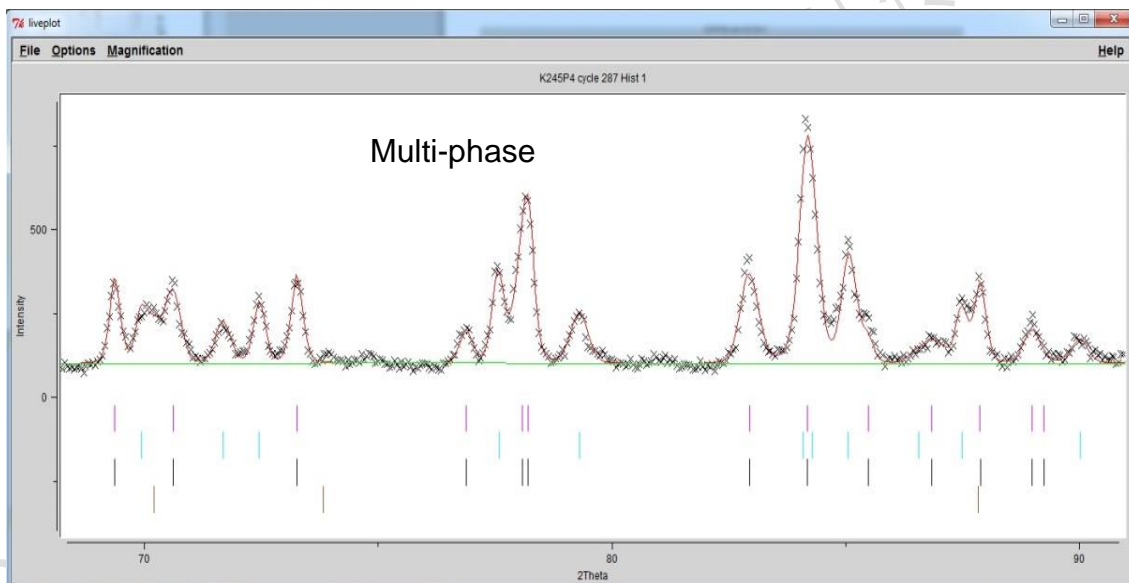
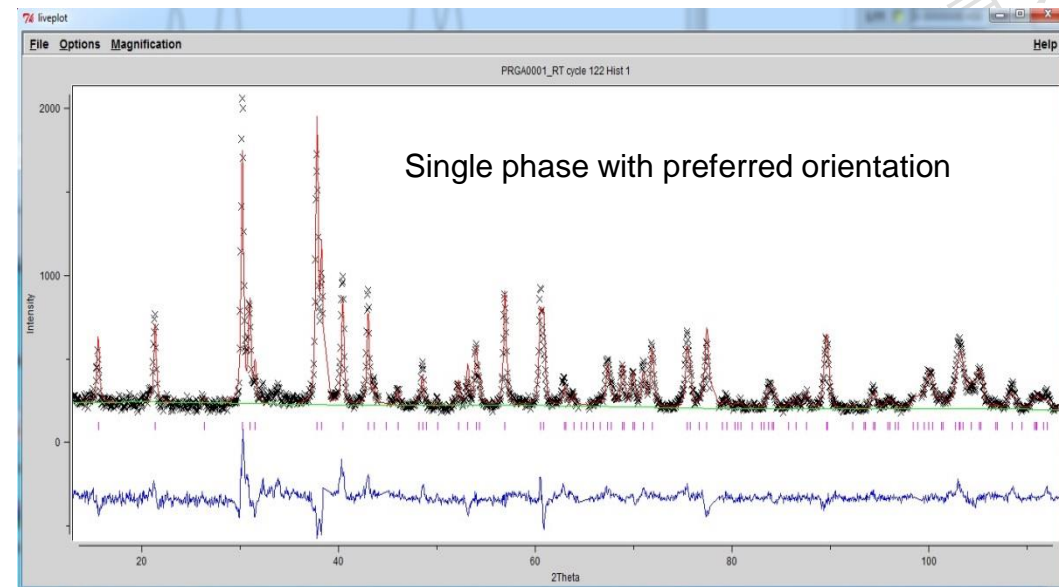
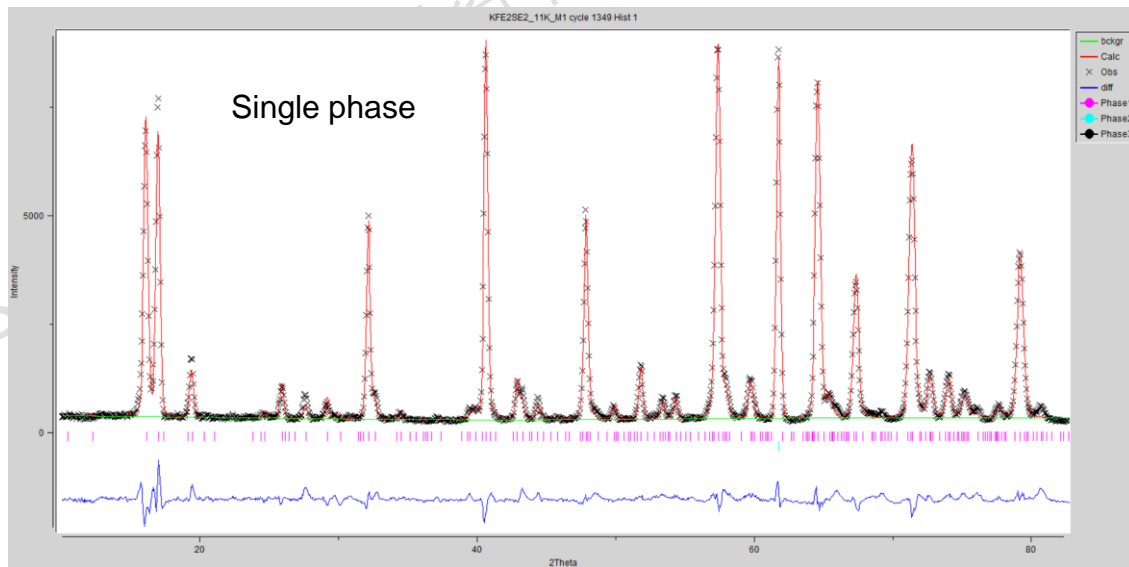
$$I_{hkl} = C|F_{hkl}|^2$$



XRD and NPD patterns for $\text{Bi}_4\text{Ge}_3\text{O}_{12}$ compound

Sample comes from Jintai Zhao's group

衍射峰重叠严重



Difficulty to get each reflection intensity
due to overlaps

- Large unit cell and low symmetry;
- Sample with multi-phase;
- Poor crystalline and particle shape;
- Defect and strain;
- Instrumental resolution
- Improper experimental condition

...

Rietveld (峰型剖面)方法分析粉末衍射数据

Rietveld (Profile) Refinement Method for the Analysis of Powder Diffraction Data



Hugo M. Rietveld and Dorothy Crowfoot Hodgkin
(She is The Nobel Prize winner in chemistry 1964).*

*Hugo M. Rietveld

The Rietveld method.

Phys. Scr. 89 (2014) 098002 (6pp)

A Profile Refinement Method for Nuclear and Magnetic Structures.

J. Appl. Cryst. 2, 65-71 (1969)

Sum of the time cited: 8720+

Line profiles of neutron powder-diffraction peaks for structure refinement.

Acta Cryst. (1967). 22, 151-152

Sum of the time cited: 1766



The Royal Swedish Academy of Sciences
has decided to award

Hugo M. Rietveld

THE GREGORI AMINOFF PRIZE

in recognition of his development of profile refinement
methods for the analysis of powder diffraction data

STOCKHOLM, MARCH 31, 1995

Kerstin Fredga
President

Carl-Olof Jacobson
Secretary General

Albinati, A., & Willis, B. T. M. *The Rietveld method. International Tables for Crystallography. Vol. C, 8.6*, 710-712(2006).

Young, A. (1993) Edited. *The Rietveld Method. International Monographs on Crystallography. 5.* Oxford University Press Inc., New York(1993).

爱明诺夫奖

格雷戈里·阿米诺夫奖 (英语: **Gregori Aminoff Prize**) 是由瑞典皇家科学院于1979年设立的国际奖项, 得名于瑞典科学家格雷戈里·阿米诺夫 (瑞典语: **Gregori Aminoff**) (1883-1947), 以奖励世界范围内在晶体学领域做出重要突出贡献的科学家, 每年颁发给不超过3人。被认为是晶体学的Nobel奖。

Chinese winner of Aminoff Prize

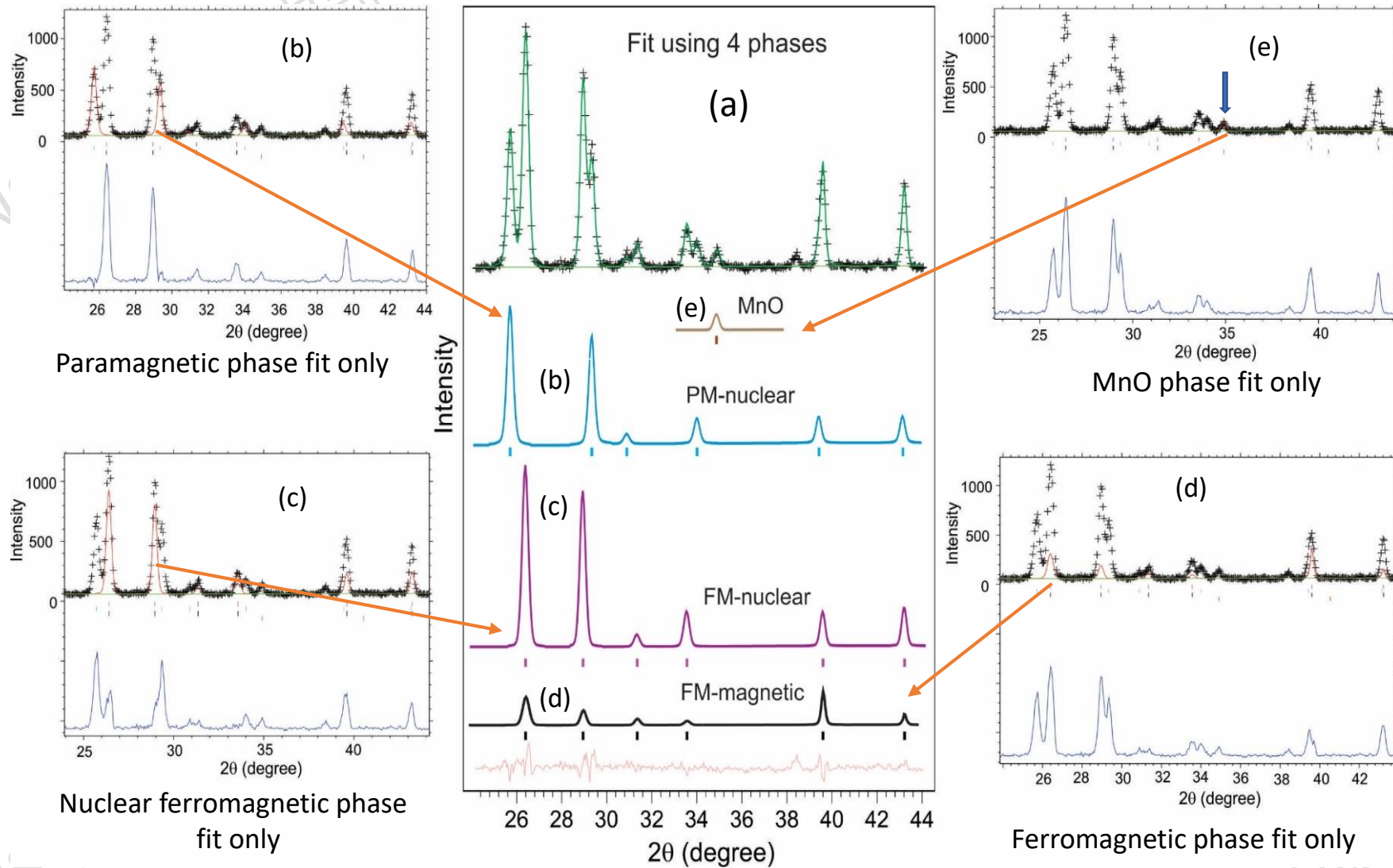
Mao Ho-kwang (毛河光)

2005, "for his pioneering research of solid materials at ultrahigh pressures and temperatures"

Shi Yigong (施一弓)

2014, "for his groundbreaking crystallographic studies of proteins and protein complexes that regulate programmed cell death"

峰型剖面精化方法定量相分析 $\text{Mn}_{1.1}\text{Fe}_{0.9}\text{P}_{0.8}\text{Ge}_{0.2}$
 Profile Refinement for Quantitative Phase Analysis of $\text{Mn}_{1.1}\text{Fe}_{0.9}\text{P}_{0.8}\text{Ge}_{0.2}$



Rietveld方法是用在已有结构模型的情况对模型进行精化，并不是用来解结构的方法。

用粉末衍射数据分析晶体结构的最大困难是如何分离部分和完全重叠的各个衍射峰强度，Hugo M. Rietveld提出用峰型剖面分析方法，改连续扫描为步进扫描收集数据，把衍射谱图数据化。有效地解决了这一困难。如图：

(1) 图 (a) 是含有4个物相的中子粉末衍射谱图。

(2) 根据这4个物相的已知模型建立4条中子粉末衍射谱图 (b, c, e, d)。

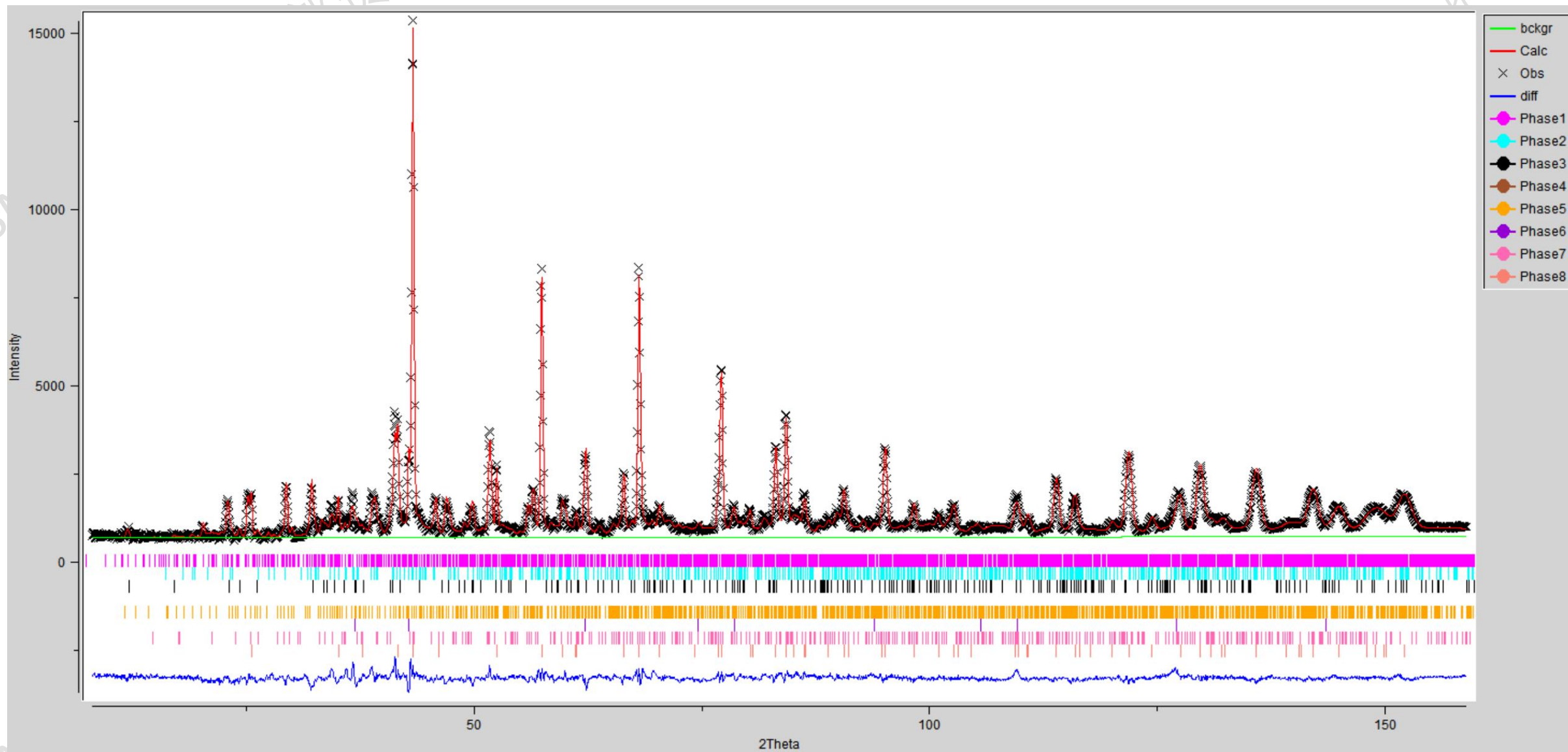
(3) 调整结构参数 (晶胞参数, 原子位置及其分布, 温度因子, 占有率等), 选择合适的峰型函数和背底函数, 以及各个相的含量, 再用最小二乘法逐步进行上述各个参数的修正, 直至计算的谱图完全拟合观察的谱图, 完成晶体结构和磁结构的精修。

粉末衍射谱图大致包含两大部分，**背底**和**衍射峰**。
 背底主要来自两部分：1) 入射光束打在**空气**，**样品盒**，**环境**，等产生；这部分的不含我们所需要的结构信息，因此应该尽量减少，来自空气的散射可通过在光路上抽真空减少空气颗粒来消除，或充氦气来减少，因为氦气颗粒小。如NIST的BT1中子谱仪在入射光路中冲氦气提高入射中子~8%；2) 来自**样品**的散射。样品的散射则主要有**康普顿**散射与样品本身的**漫散射**。**漫散射**由其样品中无序部分和晶格动力学引起。以后主要讨论与有序结构有关的衍射特征和信息提取。

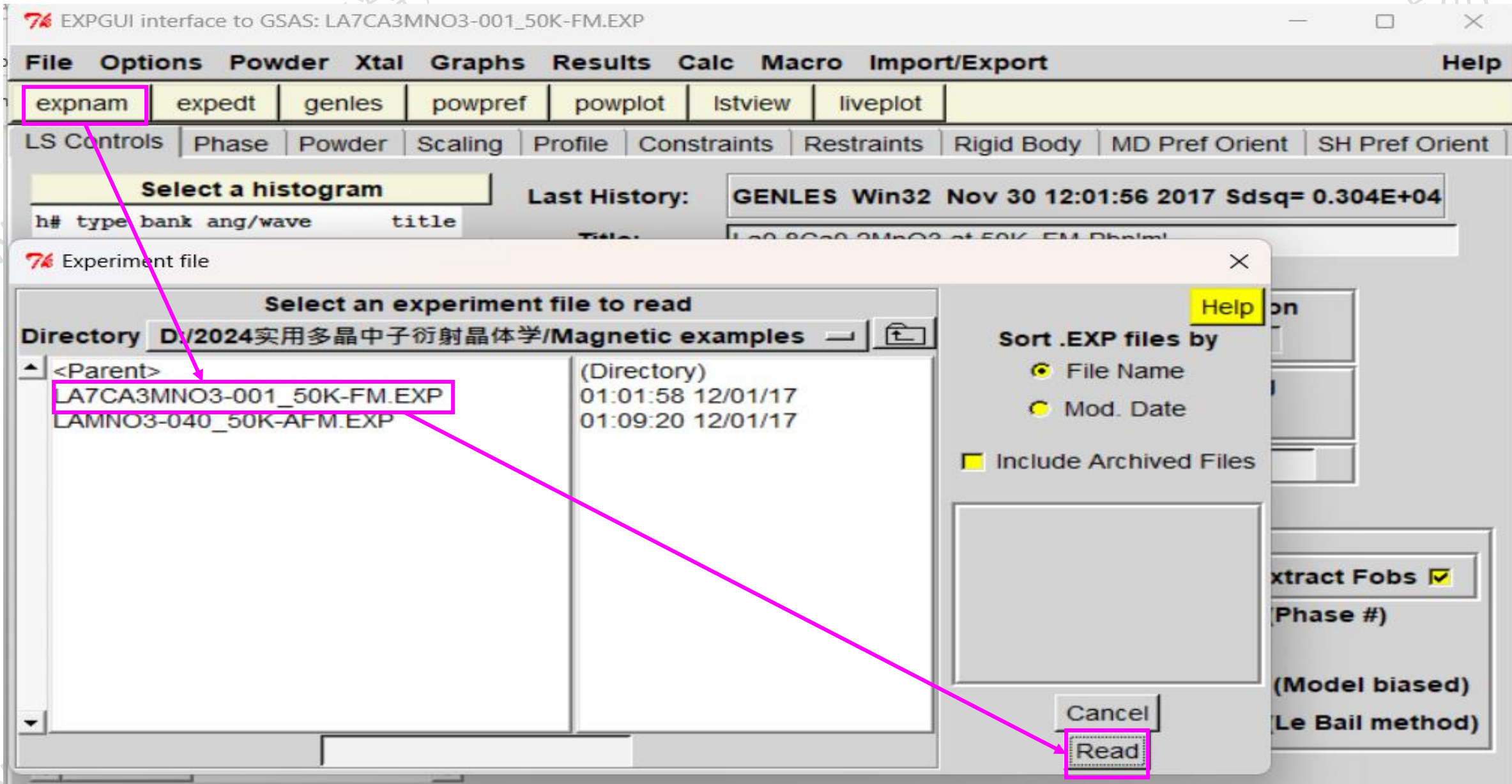


我们关注的是衍射峰所包含的结构信息。重点在：1) **峰位置**，因为它帮我们确定晶体结构中重复单元的形状和大小，对称性，以及缺陷所引起的峰位变化，如宏观应变；而**峰强**则包含结构中的化学成分，元素位置及分布，原子偏离平均位置大小（温度因子）等等。这两部分是利用衍射进行结构分析的主体。而峰形包含谱仪参数以及晶体结构不完整部分的信息和样品晶体颗粒的形状及大小。

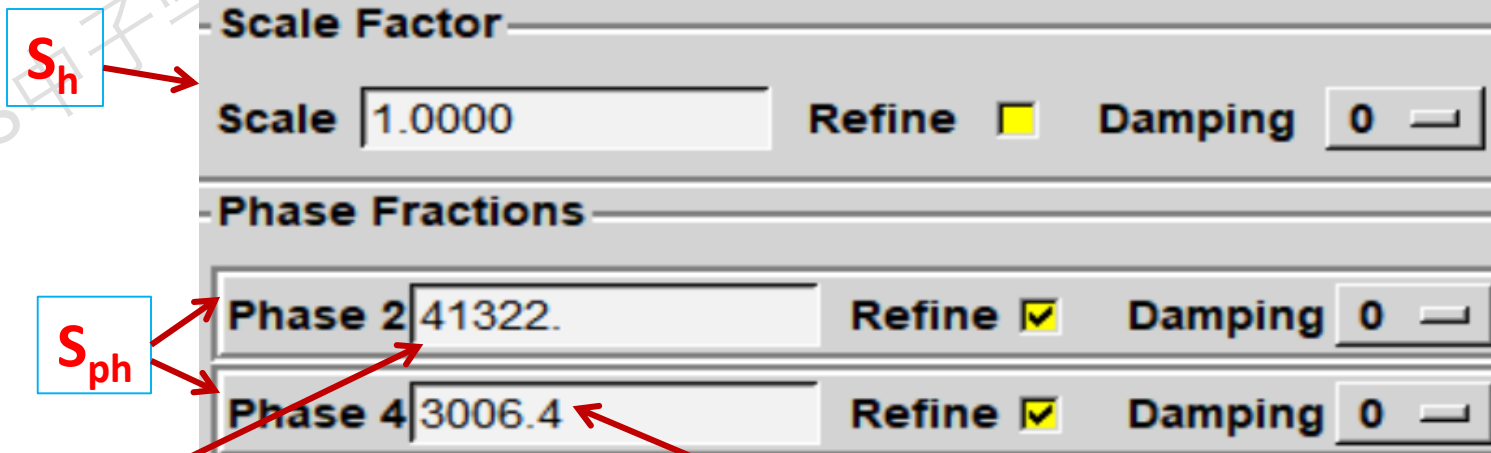
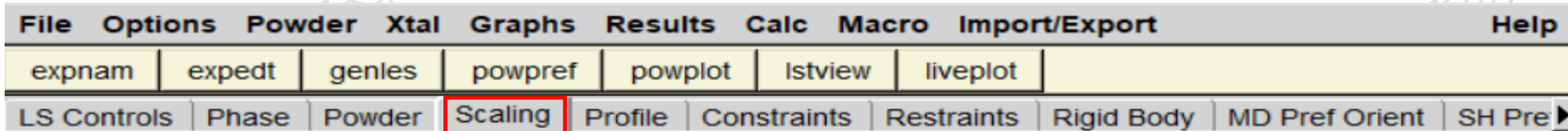
多相混合的水泥试样



GSAS-EXPGUI: expnam



GSAS-EXPGUI: Scaling



There are two types of scale factors for powder diffraction data. The histogram scale, S_h , is applied to reflections from all phases in the sample and the phase fraction scale, S_{ph} , is applied only to reflections from the p -th phase. These latter scales can be used for quantitative phase analysis for powder mixtures.

Phase/element fractions for phase no. 2

Hist Elem: 1 1 PNC
Fraction : 41322.0
Sigmas : 0.00000
Shift/esd: 0.00
Wt. Frac.: 0.95605
Sigmas : 0.00000

Phase/element fractions for phase no. 4

Hist Elem: 1 1 PNC
Fraction : 3006.41
Sigmas : 76.4528
Shift/esd: 0.00
Wt. Frac.: 0.43946E-01
Sigmas : 0.106842E-02

GSAS-EXPGUI: Powder

The screenshot shows the 'Powder' menu in the GSAS-EXPGUI software. The menu items are: LS Controls, Phase, Powder, Scaling, Profile, Constraints, Restraints, Rigid Body, MD Pref Orient, and SH Pre. The 'Powder' menu is open, showing a table with one histogram entry and several configuration panels on the right. The configuration panels include 'Background', 'Diffractometer Constants', and 'Absorption/Reflectivity Correction'. The 'Background' panel shows 'Function type 2 (12 terms)', 'Refine background' checked, and 'Damping' set to 0. The 'Diffractometer Constants' panel shows 'Refine wave' unchecked with 'wave' set to 1.5403000, and 'Refine zero' checked with 'Zero' set to -2.03475. The 'Absorption/Reflectivity Correction' panel shows 'Model #0, value: 0.0000000' and 'Refine Abs./Refl.' unchecked with 'Damping' set to 0. At the bottom of the menu, there are four buttons: 'Add New Histogram', 'Set Data Limits & Excluded Regions', 'Set Histogram Use Flags', and 'Edit $\Delta f'$ and $\Delta f''$ '. A blue box on the left contains a numbered list of five steps, with arrows pointing from the list to the corresponding buttons and menu items.

h#	type	bank	ang/wave	title
1	NC	1	1.54030	1a7c3 La0.7Ca0.3Mn

Background

Function type 2 (12 terms) Edit Background

Refine background Damping 0

Diffractometer Constants

Refine wave wave 1.5403000 Damping 0

Refine zero Zero -2.03475

Absorption/Reflectivity Correction

Model #0, value: 0.0000000

Refine Abs./Refl. Damping 0 Edit Abs./Refl.

Add New Histogram Set Data Limits & Excluded Regions Set Histogram Use Flags Edit $\Delta f'$ and $\Delta f''$

1. Add New histogram;
2. Set Histogram use flag;
3. Set Data Limits & Excluded Regions;
4. Edit background;
5. Edit Abs./Reff.

GSAS-EXPGUI: Phase

A3MNO3-001_50K-FM.EXP

File Options Powder Xtal Graphs Results Calc Macro Import/Export Help

expnam | expedt | genes | powpref | powplot | lstview | liveplot

LS Controls | **Phase** | Powder | Scaling | Profile | Constraints | Restraints | Rigid Body | MD Pref Orient | SH Pref Orient

Phase: 1 title: La_{0.8}Ca_{0.2}MnO₃ at 50K, FM-Pbn'm'

a 5.480045 b 5.465816 c 7.725897

α 90.0000 β 90.0000 γ 90.0000

* name	type	ref/damp	fractional coordinates			Mult	Occupancy	Uiso
1 La	LA	X0 U0 F0	-0.004805	0.022661	0.250000	4	0.7740	0.00403
0			0.500000	0.000000	0.000000	4	1.0000	0.00308
0			0.064748	0.491805	0.250000	4	1.0000	0.00627
0			0.723533	0.275049	0.033678	8	1.0000	0.00545
0			-0.004805	0.022661	0.250000	4	0.2260	0.00403

X U F

1. Replace phase;
2. Add new phase;
3. Edit cell parameters;
4. Add new atoms;
5. Modify atoms
6. Edit x , y , z , F , and U ;
7. Edit refinement flag.

GSAS-EXPGUI: Phase-Replace

76 EXPGUI interface to GSAS: LA7CA3MNO3-001_50K-FM.EXP

File Options Powder Xtal Graphs Results Calc Macro Import/Export Help

expnam expedt genes powpref powplot lstview liveplot

LS Controls Phase Powder Scaling Profile Constraints Restraints Rigid Body MD Pref Orient SH Pref Orient

Phase: 1 Replace title: La_{0.8}Ca_{0.2}MnO₃ at 50K, FM-Pbn'm'

76 Replace phase 1

Replacing phase #1

Current Space Group: P b n m

New Space Group:

a	5.480045	b	5.465816	c	7.725897
α	90.0000	β	90.0000	γ	90.0000

Keep atoms in phase
 Delete current atoms

Continue Cancel Import phase from: Crystallographic Information File (CIF)

Refine Cell
Cell damping 0

Add New Atoms
0 Modify Atoms

- ✓ PowderCell .CEL file
- ✓ Crystallographic Information File (CIF)
- GSAS .EXP file
- Platon .spf file
- MSI .xtl file

GSAS-EXPGUI: Phase-Add phase

EXPGUI interface to GSAS: LA7CA3MNO3-001_50K-FM.EXP

File Options Powder Xtal Graphs Results Calc Macro Import/Export Help

expnam expedt genes powpref powplot lstview liveplot

LS Controls **Phase** Powder Scaling Profile Constraints Restraints Rigid Body MD Pref Orient SH Pref Orient

Phase: 1 Replace title: La_{0.8}Ca_{0.2}MnO₃ at 50K, FM-Pbn'm'

Add Phase Magnetic & Nuclear a 5.480045 b 5.465816 c 7.725897 Edit Refine Cell
Cell damping 0

add new phase

Adding phase #2

Phase title:

Space Group:

Cell Type Any

a b c
 α 90. β 90. γ 90.

Add Cancel Help Import phase from: Crystallographic Information File (CIF)

1 La
2 Mn
3 O1
4 O2
5 Ca

0 Add New Atoms
Modify Atoms

✓ Crystallographic Information File (CIF)
PowderCell .CEL file
GSAS .EXP file
Platon .spf file
MSI .xtl file

GSAS-EXPGUI: Powder-Add New Histogram

The screenshot displays the GSAS-EXPGUI software interface. The main menu bar includes File, Options, Powder, Xtal, Graphs, Results, Calc, Macro, Import/Export, and Help. Below the menu bar, there are several sub-menus: expnam, expedt, genes, powpref, powplot, Istview, and liveplot. The 'Powder' sub-menu is highlighted with a pink box, and a pink arrow points from it to the 'Add New Histogram' dialog box.

The 'Add New Histogram' dialog box is open, showing the following fields and options:

- Adding a new histogram** (with a for Dummy Histogram)
- Data file:** [Text Field] [Select File] [Import CIF]
- Select bank** (with a dropdown menu)
- Instrument Parameter file:** [Text Field] [Select File] [Edit file]
- Select set** (with a dropdown menu)
- Usable data limit:** [Text Field]
- Radio buttons:** d-min, Q-max, TOF-min, 2-Theta Max
- Buttons:** Run RAWPLOT, Add, Cancel, Add multiple banks, Help

The background window shows a table with columns: h#, type, bank, ang/wave, title, and Background. The 'Background' section includes buttons for Edit Background, Damping (0), and Edit Abs./Refl. (0).

GSAS-EXPGUI: Powder-Set Data Limits and Excluded Regions

File Options Powder Xtal Graphs Results Calc Macro Import/Export Help

expnam expedt genes powpref powplot lstview liveplot

LS Controls Phase **Powder** Scaling Profile Constraints Restraints Rigid Body MD Pref Orient SH Pre

Selected

h#	type	bank	ang/
1	NC	1	1.5

LA7CA3MNO3-001_50K-FM-FYB... 237 Hz 14

600
400
200
10000
5000
0

36.5 37 37.5
2-Theta

Excluded Region Editing

Mouse click action	zoom	Add region	Delete region	Save & Finish			
Excluded Regions	<10.000	37.100 to 37.600	53.350 to 54.150	66.950 to 67.650	79.600 to 79.750	103.000 to 103.750	>162.026

Help

Background

Sampling

Edit Abs./Ref.

Add New Histogram **Set Data Limits & Excluded Regions** Set Histogram Use Flags Edit Δ^* and Δ°

Set background function and number of coefficients

The screenshot shows the GSAS-EXPGUI interface with the 'Powder' tab selected. The 'Background' section is active, showing 'Function type 2 (3 terms)' and 'Refine background' checked. The 'Edit Background' dialog is open, showing 'Setting background terms for histogram 2' with 'Function type' set to 2 and 'Number of terms' set to 3. A list of function types is shown on the left, with '2 - Cosine Fourier series' selected.

Select a histogram

h#	type	bank	ang/wave	title
2	NC	1	1.54030	Automatically generated file

Background

Function type 2 (3 terms) **Edit Background**

Refine background Damping 0

Edit Background

Setting background terms for histogram 2 **Fit Background Graphically**

Function type **2** Number of terms **3**

1 | 1.0 2 | 0.0 3 | 0.0

Function Types List:

- 1 - Shifted Chebyshev
- ✓ 2 - Cosine Fourier series
- 4 - Power series in $Q^{2n}/n!$
- 5 - Power series in $n!/Q^{2n}$
- 6 - Power series in $Q^{2n}/n!$ and $n!/Q^{2n}$
- 7 - Linear interpolation function
- 8 - Reciprocal interpolation function

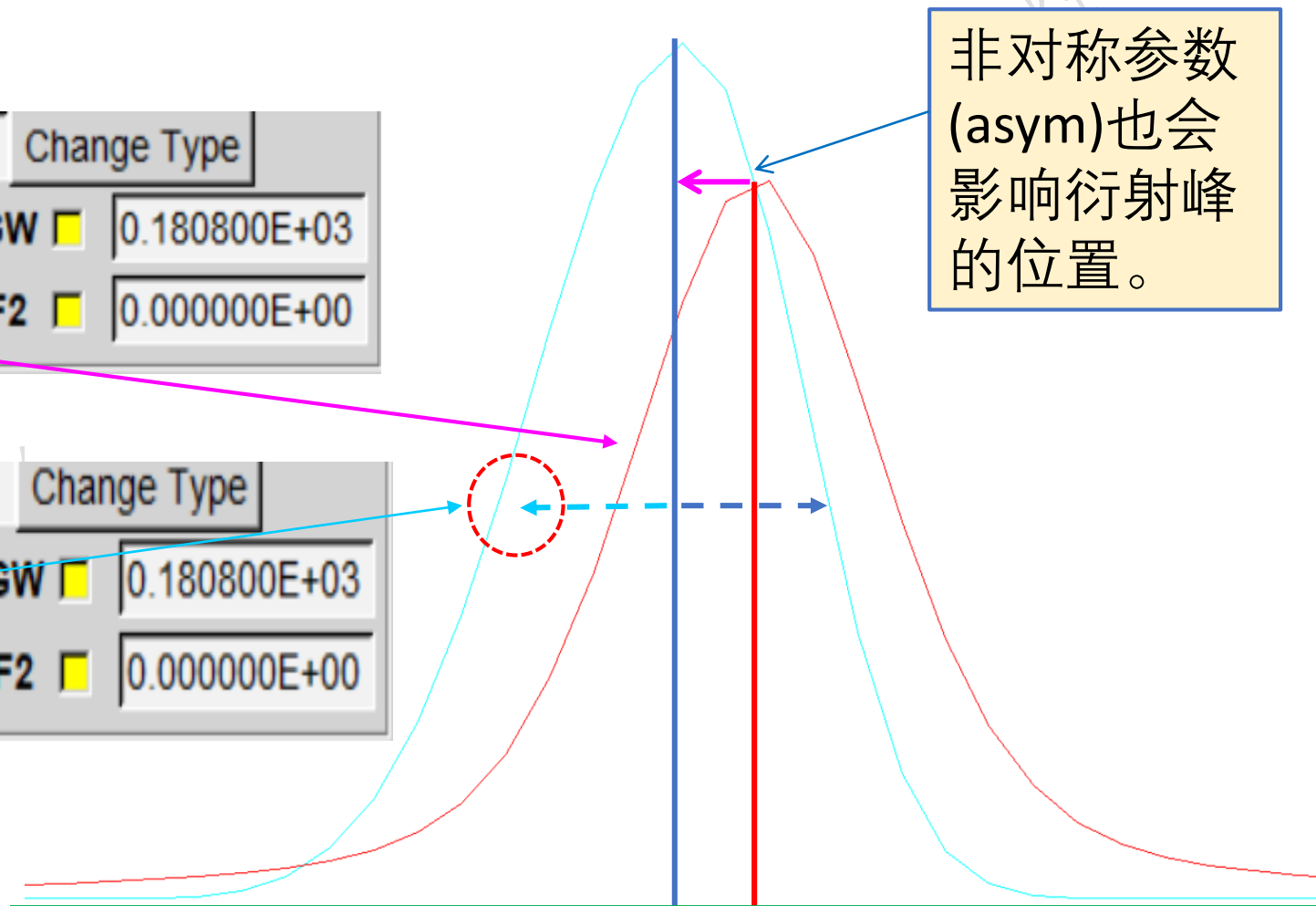
CW profile function type 1

Gaussian function

CW-Profile

Damping	0	Peak cutoff	0.00500	Change Type	
GU	<input type="checkbox"/> 0.239700E+03	GV	<input type="checkbox"/> -0.298200E+03	GW	<input type="checkbox"/> 0.180800E+03
asym	<input type="checkbox"/> 0.000000E+01	F1	<input type="checkbox"/> 0.000000E+00	F2	<input type="checkbox"/> 0.000000E+00

Damping	0	Peak cutoff	0.00500	Change Type	
GU	<input type="checkbox"/> 0.239700E+03	GV	<input type="checkbox"/> -0.298200E+03	GW	<input type="checkbox"/> 0.180800E+03
asym	<input type="checkbox"/> 20.000000	F1	<input type="checkbox"/> 0.000000E+00	F2	<input type="checkbox"/> 0.000000E+00



Rietveld and others (H.M. Rietveld, *J. Appl. Cryst.*, **2**, 65-71, 1969; Cooper & Sayer, *J. Appl. Cryst.*, **8**, 615-618, 1975; & Thomas, *J. Appl. Cryst.*, **10**, 12-13, 1977).

GSAS-EXPGUI: Profile

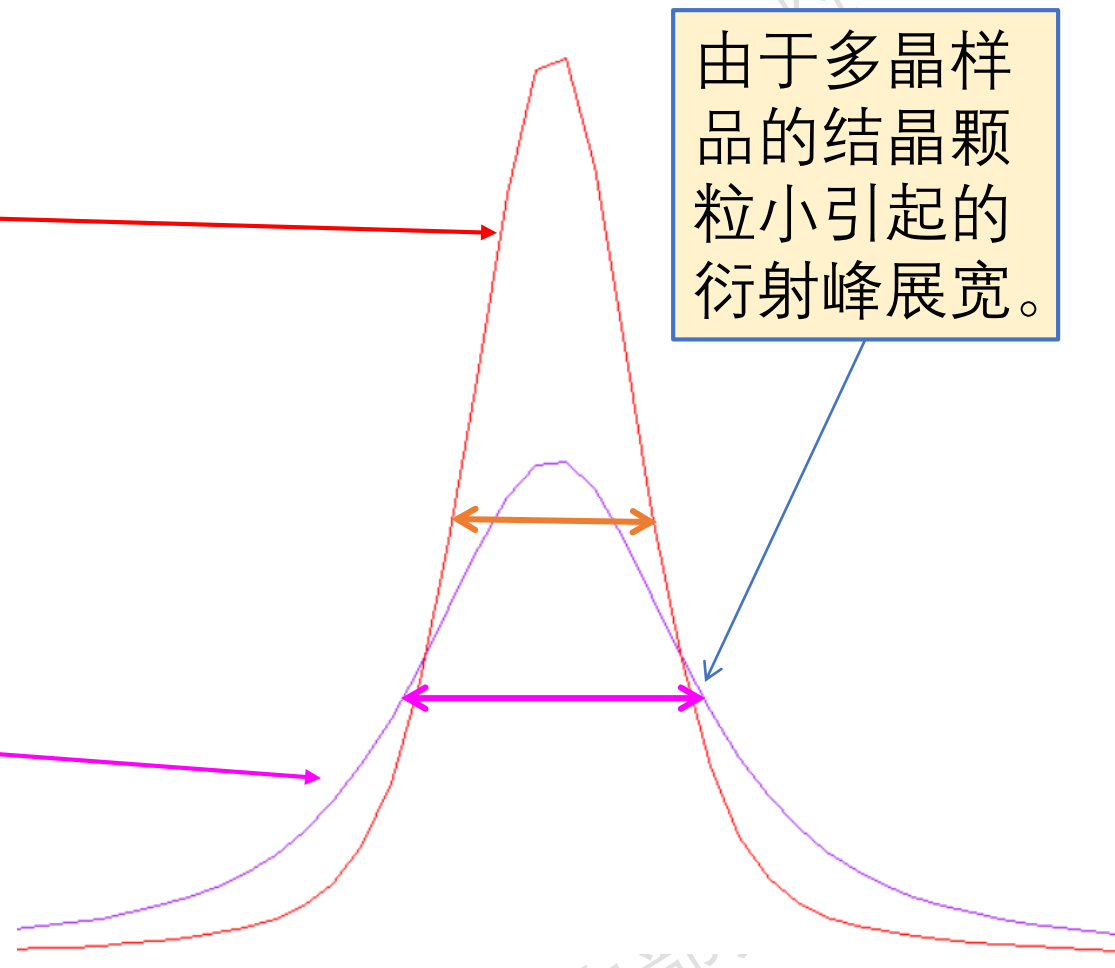
CW profile function type 2

Multi-term Simpson's rule integration

CW-Profile

Damping 0		Peak cutoff 0.00500		Change Type				
GU	<input type="checkbox"/>	239.700000	GV	<input type="checkbox"/>	-298.200000	GW	<input type="checkbox"/>	180.800000
LX	<input type="checkbox"/>	0.000000	LY	<input type="checkbox"/>	0.000000	trns	<input type="checkbox"/>	0.000000
asym	<input type="checkbox"/>	0.000000	shft	<input type="checkbox"/>	0.000000	GP	<input type="checkbox"/>	0.000000
stec	<input type="checkbox"/>	0.000000	ptec	<input type="checkbox"/>	0.000000	sfec	<input type="checkbox"/>	0.000000
L11	<input type="checkbox"/>	0.000000	L22	<input type="checkbox"/>	0.000000	L33	<input type="checkbox"/>	0.000000
L12	<input type="checkbox"/>	0.000000	L13	<input type="checkbox"/>	0.000000	L23	<input type="checkbox"/>	0.000000

Damping 0		Peak cutoff 0.00500		Change Type				
GU	<input type="checkbox"/>	239.700000	GV	<input type="checkbox"/>	-298.200000	GW	<input type="checkbox"/>	180.800000
LX	<input checked="" type="checkbox"/>	40.000000	LY	<input type="checkbox"/>	0.000000	trns	<input type="checkbox"/>	0.000000
asym	<input type="checkbox"/>	0.000000	shft	<input type="checkbox"/>	0.000000	GP	<input type="checkbox"/>	0.000000
stec	<input type="checkbox"/>	0.000000	ptec	<input type="checkbox"/>	0.000000	sfec	<input type="checkbox"/>	0.000000
L11	<input type="checkbox"/>	0.000000	L22	<input type="checkbox"/>	0.000000	L33	<input type="checkbox"/>	0.000000
L12	<input type="checkbox"/>	0.000000	L13	<input type="checkbox"/>	0.000000	L23	<input type="checkbox"/>	0.000000



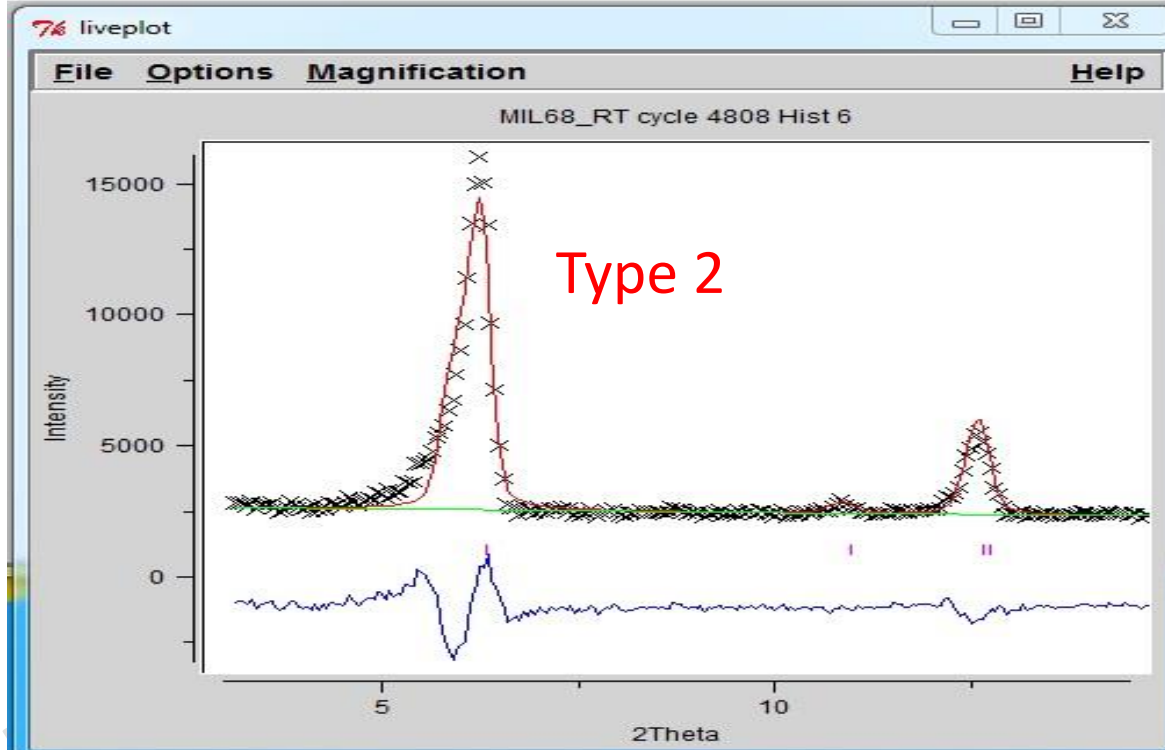
C.J. Howard (J. Appl. Cryst., **15**, 615-620, 1982) of the pseudo-Voigt, $F(\Delta T)$, described by Thompson, et al. and used above in the second and third functions for TOF data.

The third CW profile function is a similar variation on the pseudo-Voigt function used in the second function. However, it uses a much more successful description of the reflection asymmetry due to axial described by Finger, Cox & Jephcoat, J. Appl. Cryst., 27,892-900 (1994) as an implementation of the peak shape function described by Van Laar & Yelon, J. Appl. Cryst., 17, 47-54 (1984).

Hist 6 -- Phase 1 (type 2)

Damping 0 Peak cutoff 0.00100 Change Type

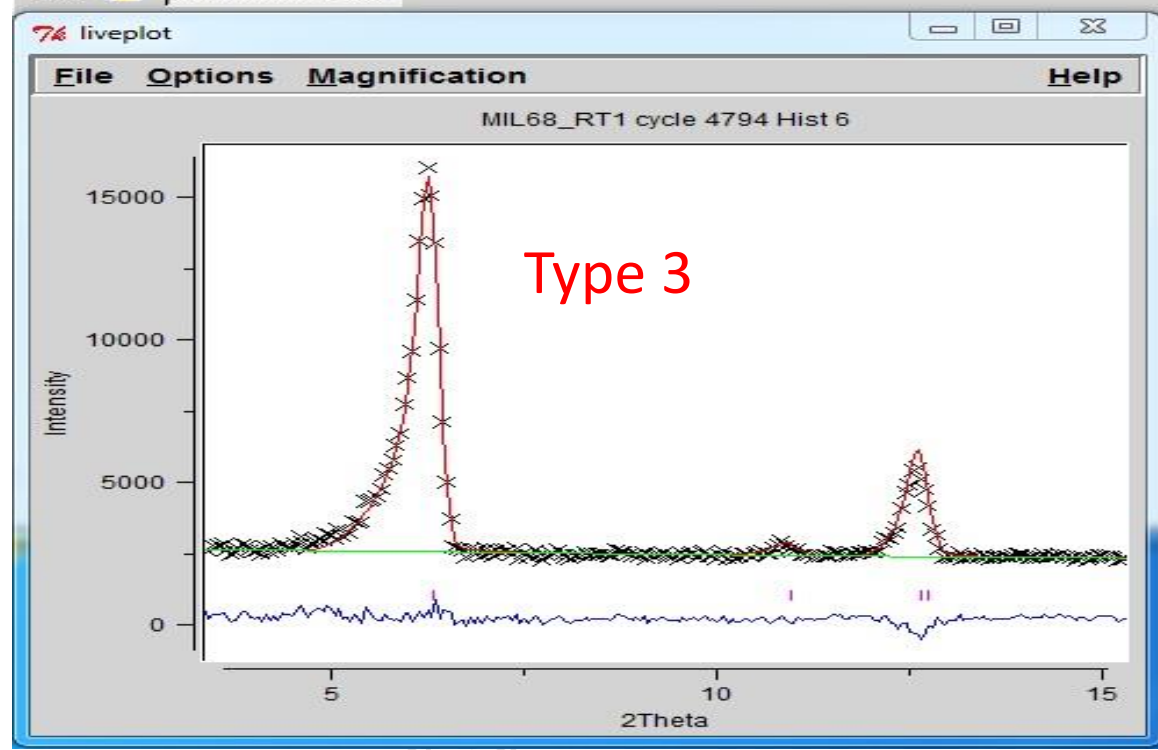
GU <input checked="" type="checkbox"/>	0.668891E+03	GV <input checked="" type="checkbox"/>	-0.213626E+03	GW <input checked="" type="checkbox"/>	0.770997E+02
LX <input checked="" type="checkbox"/>	0.810699E+01	LY <input checked="" type="checkbox"/>	0.201073E+02	trns <input type="checkbox"/>	0.000000E+00
asym <input checked="" type="checkbox"/>	0.652199E+01	shft <input type="checkbox"/>	0.000000E+00	GP <input type="checkbox"/>	0.000000E+00
stec <input type="checkbox"/>	0.000000E+00	ptec <input type="checkbox"/>	0.000000E+00	sfec <input type="checkbox"/>	0.000000E+00
L11 <input type="checkbox"/>	0.000000E+00	L22 <input type="checkbox"/>	0.000000E+00	L33 <input type="checkbox"/>	0.000000E+00
L12 <input type="checkbox"/>	0.000000E+00	L13 <input type="checkbox"/>	0.000000E+00	L23 <input type="checkbox"/>	0.000000E+00



Hist 6 -- Phase 1 (type 3)

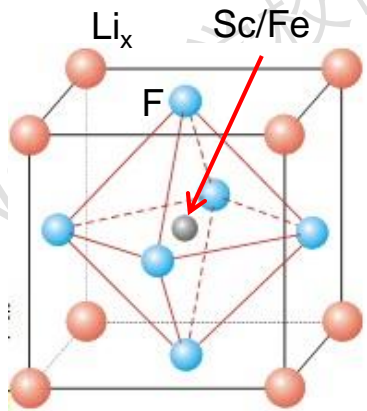
Damping 0 Peak cutoff 0.00200 Change Type

GU <input type="checkbox"/>	0.512608E+03	GV <input type="checkbox"/>	-0.306205E+03	GW <input type="checkbox"/>	0.122921E+03
GP <input type="checkbox"/>	0.000000E+00	LX <input type="checkbox"/>	0.000000E+00	LY <input type="checkbox"/>	0.413402E+02
S/L <input type="checkbox"/>	0.400000E-01	H/L <input type="checkbox"/>	0.300000E-01	trns <input type="checkbox"/>	0.000000E+00
shft <input type="checkbox"/>	0.000000E+00	stec <input type="checkbox"/>	0.000000E+00	ptec <input type="checkbox"/>	0.000000E+00
sfec <input type="checkbox"/>	0.000000E+00	L11 <input type="checkbox"/>	0.000000E+00	L22 <input type="checkbox"/>	0.000000E+00
L33 <input type="checkbox"/>	0.000000E+00	L12 <input type="checkbox"/>	0.000000E+00	L13 <input type="checkbox"/>	0.000000E+00
L23 <input type="checkbox"/>	0.000000E+00				

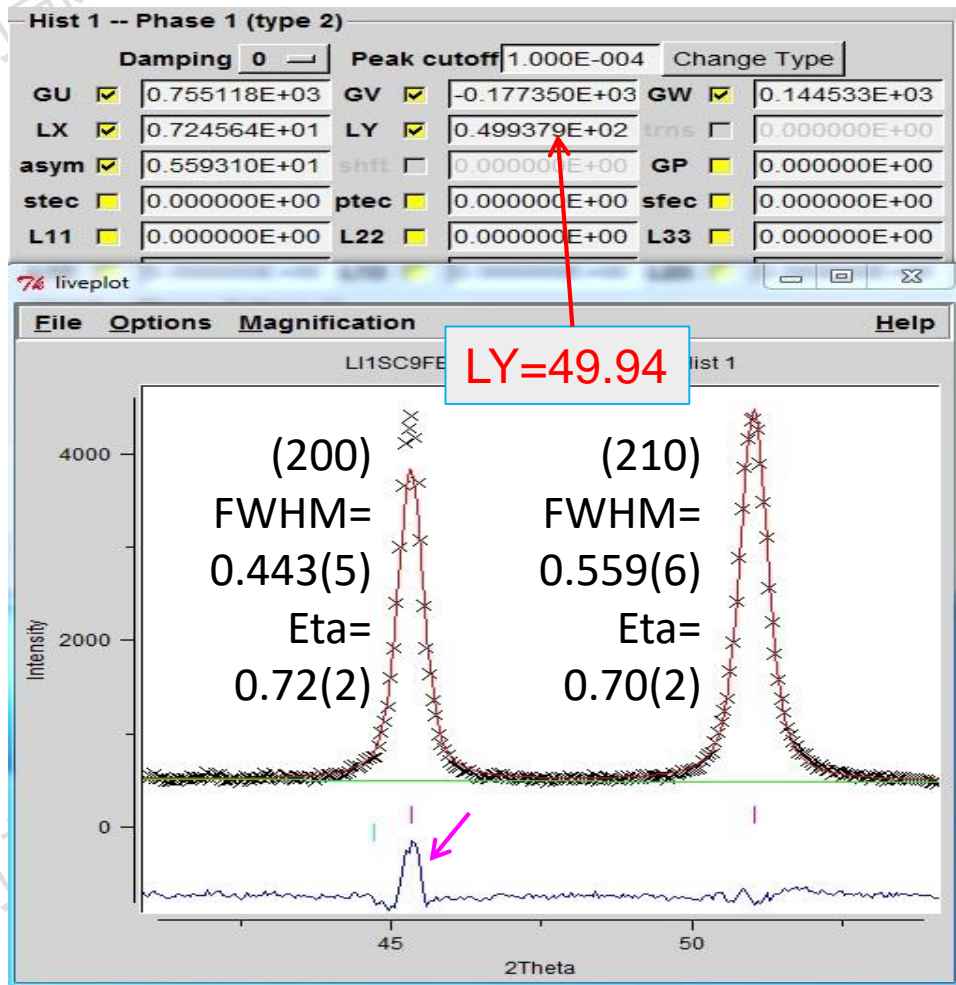


CW profile function type 4

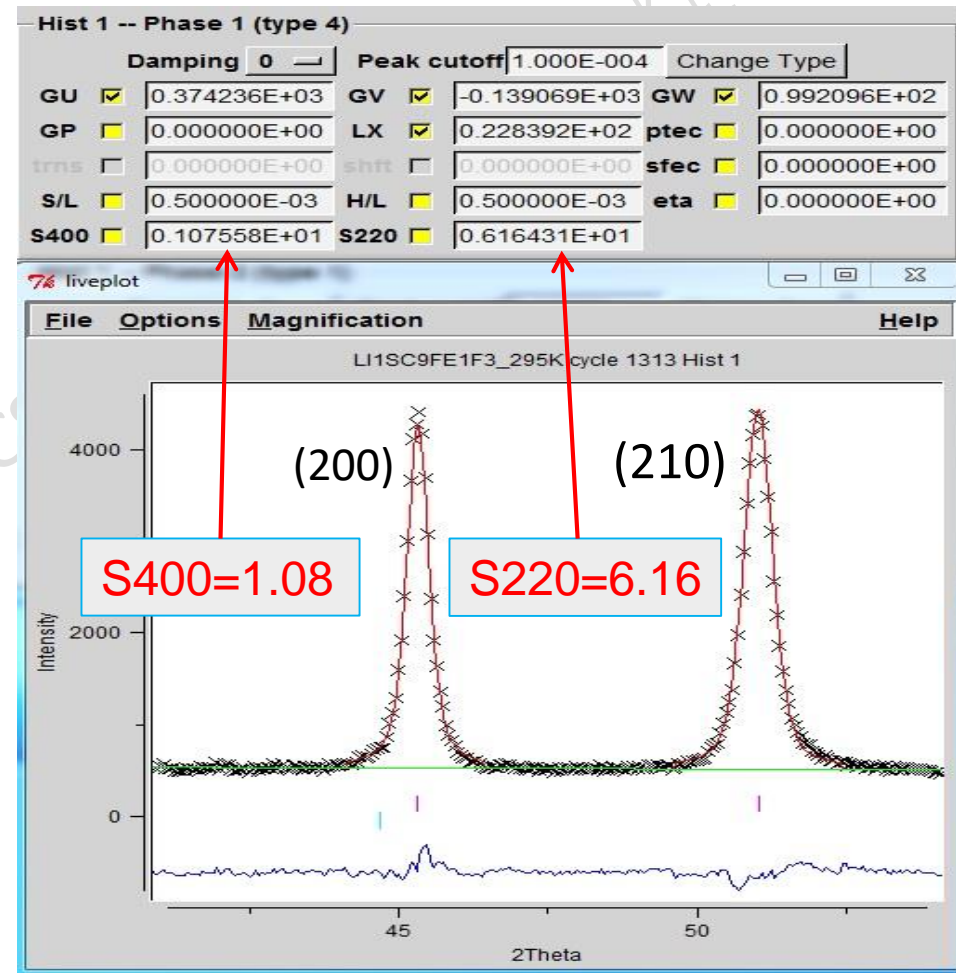
Multi-term Simpson's rule integration



Ionic radius	
Li ⁺ :	0.60
Sc ³⁺ :	0.81
Fe ³⁺ :	0.64
Fe ²⁺ :	0.76
F ⁻ :	1.36



Fit by profile #2, isotropic strain broadening



Fit by profile #4, anisotropic strain broadening

CW profile function type 5

The fifth CW profile function is a modification of the third function to include generalized macroscopic strain. In place of the six γ_{ij} coefficients used in that function to describe an empirical anisotropic microstrain broadening there are a maximum of six δ_{ij} coefficients to describe an offset of each reflection position. This profile function thus becomes useful for refinements with multiple data sets taken at a, for example, range of temperatures where the δ_{ij} can describe the effect of thermal expansion on the lattice parameters. In this case set the $\delta_{ij}=0$ for one histogram and allow them to vary for the others.

Hist 2 -- Phase 1 (type 5)

Damping Peak cutoff

GU <input type="checkbox"/>	<input type="text" value="239.700000"/>	GV <input type="checkbox"/>	<input type="text" value="-298.200000"/>	GW <input type="checkbox"/>	<input type="text" value="180.800000"/>
GP <input type="checkbox"/>	<input type="text" value="0.000000"/>	LX <input type="checkbox"/>	<input type="text" value="0.000000"/>	LY <input type="checkbox"/>	<input type="text" value="0.000000"/>
S/L <input type="checkbox"/>	<input type="text" value="0.000000"/>	H/L <input type="checkbox"/>	<input type="text" value="0.000000"/>	trns <input type="checkbox"/>	<input type="text" value="0.000000"/>
shft <input type="checkbox"/>	<input type="text" value="0.000000"/>	stec <input type="checkbox"/>	<input type="text" value="0.000000"/>	ptec <input type="checkbox"/>	<input type="text" value="0.000000"/>
sfec <input type="checkbox"/>	<input type="text" value="0.000000"/>	D1 <input type="checkbox"/>	<input type="text" value="0.000000"/>	D2 <input type="checkbox"/>	<input type="text" value="0.000000"/>
D3 <input type="checkbox"/>	<input type="text" value="0.000000"/>	D4 <input type="checkbox"/>	<input type="text" value="0.000000"/>	D5 <input type="checkbox"/>	<input type="text" value="0.000000"/>
D6 <input type="checkbox"/>	<input type="text" value="0.000000"/>				

type 5

Hist 2 -- Phase 1 (type 3)

Damping Peak cutoff

GU <input type="checkbox"/>	<input type="text" value="239.700000"/>	GV <input type="checkbox"/>	<input type="text" value="-298.200000"/>	GW <input type="checkbox"/>	<input type="text" value="180.800000"/>
GP <input type="checkbox"/>	<input type="text" value="0.000000"/>	LX <input type="checkbox"/>	<input type="text" value="0.000000"/>	LY <input type="checkbox"/>	<input type="text" value="0.000000"/>
S/L <input type="checkbox"/>	<input type="text" value="0.040000"/>	H/L <input type="checkbox"/>	<input type="text" value="0.030000"/>	trns <input type="checkbox"/>	<input type="text" value="0.000000"/>
shft <input type="checkbox"/>	<input type="text" value="0.000000"/>	stec <input type="checkbox"/>	<input type="text" value="0.000000"/>	ptec <input type="checkbox"/>	<input type="text" value="0.000000"/>
sfec <input type="checkbox"/>	<input type="text" value="0.000000"/>	L11 <input type="checkbox"/>	<input type="text" value="0.000000"/>	L22 <input type="checkbox"/>	<input type="text" value="0.000000"/>
L33 <input type="checkbox"/>	<input type="text" value="0.000000"/>	L12 <input type="checkbox"/>	<input type="text" value="0.000000"/>	L13 <input type="checkbox"/>	<input type="text" value="0.000000"/>
L23 <input type="checkbox"/>	<input type="text" value="0.000000"/>				

type 3

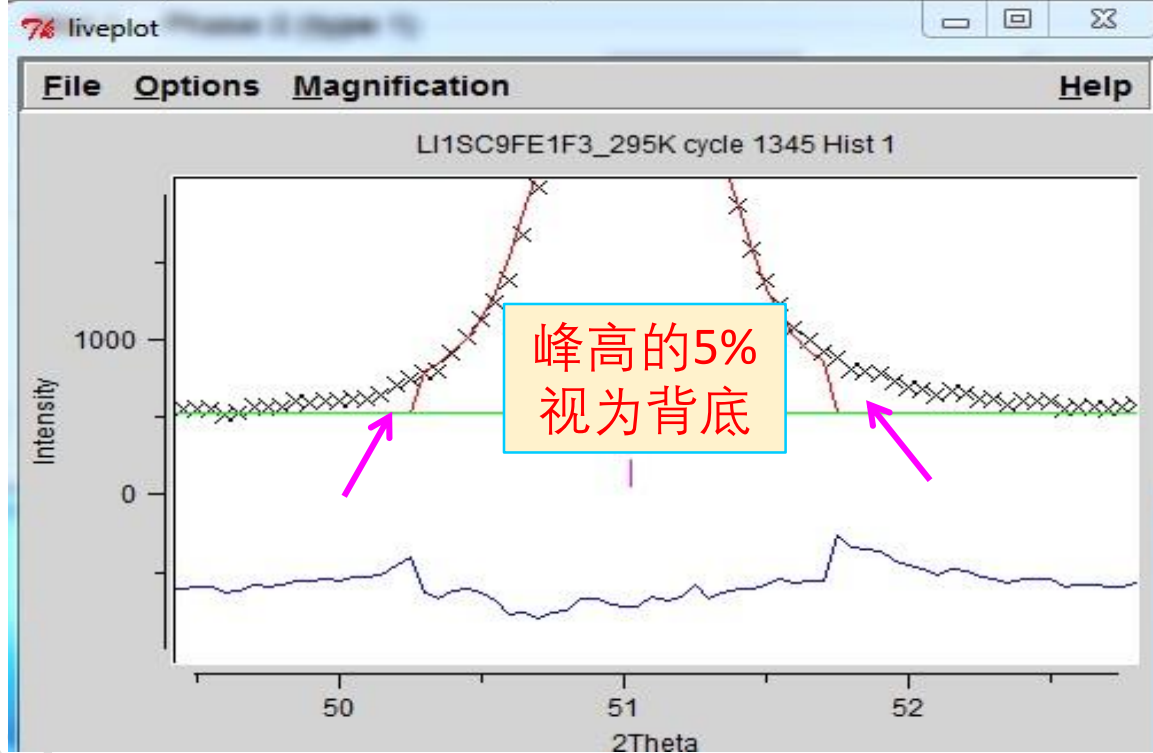
Set peak cutoff

Hist 1 -- Phase 1 (type 4)

Damping Peak cutoff

GU	<input checked="" type="checkbox"/>	<input type="text" value="0.603080E+03"/>	GV	<input checked="" type="checkbox"/>	<input type="text" value="-0.154326E+03"/>	GW	<input checked="" type="checkbox"/>	<input type="text" value="0.836753E+02"/>
GP	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>	LX	<input checked="" type="checkbox"/>	<input type="text" value="0.249161E+02"/>	ptec	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>
trns	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>	shft	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>	sfec	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>
S/L	<input type="checkbox"/>	<input type="text" value="0.500000E-03"/>	H/L	<input type="checkbox"/>	<input type="text" value="0.500000E-03"/>	eta	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>
S400	<input checked="" type="checkbox"/>	<input type="text" value="0.147940E-01"/>	S220	<input checked="" type="checkbox"/>	<input type="text" value="0.509791E+01"/>			

0.05

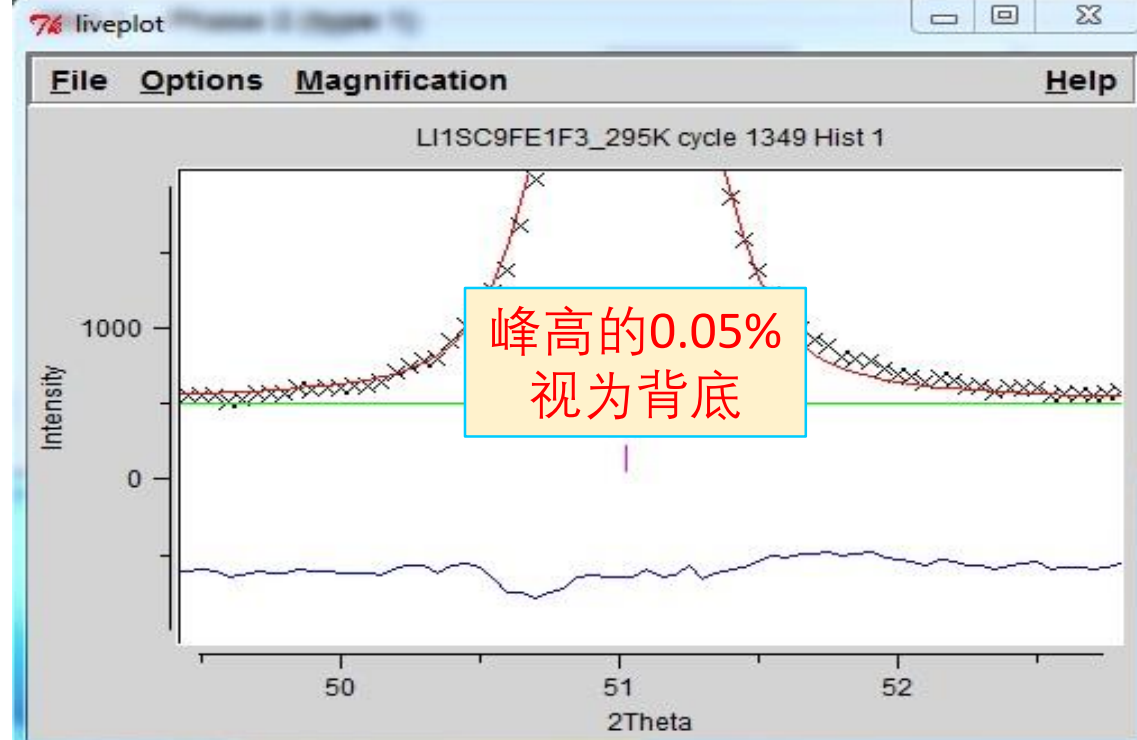


Hist 1 -- Phase 1 (type 4)

Damping Peak cutoff

GU	<input checked="" type="checkbox"/>	<input type="text" value="0.541946E+03"/>	GV	<input checked="" type="checkbox"/>	<input type="text" value="-0.130122E+03"/>	GW	<input checked="" type="checkbox"/>	<input type="text" value="0.957231E+02"/>
GP	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>	LX	<input checked="" type="checkbox"/>	<input type="text" value="0.232900E+02"/>	ptec	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>
trns	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>	shft	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>	sfec	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>
S/L	<input type="checkbox"/>	<input type="text" value="0.500000E-03"/>	H/L	<input type="checkbox"/>	<input type="text" value="0.500000E-03"/>	eta	<input type="checkbox"/>	<input type="text" value="0.000000E+00"/>
S400	<input checked="" type="checkbox"/>	<input type="text" value="0.345154E+00"/>	S220	<input checked="" type="checkbox"/>	<input type="text" value="0.580268E+01"/>			

0.005



File Options Powder Xtal Graphs Results Calc Macro Import/Export Help

expnam expedt genes powpref powplot Istview liveplot

LS Controls Phase Powder Scaling Profile Constraints Restraints Rigid Body MD Pref Orient SH Pre

Select a histogram

h# 76 Edit Absorption/Reflectivity

1

Changing settings for histogram 1

Absorption Coefficient(s)		Absorption Function
1	2	
0.0000000	0.0000000	0

Correction for cylindrical samples [Lobanov & Alte da Veiga]. OK for all data types, but not for Bragg-Brentano flat-plate geometry. Set term 1 to μ^*R/λ (TOF: μ^*R for $\lambda=1$). For CW x-ray/neutron, do not refine!

Continue Cancel Help

Function type 2 (12 terms) Edit Background

Refine background Damping 0

Compton Constants

wave wave 1.5403000 Damping 0

zero Zero -2.03475

Absorption/Reflectivity Correction

Model #0, value: 0.0000000

Refine Abs./Refl. Damping 0 Edit Abs./Refl.

Add New Histogram Set Data Limits & Excluded Regions Set Histogram Use Flags Edit Δ^* and Δ^{θ}

GSAS-EXPGUI: Powder-Edit Edit Abs./Refl-Absorption

样品的吸收校正不足

样品含有大量的元素Tb和Co，室温中子数据在中国散裂中子源高分辨中子通用粉末衍射仪(GPPD)上收集。

File Options Powder Xtal Graphs Results Calc Macro Import/Export Help

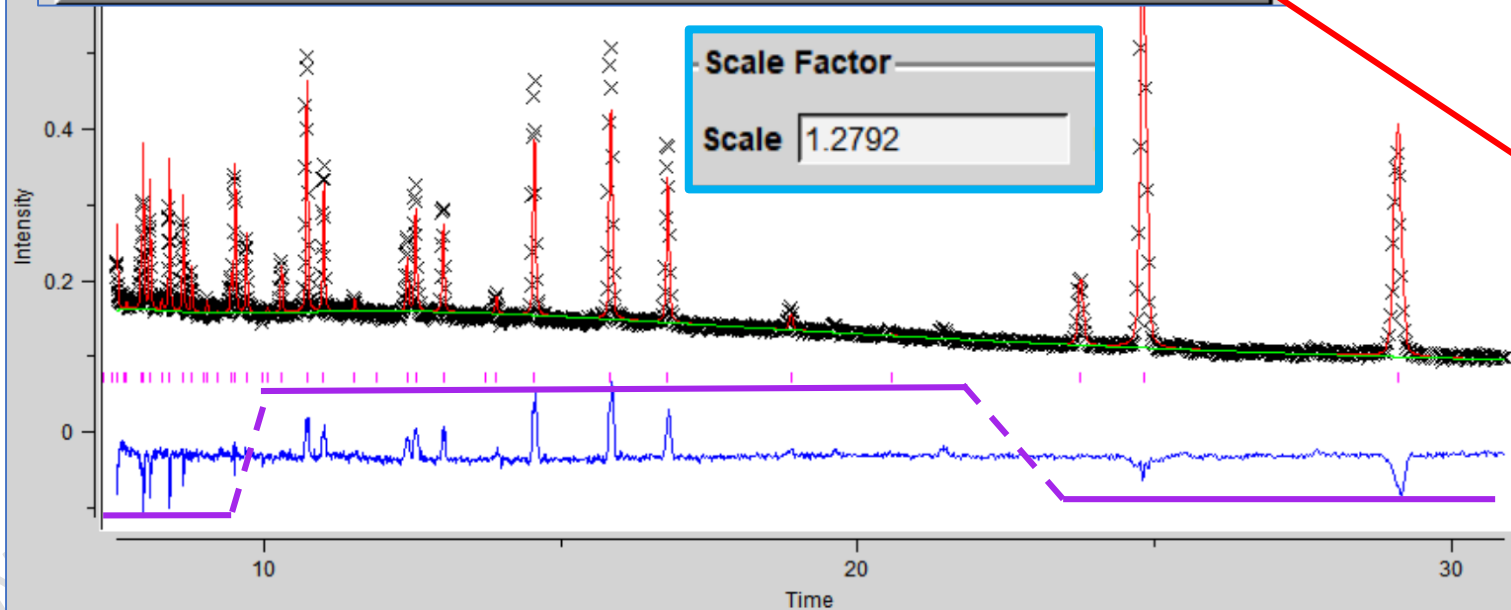
expnam expdet genes powpref powplot lstview liveplot

LS Controls Phase Powder Scaling Profile Constraints Restraints Rigid Body MD Pref

Phase: 1 Replace title: from G:/exp/RCo2highentropyVTN/300K/bank2/ErCo2.cif

Add Phase a 7.187549 b 7.187549 c 7.187549 Edit Refine Cell
α 90.0000 β 90.0000 γ 90.0000 Cell damping 0

* name	type	ref/damp	fractional coordinates			Mult	Occupancy	Uiso
1		0 0 0	0.375000	0.375000	0.375000	8	0.3500	-0.01722
2		0 0 0	0.000000	0.000000	0.000000	16	0.9000	-0.01838
3		0 0 0	0.375000	0.375000	0.375000	8	0.3000	-0.01722
4		0 0 0	0.375000	0.375000	0.375000	8	0.3500	-0.01722
5		0 0 0	0.000000	0.000000	0.000000	16	0.1000	-0.01838



File Options Powder Xtal Graphs Results Calc Macro Import/Export Help

expdet genes powpref powplot lstview liveplot

Phase Powder Scaling Profile Constraints Restraints Rigid Body MD Pref

Background

Function type 4 (5 terms) Edit Background

Refine background Damping 0

Compton Constants

DIFC DIFC 1436.36

DIFA DIFA 3.57 Damping 0

Zero Zero 14.29

Abs./Reflectivity Correction

#0 value: 0.000000

Abs./Refl. Damping 0 Edit Abs./Refl.

New Set Data Limits & Set Histogram Edit
Program Excluded Regions Use Flags Δf and Δf'

*中科院物理所田正营提供数据

GSAS-EXPGUI: Powder-Edit Edit Abs./Refl-Absorption

样品的吸收校正恰当

* name	type	ref/damp	fractional coordinates			Mult	Occupancy	Uiso
		0 0 0	0.375000	0.375000	0.375000	8	0.3500	0.00332
		0 0 0	0.000000	0.000000	0.000000	16	0.9000	0.00321
		0 0 0	0.375000	0.375000	0.375000	8	0.3000	0.00332
		0 0 0	0.375000	0.375000	0.375000	8	0.3500	0.00332
		0 0 0	0.000000	0.000000	0.000000	16	0.1000	0.00321

CW-N

Absorption Coefficient(s) Absorption Function

1	2	Function
0.00000	0.0000000	0

Correction for cylindrical samples [Lobanov & Alte da Veiga]. OK for all data types, but not for Bragg-Brentano flat-plate geometry. Set term 1 to μ^*R/λ (TOF: μ^*R for $\lambda=1$). For CW x-ray/neutron, do not refine!

Absorption Coefficient(s) Absorption Function

1	2	Function
0.00000	0.0000000	1

Wavelength-dependent correction for container penetration. Use with TOF & Energy Disp x-ray only.

TOF-N

Absorption Coefficient(s) Absorption Function

1	2	Function
0.00000	0.0000000	2

Surface roughness correction [Pitschke, Hermann & Mutton]. Use with flat-plate reflection geometry (usually Bragg-Brentano) only.

Absorption Coefficient(s) Absorption Function

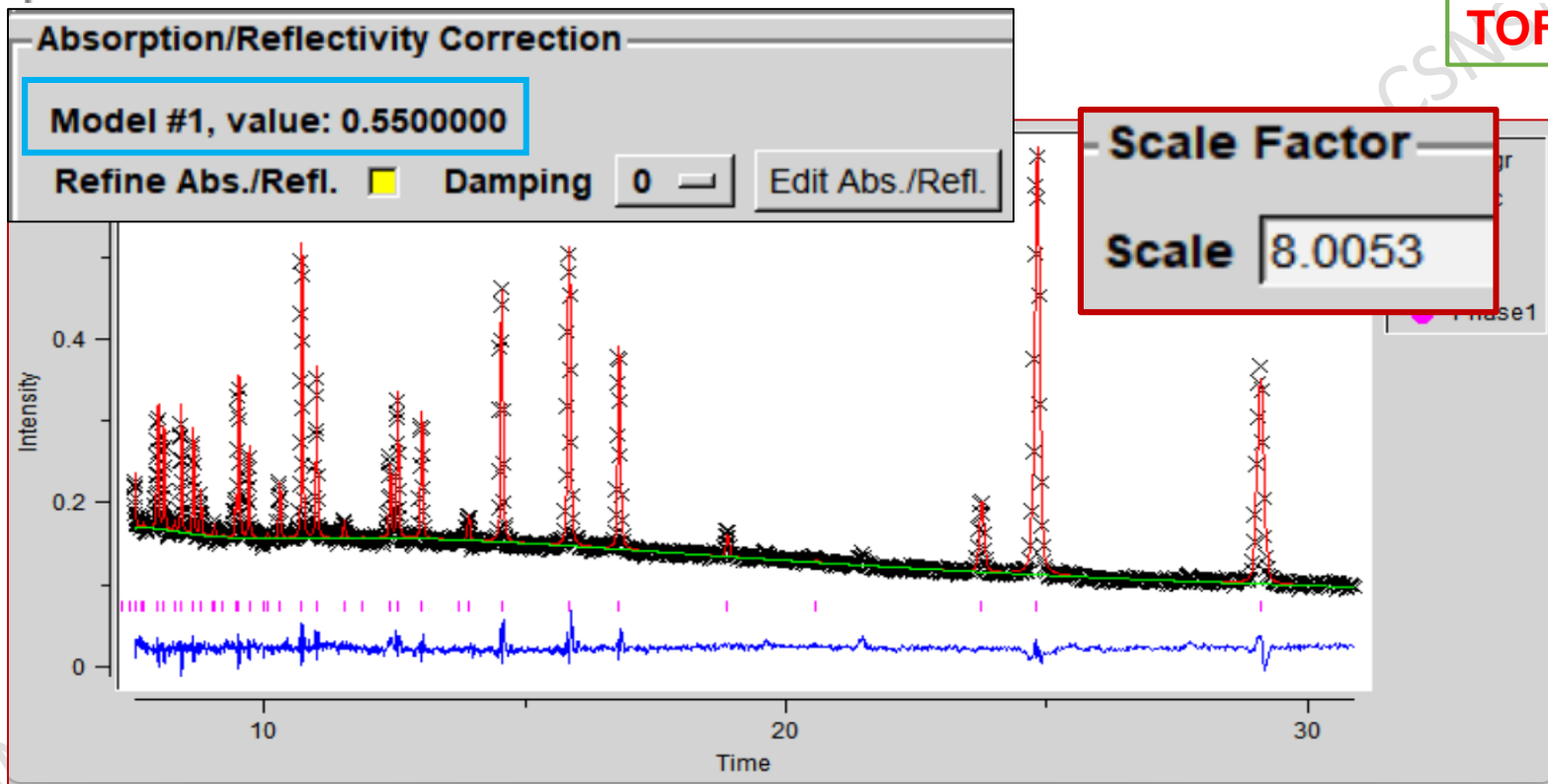
1	2	Function
0.00000	0.0000000	3

Surface roughness correction, [Suortti]. Use with flat-plate reflection geometry (usually Bragg-Brentano) only.

Absorption Coefficient(s) Absorption Function

1	2	Function
0.00000	0.0000000	4

Flat plate samples in transmission mode. OK for all data types, but not Bragg-Brentano geometry. Term 2 is angle w/r to beam (usually 0). For CW, do not refine.



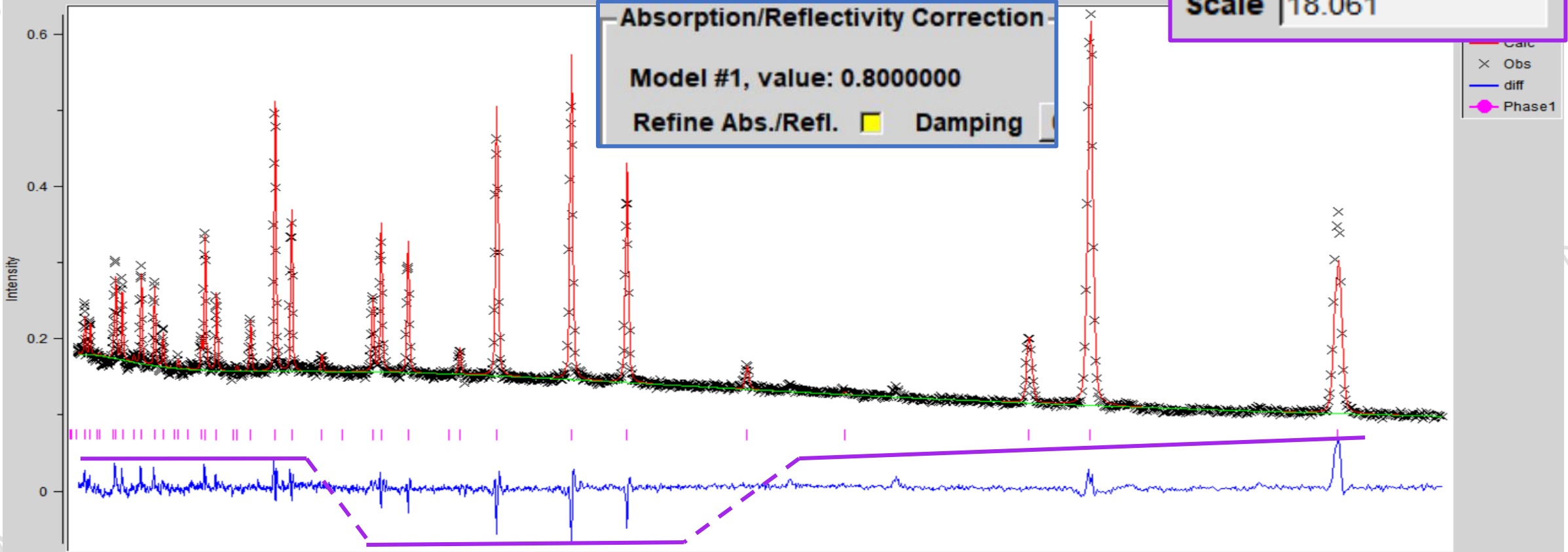
GSAS-EXPGUI: Edit Background-Absorption

样品的吸收校正过头

* name	type	ref/damp	fractional coordinates			Mult	Occupancy	Uiso
1		0 0 0	0.375000	0.375000	0.375000	8	0.3500	0.01413
2		0 0 0	0.000000	0.000000	0.000000	16	0.9000	0.01371
3		0 0 0	0.375000	0.375000	0.375000	8	0.3000	0.01413
4		0 0 0	0.375000	0.375000	0.375000	8	0.3500	0.01413
5		0 0 0	0.000000	0.000000	0.000000	16	0.1000	0.01371

Scale Factor
Scale 18.061

Absorption/Reflectivity Correction
Model #1, value: 0.800000
Refine Abs./Ref. Damping



GSAS-EXPGUI: Constraints

File Options Powder Xtal Graphs Results Calc Macro Import/Export Help																			
expnam		expedt		genes		powpref		powplot		lstview		liveplot							
LS Controls		Phase		Powder		Scaling		Profile		Constraints		Restraints		Rigid Body		MD Pref Orient		SH Pref Orient	
#		Phase	Atom(s)	Variable	Multiplier	Atom(s)	Variable	Multiplier	Delete										
1	edit	1	1,5	X	x 1.0000				<input type="checkbox"/>										
2	edit	1	1,5	Y	x 1.0000				<input type="checkbox"/>										
3	edit	1	1,5	UI50	x 1.0000				<input type="checkbox"/>										
4	edit	1	5	FRAC	x -1.0000	1	FRAC	x 1.0000	<input type="checkbox"/>										

* name	type	ref/damp	fractional coordinates			Mult	Occupancy	Uiso
1 La	LA	X0 U0 F0	-0.004805	0.022661	0.250000	4	0.7740	0.00403
2 Mn	MN+3	X0 U0 0	0.500000	0.000000	0.000000	4	1.0000	0.00308
3 O1	O	X0 U0 0	0.064747	0.491805	0.250000	4	1.0000	0.00627
4 O2	O	X0 U0 0	0.723533	0.275049	0.033679	8	1.0000	0.00545
5 Ca	CA	X0 U0 F0	-0.004805	0.022661	0.250000	4	0.2260	0.00403

Carbon Content and Temperature Dependence of the T_c and Structure Parameters in the Non-oxide Perovskite Superconductor MgC_xNi₃

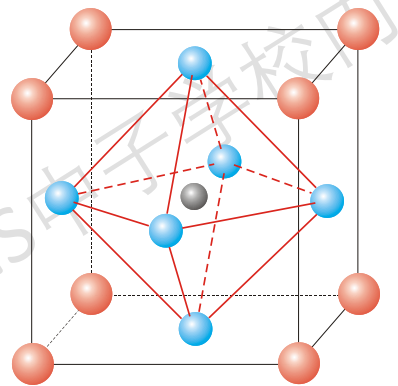
He T., Huang Q. Z., *et al.*, *Nature*. **411**, 2001, 54-56.

CSNS
Qingzhen Huang

China Spallation Neutron Source

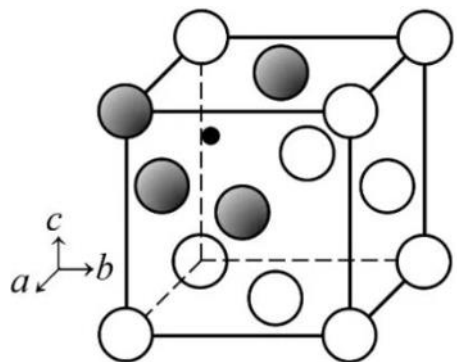
CSNS中子学校内部资料, 请勿外传

T_c vs C content for MgC_xNi₃

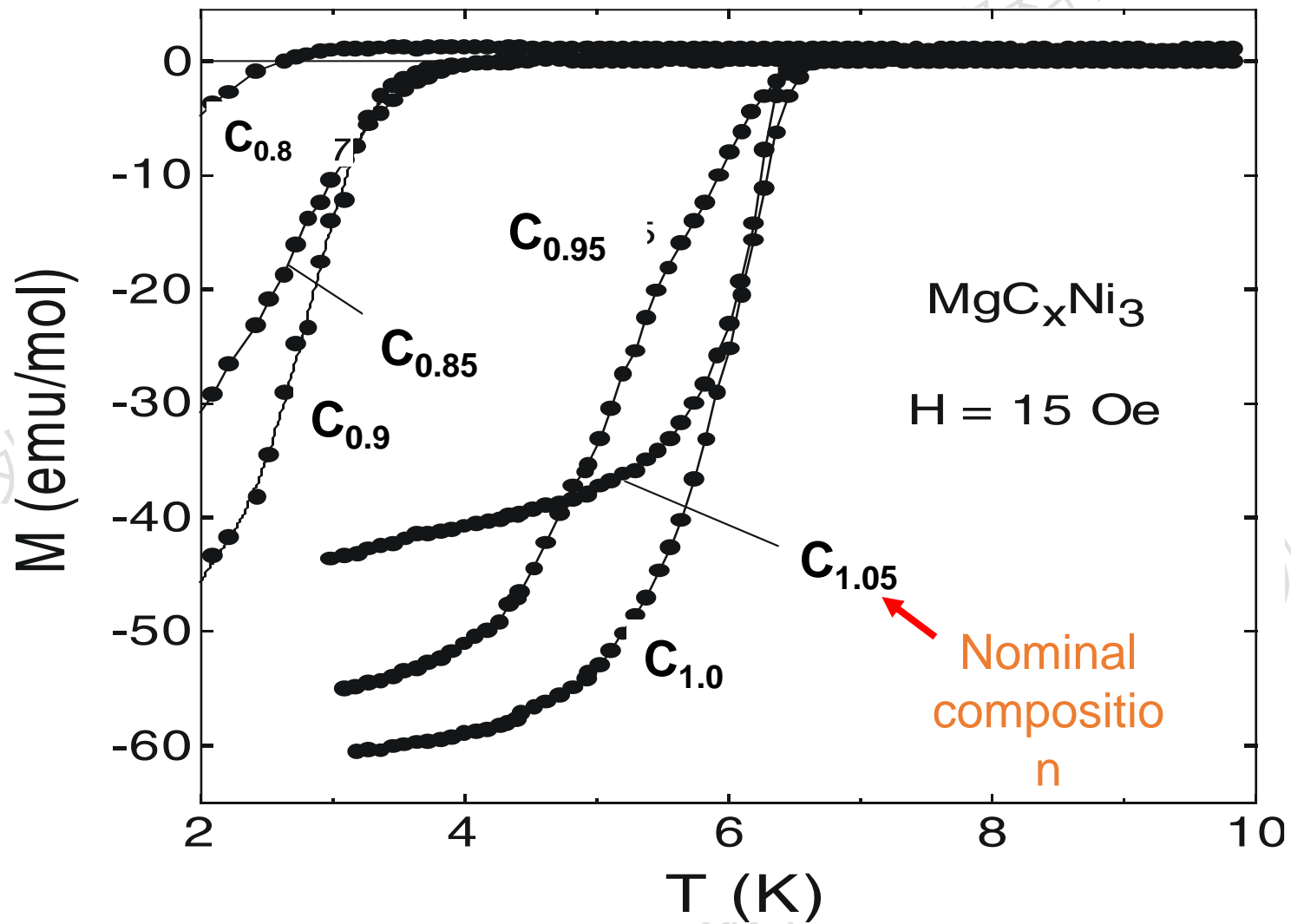


Structure of
Perovskite
ABX₃

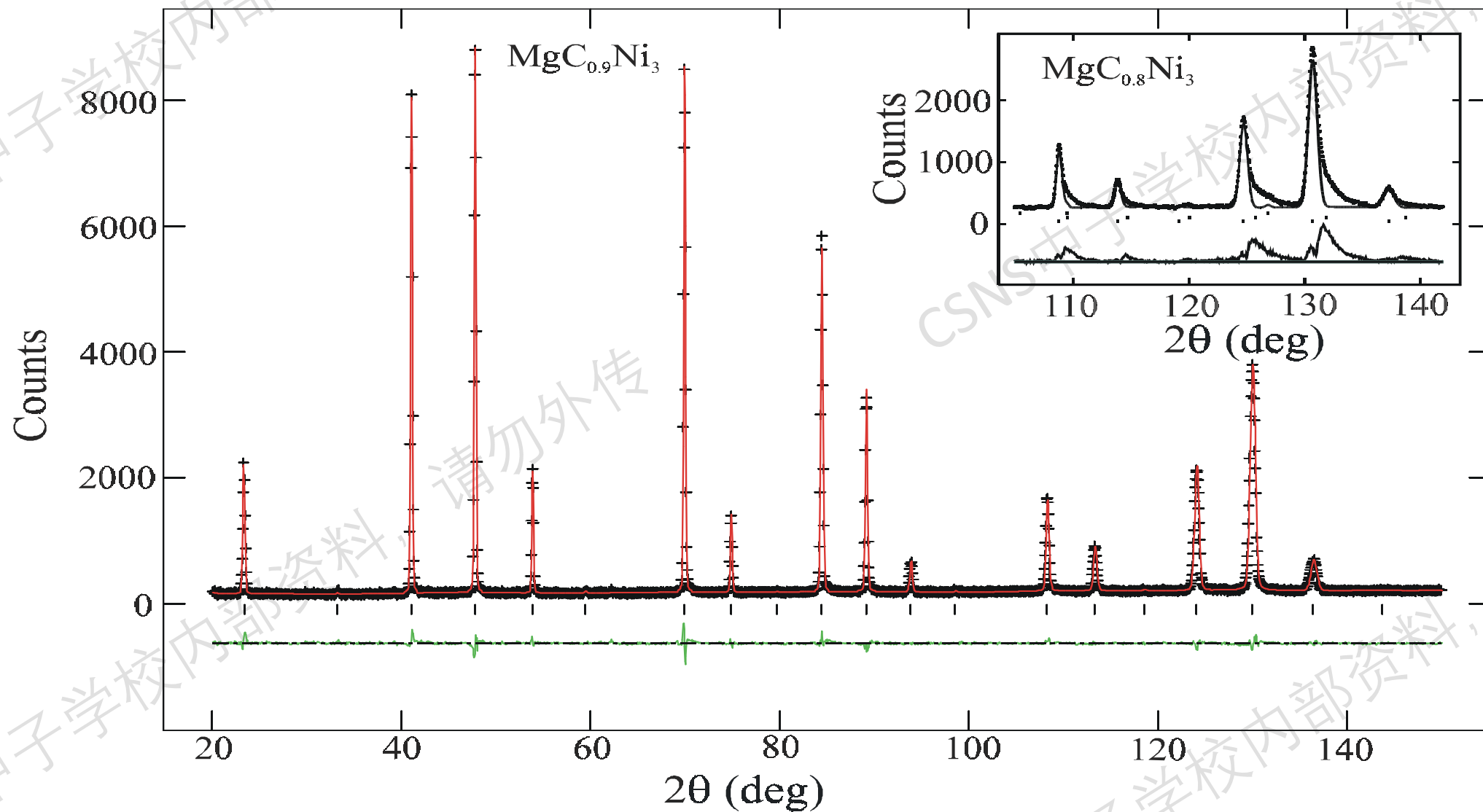
● Mg ● C ● Ni



面心立方的四面体间隙及空间排布



Neutron diffraction pattern of MgC_xNi_3



Constant neutron scattering length gives stronger intensities in higher angle range.

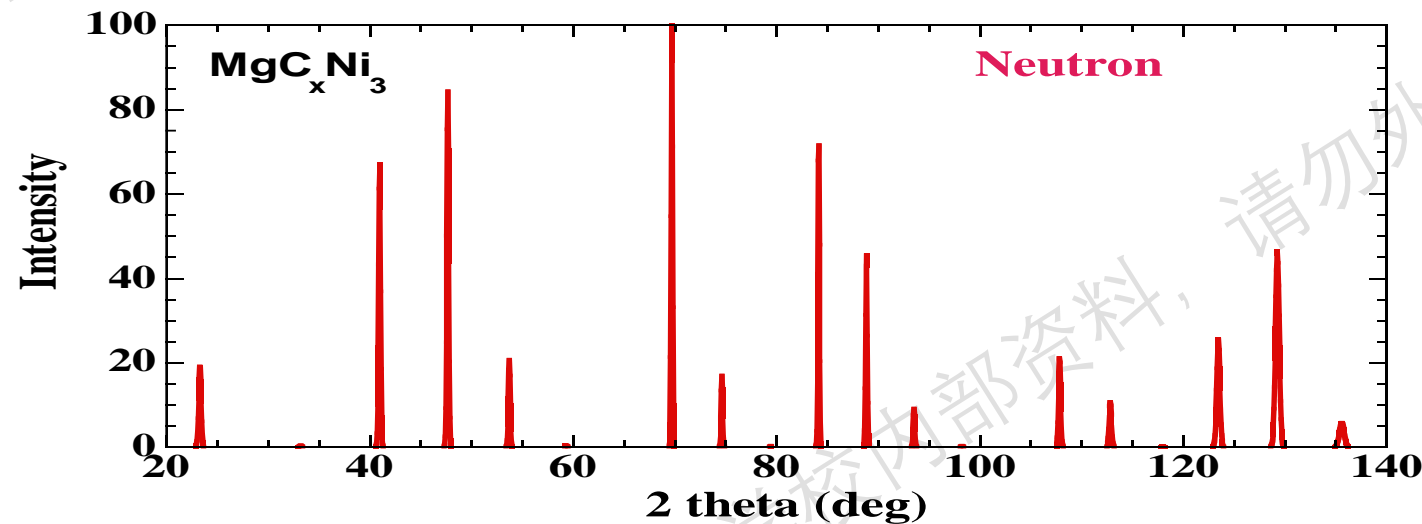
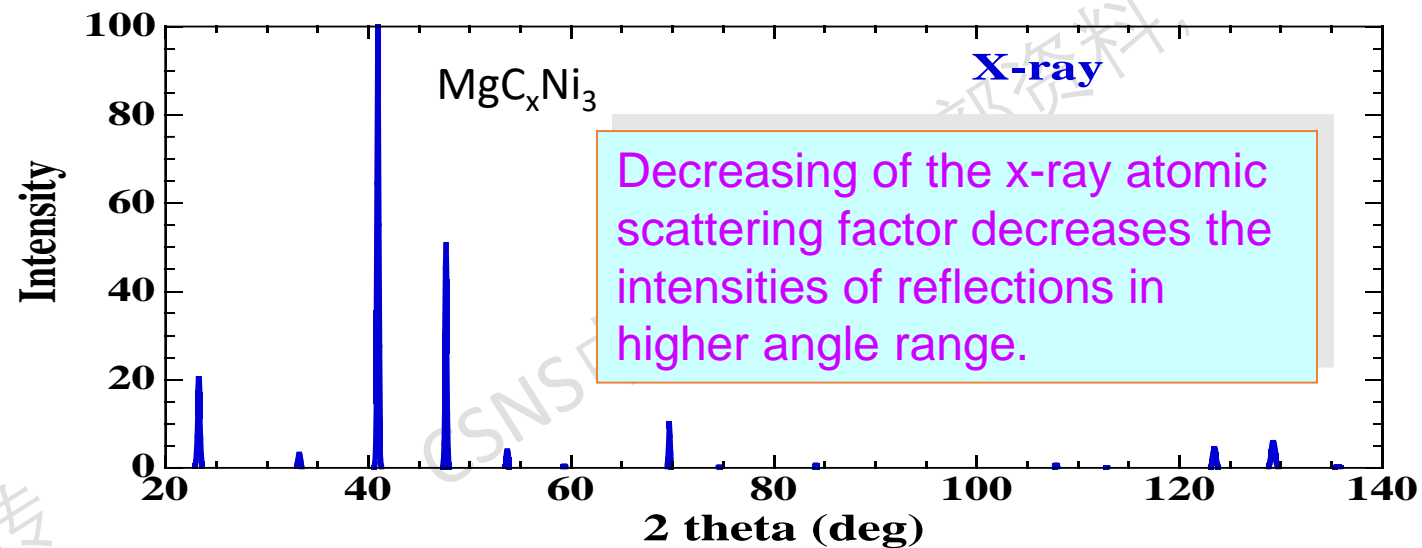
	C	Mg	Ni
X-ray #	6	12	28
Neutron <i>b</i>	0.665	0.583	1.03

Ne(Mg):Ne(C):Ne(Ni) 0.429:0.214:1

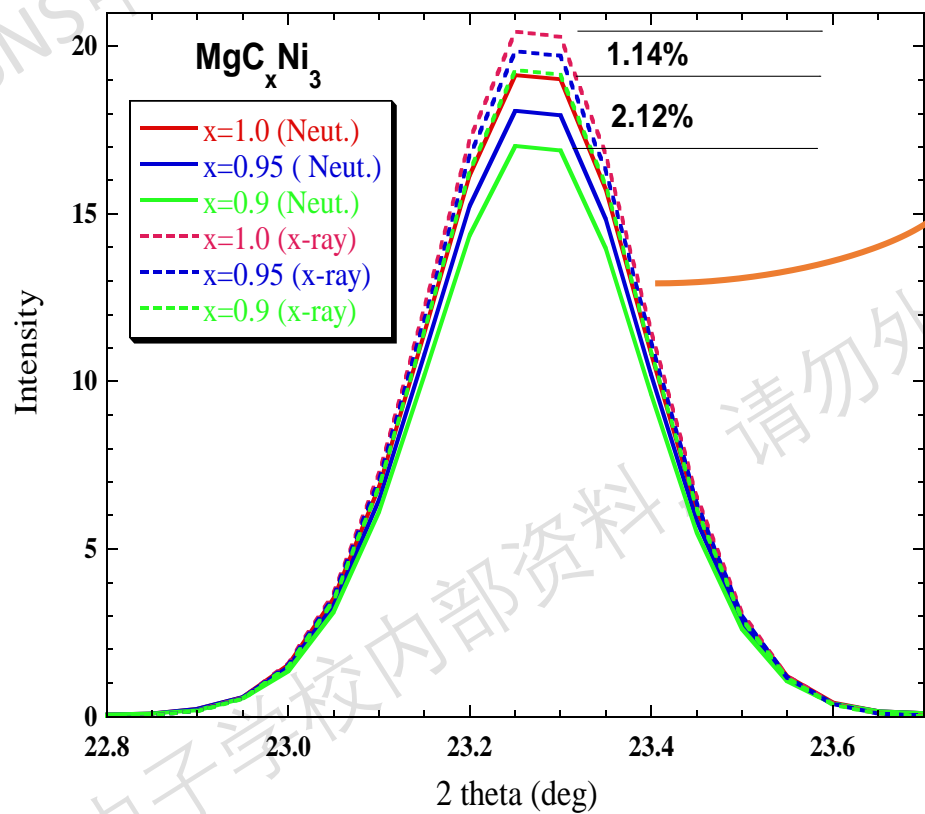
b(Mg):*b*(C):*b*(Ni) 0.522:0.646:1

X-ray: $|F_{hkl}|^2 = |\sum f_j \exp(2\pi i (hx + ky + lz))|^2 e^{-2W}$

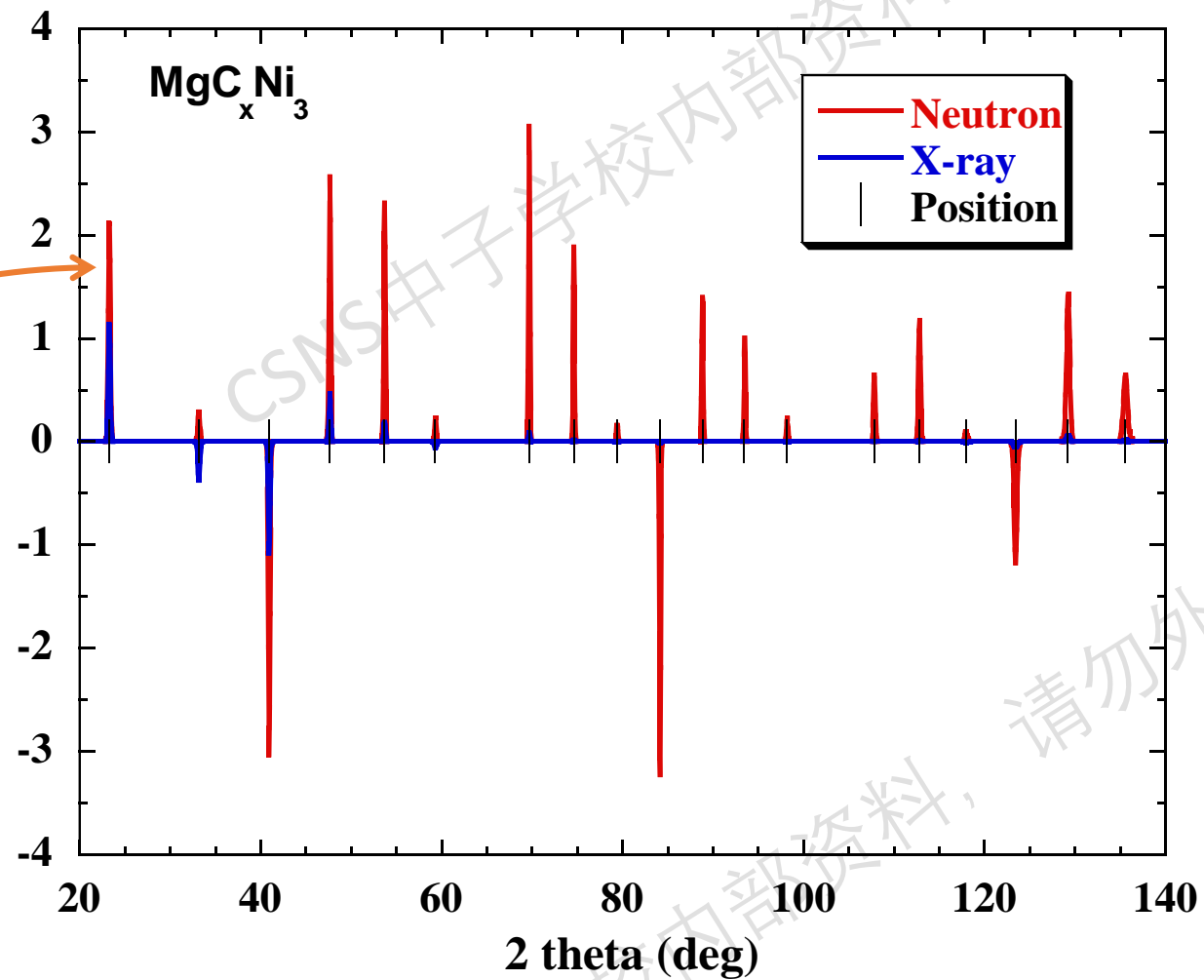
Neutron: $|F_{hkl}|^2 = \sum b_j \exp(2\pi i (hx + ky + lz))|^2 e^{-2W}$



Differences in Intensities



$I(C_{1.0}) - I(C_{0.9})$ (%)



MgC_{0.978}Ni₃, T_c=7.30 K

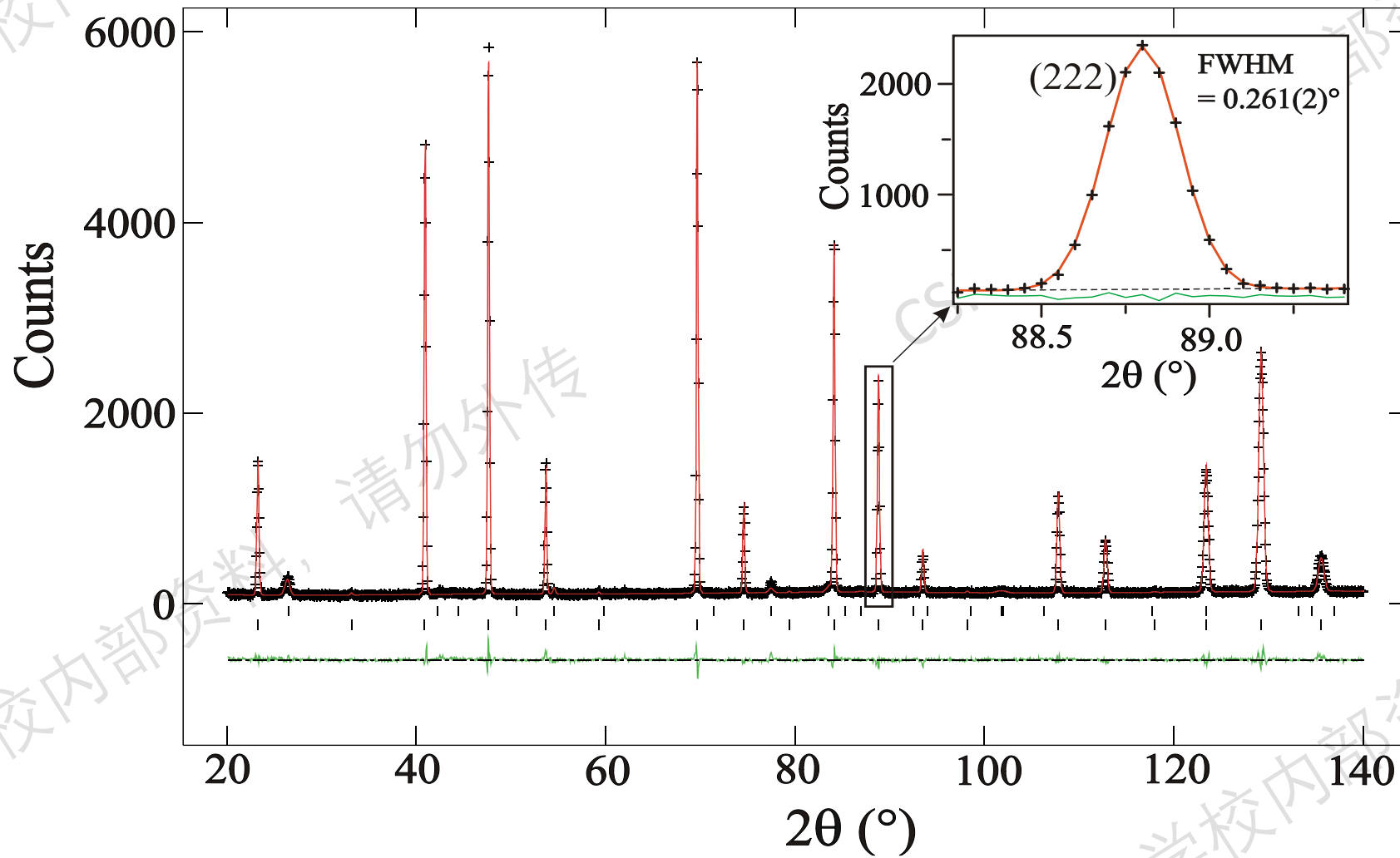
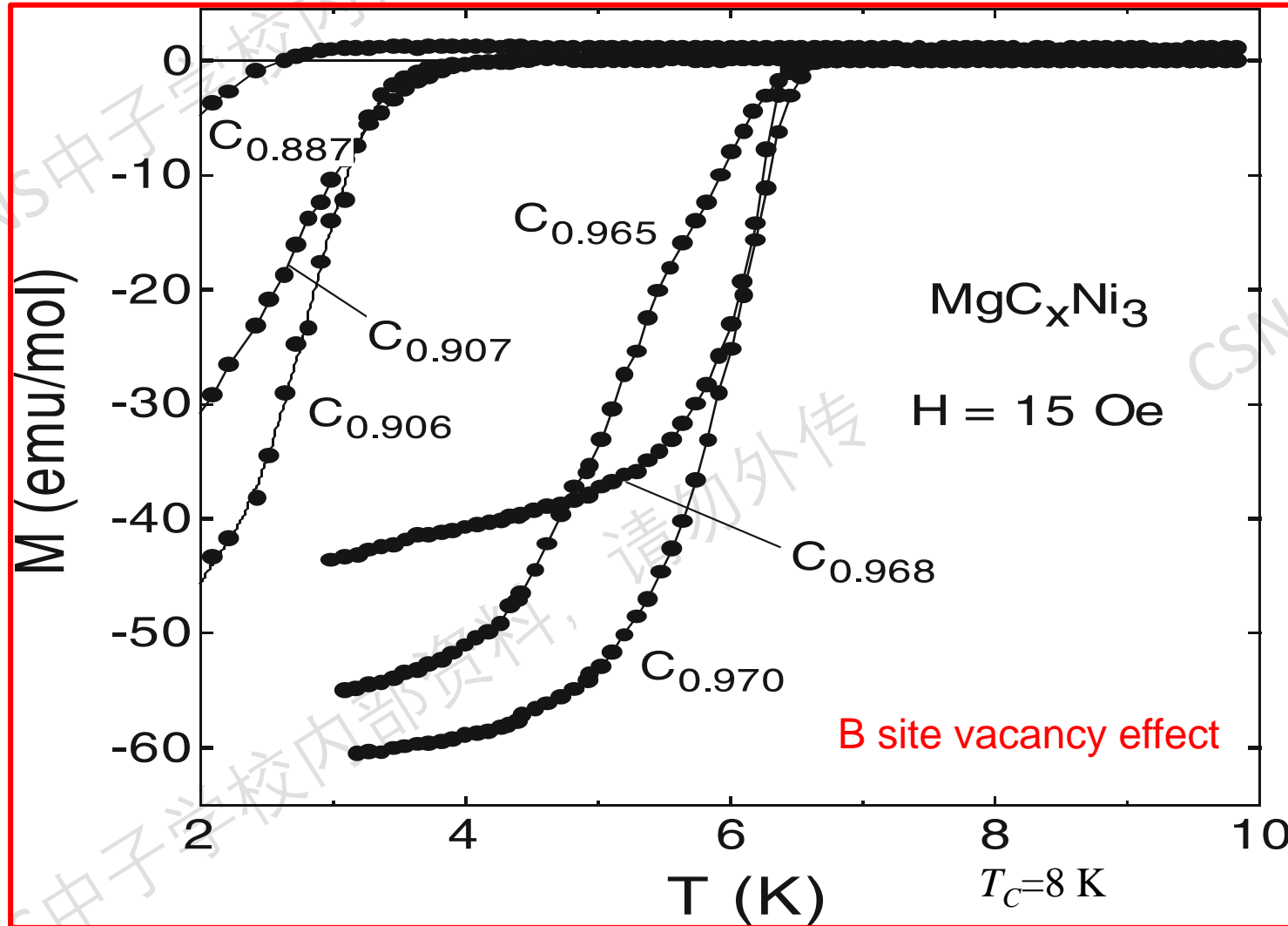


Table. Structure Information of the MgC_xNi₃ Sample

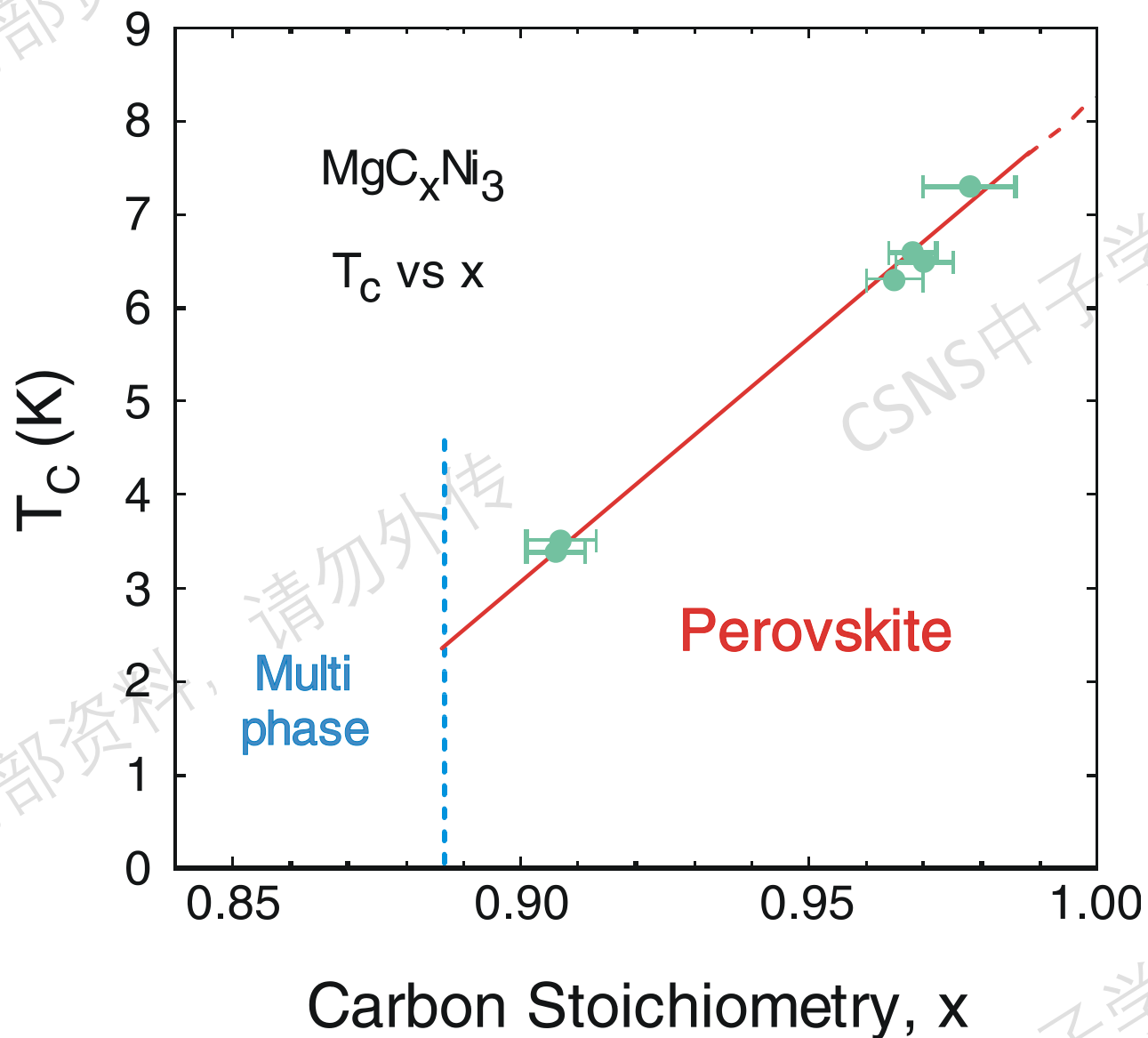
X _{nominal}	T _c (K)	X _{neutron}	Phase Fraction (%)			a (Å)		χ ²	R _p (%)	R _{wp} (%)
			I	II	III	I	II			
1.25	7.30	0.978(8)	98.0	0.0	2.0	3.81221(5)	-	1.191	4.49	5.95
1.05	6.59	0.968(4)	97.3	0.0	2.7	3.81207(2)	-	1.166	3.32	3.84
1.0	6.49	0.970(5)	96.8	0.0	3.2	3.81060(2)	-	1.174	3.79	4.76
0.95	6.30	0.965(5)	98.5	0.0	1.5	3.80990(2)	-	1.413	4.15	5.36
0.9	3.38	0.906(5)	100	0	0.0	3.80014(2)	-	1.612	4.28	5.53
0.85	3.51	0.907(6)	64.0	34.6	1.4	3.80147(5)	3.78844(23)	1.311	4.13	5.31
0.8	<1.8	0.887(7)	58.5	38.0	3.5	3.79515(5)	3.77802(25)	1.690	3.85	4.89

T_c vs C content

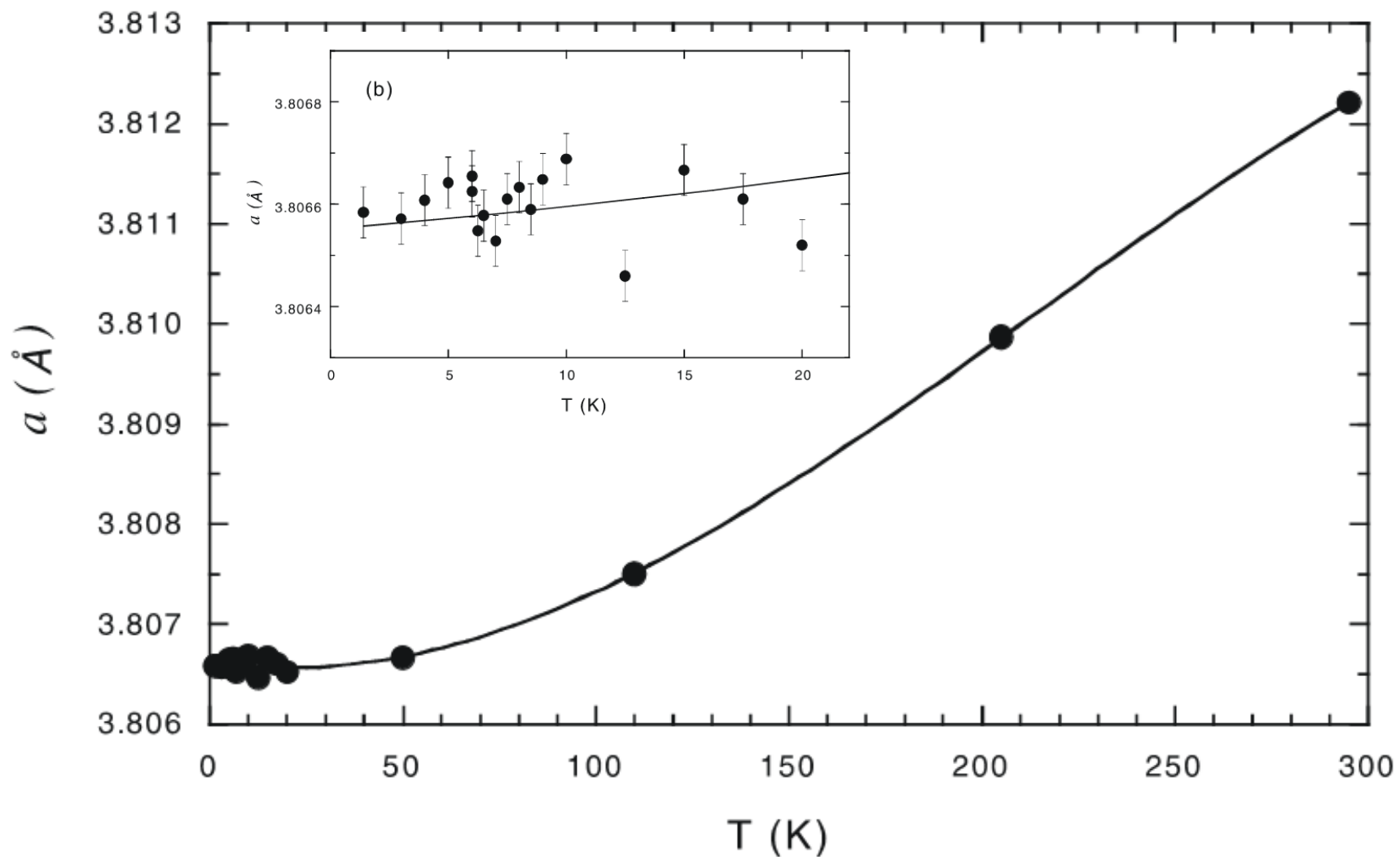


X_{nominal}	T_c (K)	X_{neutron}
1.25	7.30	0.978(8)
1.05	6.59	0.968(4)
1.0	6.49	0.970(5)
0.95	6.30	0.965(5)
0.9	3.38	0.906(5)
0.85	3.51	0.907(6)
0.8	<1.8	0.887(7)

Plot of T_c as a Function of Carbon Content x in MgC_xNi₃

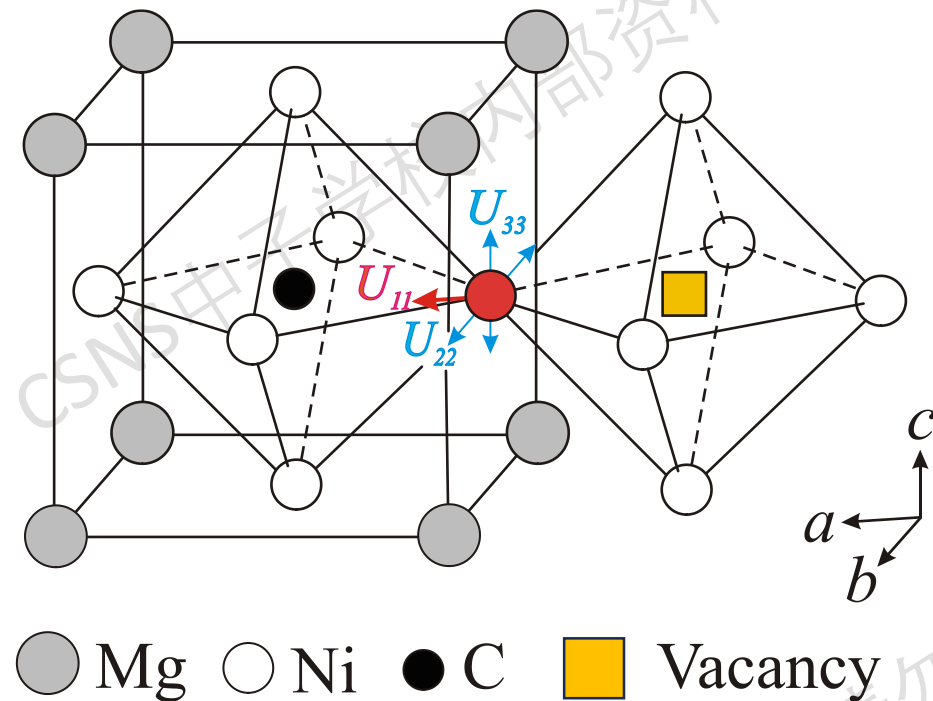
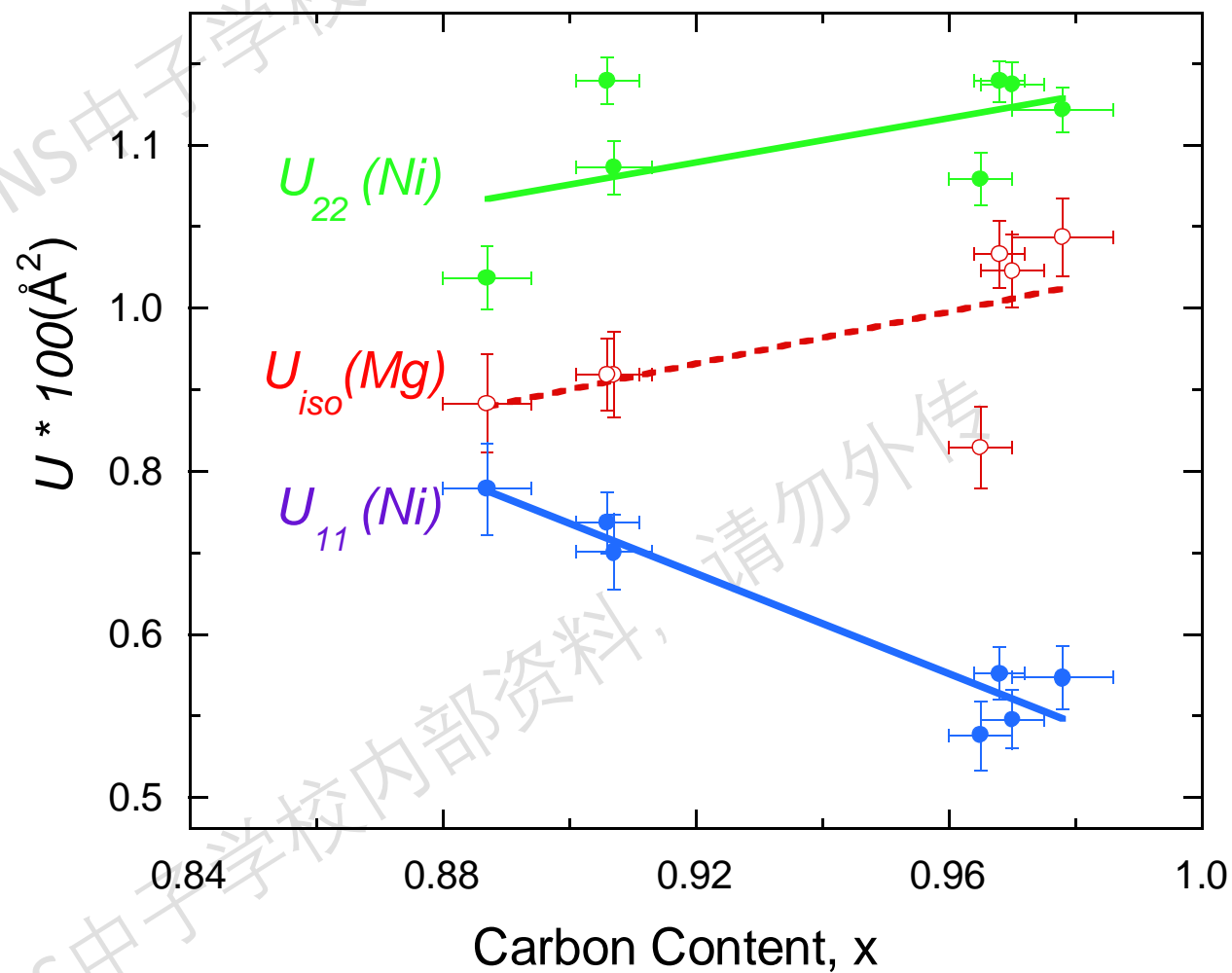


No Detectable Structural Transition Observed Between 2 and 300 K for MgC_xNi₃



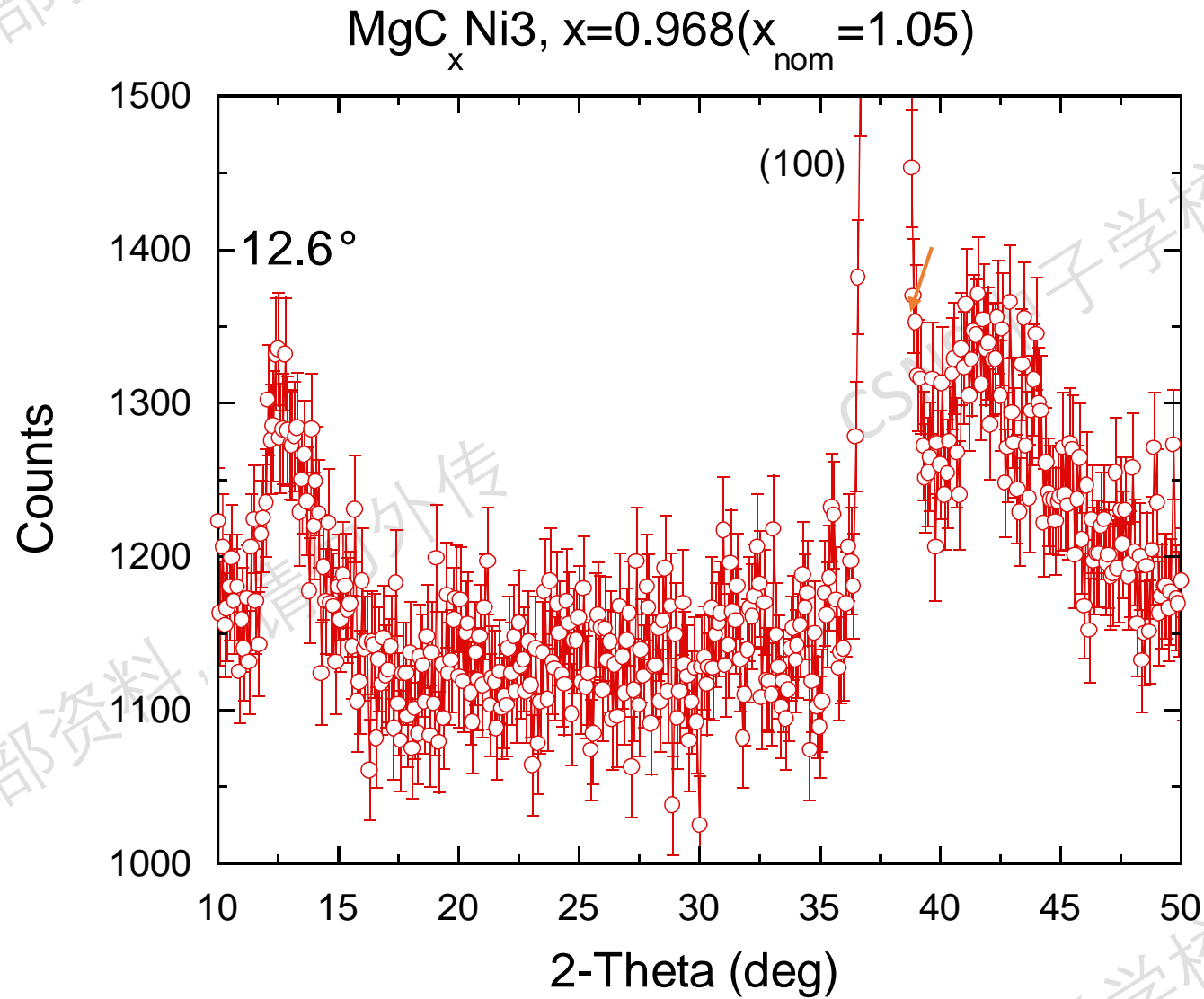
MgC_xNi₃中C的占有率引起的温度因子变化

Variation of the Temperature Factors as a Function of Carbon Content x for MgC_xNi₃



在室温，右图中所示的温度因子 $U_{11}(\text{Ni})$ 随着C的占有率减少而增大，说明当产生C缺位时，近邻Ni向邻近的C原子位移，如上图。

Neutron intensity showing the short-range graphite peak



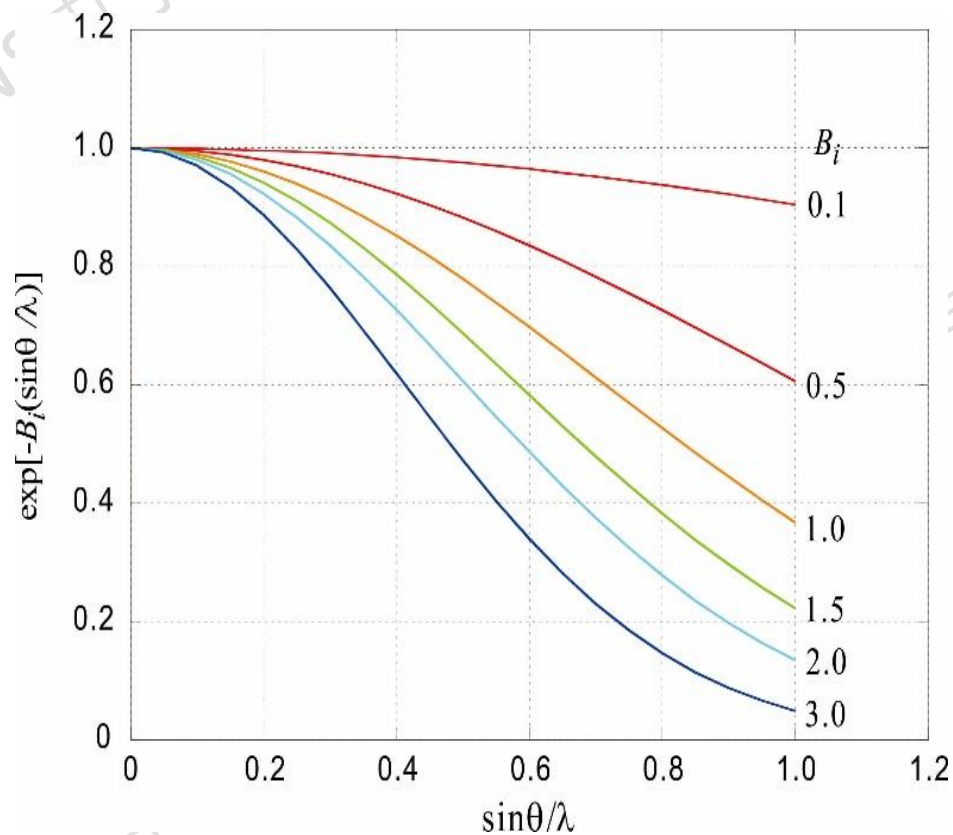
Summary

- * Perovskite phase stability range $0.88 < x < 1.0$ for MgC_xNi₃.
- * T_c monotonically decreases with decreasing x .
- * Carbon vacancy affects the position of the Ni atoms.
- * **No detectable structural transition and magnetic order.**

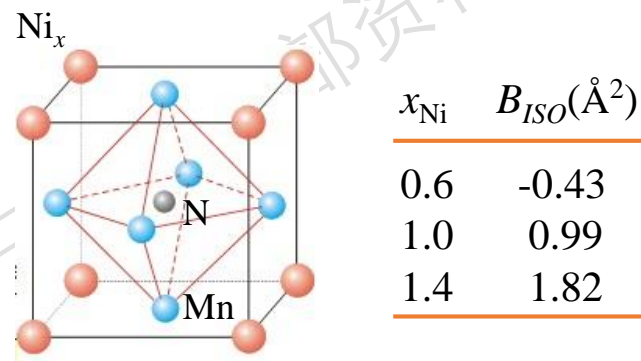
结构精化中反常的温度因子与结构的相关性

Correlation between abnormal temperature factors and structure in structure refinement

$$|F_{hkl}|^2 = \left| \sum b_j \exp(2\pi i (hx + ky + lz)) \right|^2 e^{-2W}$$



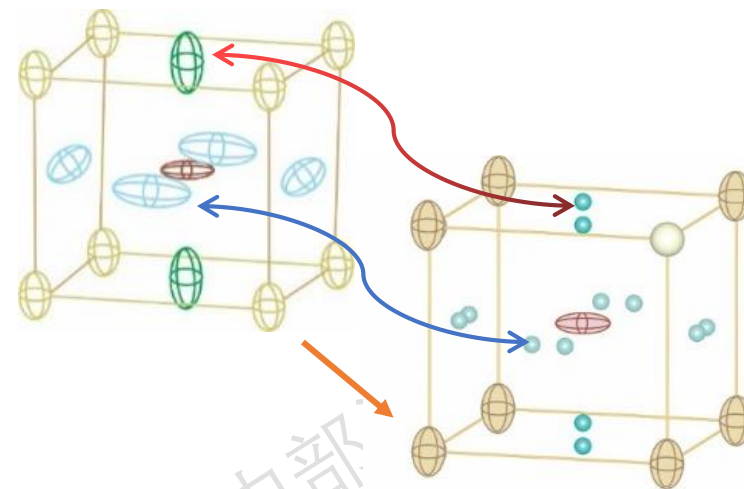
(a) B_i 与 $f_i(b, q_i)$ 成正比：因此当结构中某个结晶学位置上有空位，或被散射因子小的元素占有，即是该位置的散射本领偏小，这时 B_i 减小，可以考虑占有率或占位问题；



(b) 与原子位置的位移相关：原子位置偏离，或静态位移，或劈裂；

(c) 与原子动态位移有关：配位多面体，低键联，轻元素，各向异性环境等等；

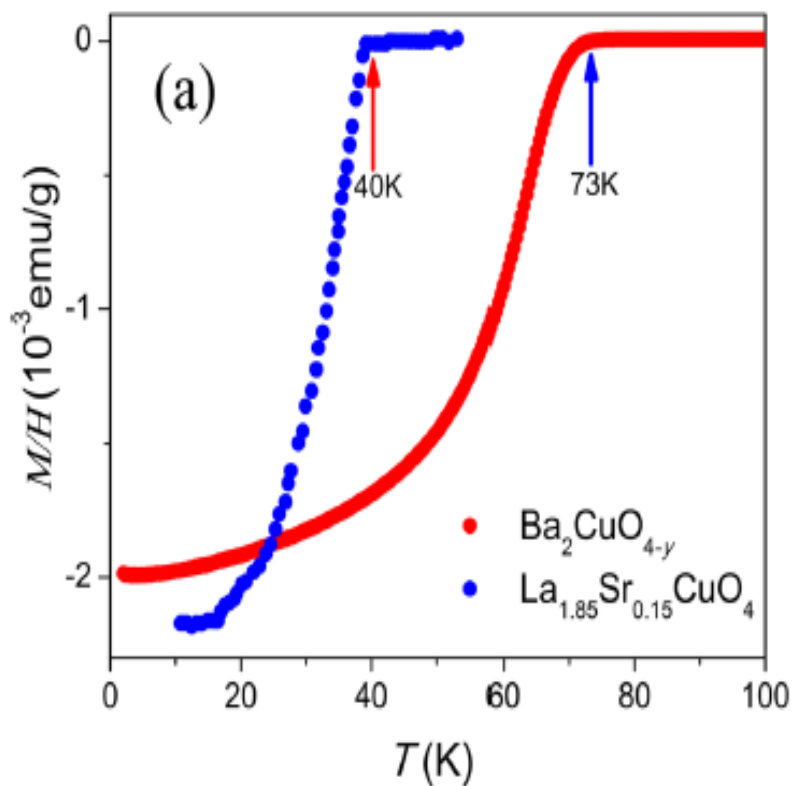
(d) 与物性有关：声子振动等。



大各向异性温度因子与原子位置的静态位移和占有率

Strong Anisotropic Temperature Factor Associated with Atomic Shift and Occupancy

探询Ba₂CuO_{4-x} 比La_{1.85}Sr_{0.15}CuO₄ 的超导温度高出33K的原因，了解其晶体结构异同能够提供有力的信息。



Temperature dependence of field-cooling (FC) dc magnetic susceptibility of Ba214 and optimum doped La_{1.85}Sr_{0.15}CuO₄.

Table 1. Refined structure parameters using three different models for Ba₂CuO_{4-x} at 6 K. Space group *I4/mmm* (#139), atomic position: Ba: 4*e* (0 0 *z*); Cu: 2*a* (0 0 0); O(1): 4*e* (0 0 *z*); O(2): 4*c* (0, ½, 0); O'(2): 8*j* (*x*, ½, 0); O(11): 4*e* (0 0 *z*); and O(12): 4*e* (0 0 *z*).

Model I. Isotropic temperature and full occupancy.

$$a = b = 3.9931(2) \text{ (\AA)}, c = 12.921(1) \text{ (\AA)}$$

Atom	<i>z</i>	<i>n</i>	100 × <i>U</i> _{iso} (Å ²)
Ba	0.3570(5)	1	-1.1(2)
Cu	0	1	-0.5(2)
O(1)	0.1453(4)	1	0.9(2)
O(2)	0	1	9.9(4)

$$R_{WP} = 5.03\%, R_p = 3.97\%, \chi^2 = 2.200$$

O(2)的温度因子大暗示其占有率或/和偏离现有的位置

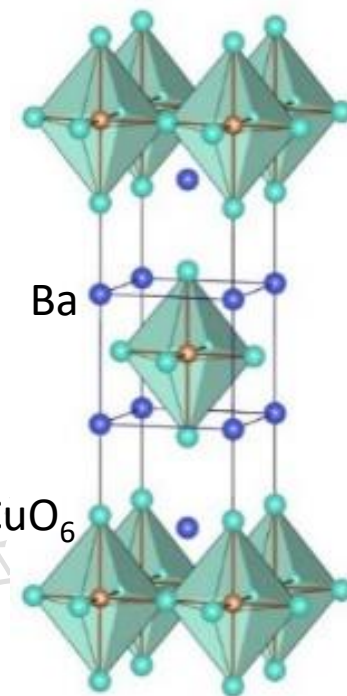


Table. Refined structure parameters using three different models for $\text{Ba}_2\text{CuO}_{4-x}$ at 6 K. Space group $I4/mmm$ (#139), atomic position: Ba: $4e$ (0 0 z); Cu: $2a$ (0 0 0); O(1): $4e$ (0 0 z); O(2): $4c$ (0, $\frac{1}{2}$, 0); O'(2): $8j$ (x , $\frac{1}{2}$, 0); O(11): $4e$ (0 0 z); and O(12): $4e$ (0 0 z).

Model II. Isotropic temperature.

$a = b = 3.9931(2)$ (Å), $c = 12.921(1)$ (Å)

Atom	z	n	$100 \times U_{iso}$ (Å ²)
Ba	0.3580(4)	1	0.3(1)
Cu		1	0.7(2)
O(1)	0.1436(4)	1	2.3(2)
O(2)			0.58(2) 3.6(4)

$R_{WP} = 4.37\%$, $R_p = 3.47\%$, $\chi^2 = 1.665$

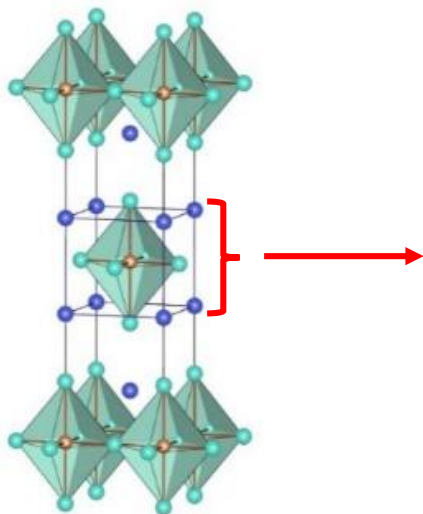
Model III. Anisotropic temperature.

$a = b = 3.9932(2)$ (Å), $c = 12.919(1)$ (Å)

Atom	z	n	U_{11} (Å ²)	U_{22} (Å ²)	U_{33} (Å ²)
Ba	0.3576(4)	1	0.2(2)	$=U_{11}$	0.9(4)
Cu	0	1	1.8(3)	$=U_{11}$	0
O(1)	0.1446(4)	1	1.6(2)	$=U_{11}$	4.7(5)
O(2)	0		0.58(2)	13(1)	2.2(7)

$R_{WP} = 4.19\%$, $R_p = 3.29\%$, $\chi^2 = 1.532$

Model II 的结构中O(2)有42%的空位，但O(1)和O(2)的各向同性温度因子偏大，这可能是由于空位引起原子位置的静态位移。



Model III 各个原子位置各向异性温度因子的精修结果， R 因子有了明显减小。其中BaO层的Ba原子和O(1)的 U_{33} 明显偏大，而CuO₂层中的Cu原子和O(2)的 U_{11} 也明显偏大。

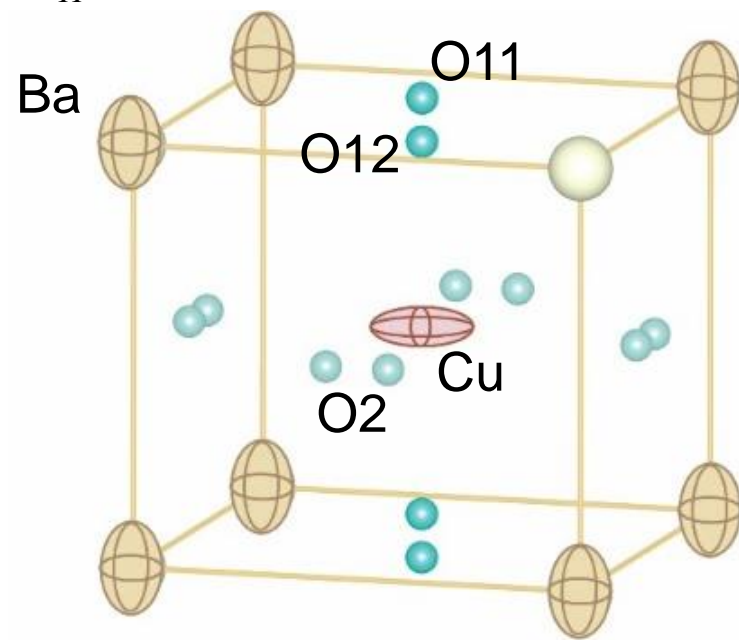
Table. Refined structure parameters using three different models for Ba₂CuO_{4-x} at 6 K. Space group *I4/mmm* (#139), atomic position: Ba: 4*e* (0 0 *z*); Cu: 2*a* (0 0 0); O(1): 4*e* (0 0 *z*); O(2): 4*c* (0, ½, 0); O'(2): 8*j* (*x*, ½, 0); O(11): 4*e* (0 0 *z*); and O(12): 4*e* (0 0 *z*).

Model IV. Oxygen atoms splitting. $R_{WP} = 4.16\%$, $R_p = 3.26\%$, $\chi^2 = 1.511$

$a = b = 3.9932(2)$ (Å), $c = 12.919(1)$ (Å).

Atom	<i>x</i>	<i>z</i>	<i>n</i>	U_{iso} (Å ²)	U_{11} (Å ²)	U_{22} (Å ²)	U_{33} (Å ²) × 100
Ba	0	0.3577(4)	1		0.0	= U_{11}	0.9(3)
Cu	0	0	1		1.8(3)	= U_{11}	0
O(11)	0	0.132(1)	0.5	1.5(2)			
O(12)	0	0.157(1)	0.5	1.5(2)			
O'(2)	0.077(4)	0	0.297(8)	0.8(5)			

Model IV 的结构中O(1)以及Cu和O(2)各沿着向 U_{33} 和及 U_{11} 的方向劈裂成两个位置，如上图所示。



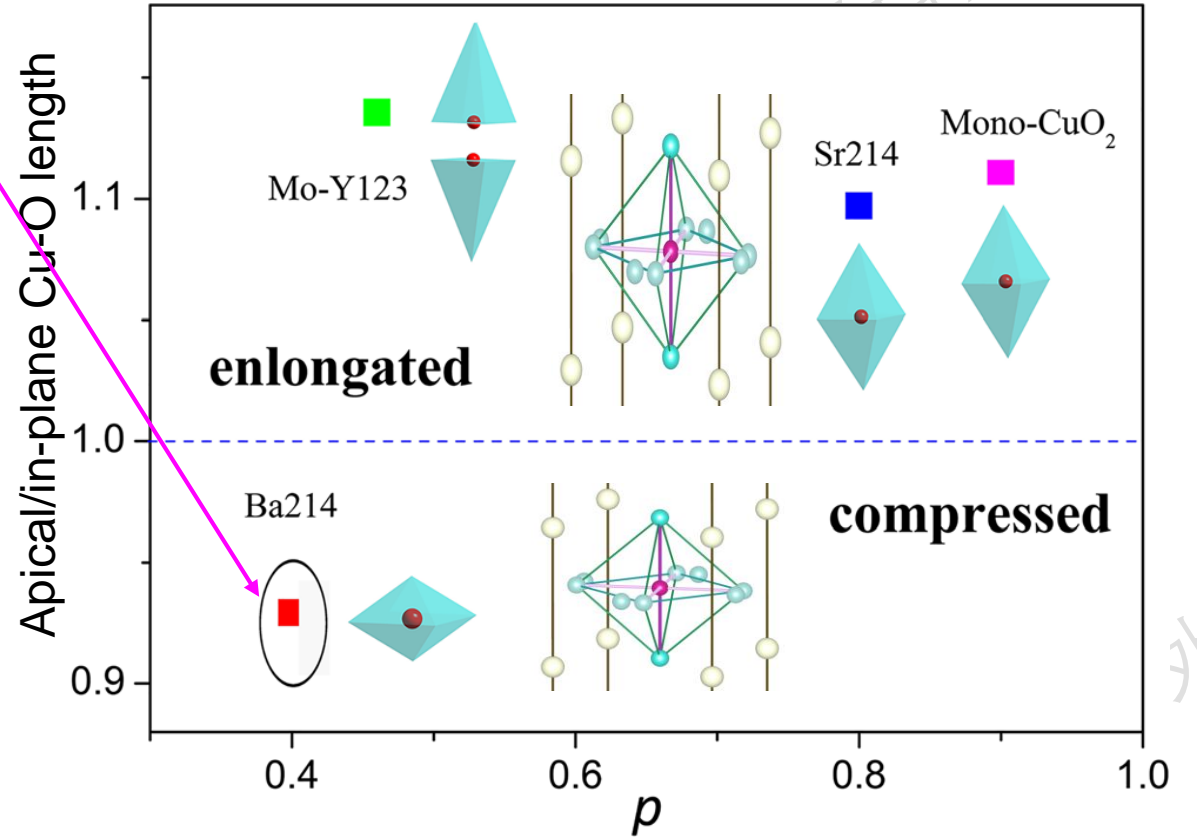
温度因子反映原子位置的静态位移和占有率

$\text{Ba-O1: } 2.751(6)$ $\text{Cu-O1: } 1.868(5) \times 2;$
 $2.8238(2) \times 4$ $\text{Cu-O2: } 1.9967(1) \times 2.4$
 $\text{Ba-O2: } 2.715(4) \times 2.4$

 $V(\text{Cu}^{2+}\text{-O11})=0.920 \times 2=1.84$
 $V(\text{Cu}^{2+}\text{-O12})=0.387 \times 2=0.774$
 $V(\text{Cu}^{1+}\text{-O11})=0.728 \times 2=1.456$
 $V(\text{Cu}^{1+}\text{-O12})=0.307 \times 2=0.307$

 $V(\text{Cu}^{2+}\text{-O2})= 0.398 \times 2.4=0.955$
 $V(\text{Cu}^{1+}\text{-O12})=0.315 \times 2.4=0.756$

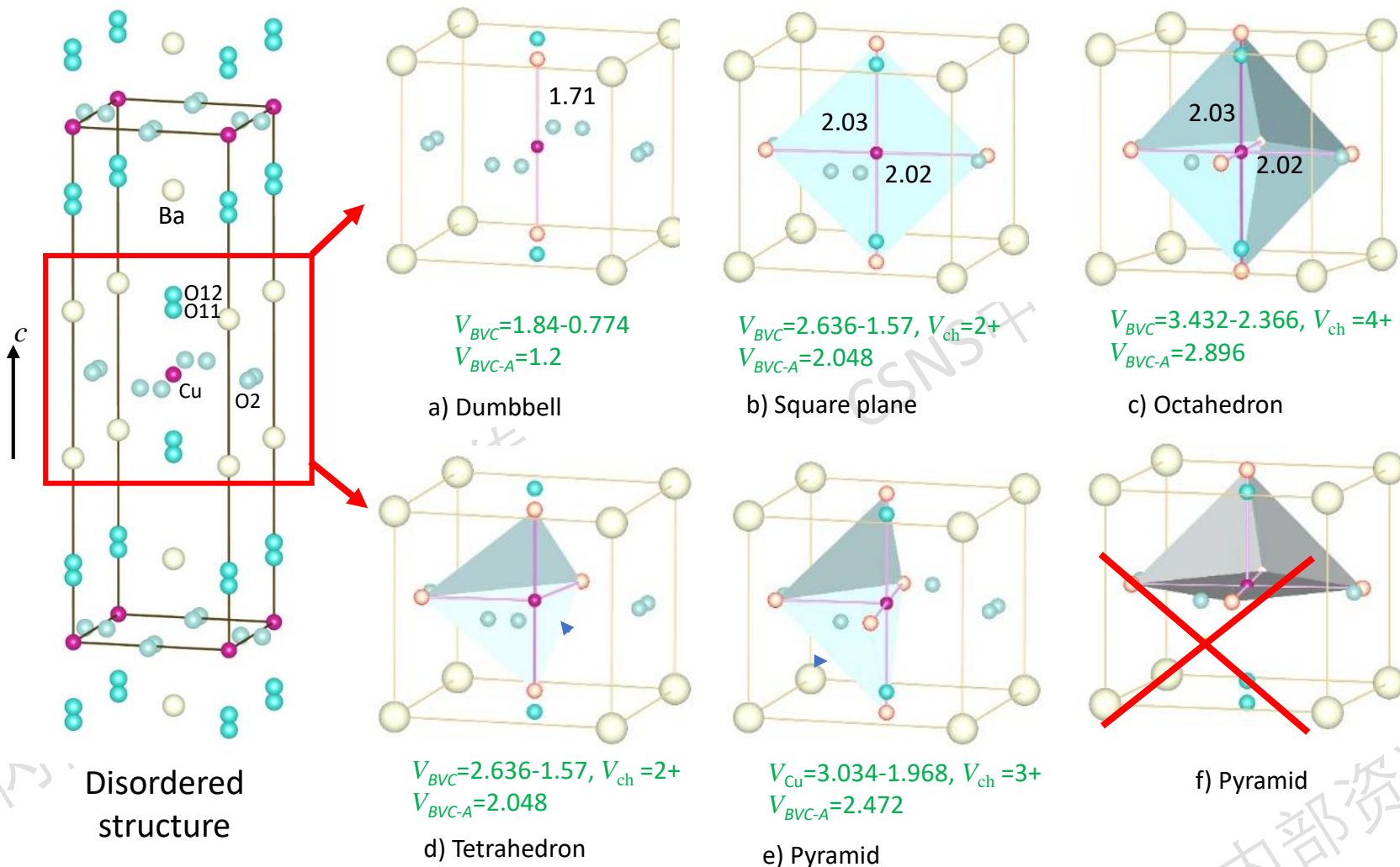
 $V(\text{Cu}^{2+})=2.217$
 $V(\text{Cu}^{1+})=1.711$



晶体结构研究发现Ba214与其它214等在结构上最大的不同是氧八面体沿着c方向被压缩，使得在面上的Cu-O键长比顶端的Cu-O键长更长。

The ratio between the bond lengths of in-plane Cu–O and copper apical oxygen of heavily over doped cuprates $\text{Cu}_{0.75}\text{Mo}_{0.25}\text{Sr}_2\text{YCu}_2\text{O}_{7.54}$ (Mo-Y123), monolayer CuO_2 (Mono-CuO₂), and $\text{Sr}_2\text{CuO}_{4-\delta}$ (Sr214), compared with Ba214.

Crystal structure and possible Cu-O_x polyhedron in Ba₂CuO_{3.2}



Possible polyhedron

根据实验测得的键长，用表中的数据计算离子的价态

$$V_i = \sum_j v_{ij}, v_{ij} = \exp[(R_{ij} - d_{ij})/b], e = 2.71828, b = 0.37, d_{ij} \text{ is bond distance}$$

Recommended bond-valence parameters for oxides, fluorides and chlorides

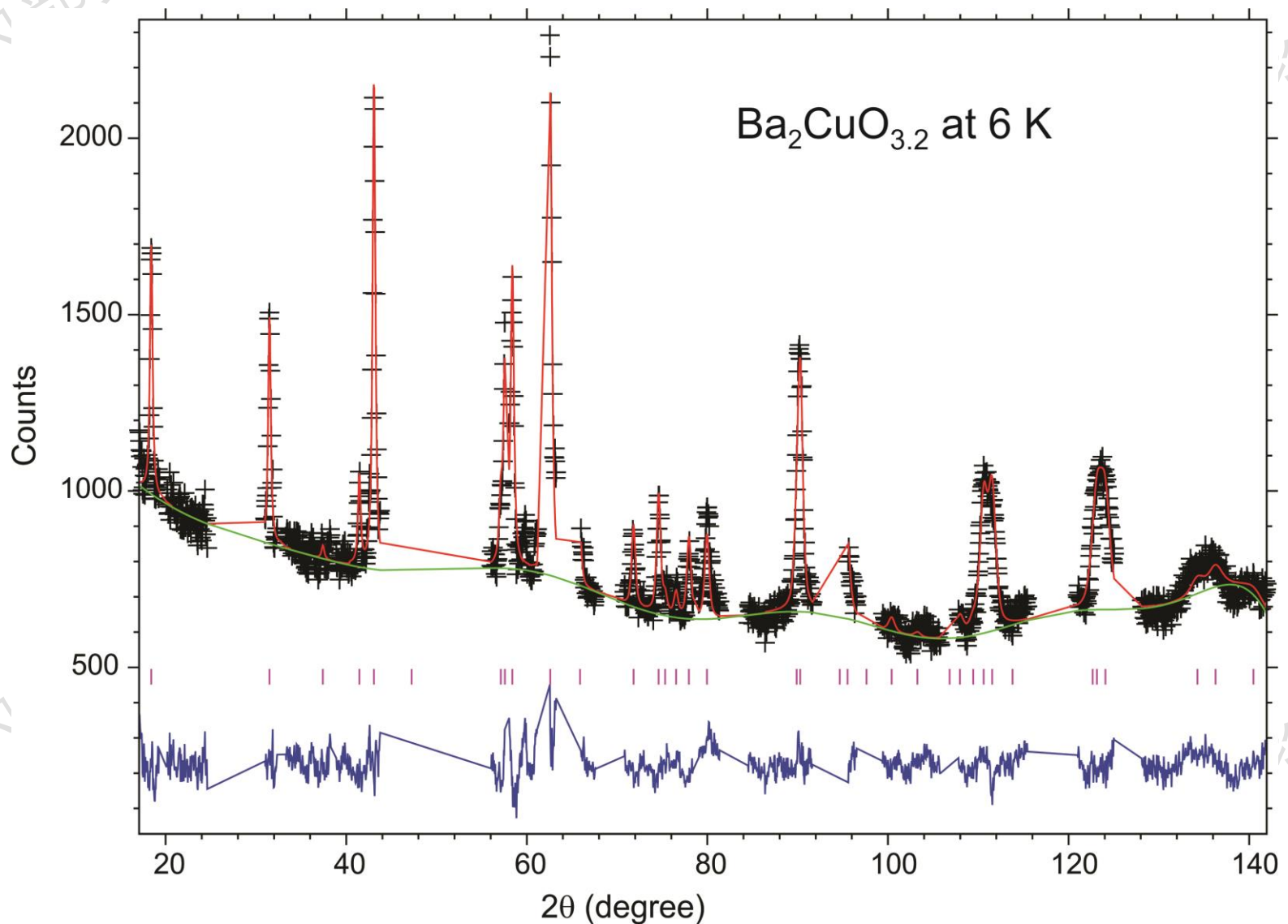
Cation	O	F	Cl	Cation	O	F	Cl
Ac ^{III}	2.24	2.13	2.63	Mn ^{IV}	1.753	1.71	2.13
Ag ^I	1.805	1.80	2.09	Mn ^{VII}	1.79	1.72	2.17
Al ^{III}	1.651	1.545	2.03	Mo ^{VI}	1.907	1.81	2.28
Am ^{III}	2.11	2.00	2.48	N ^{III}	1.361	1.37	1.75
As ^{III}	1.789	1.70	2.16	N ^V	1.432	1.36	1.80
As ^V	1.767	1.62	2.14	Na ^I	1.80	1.677	2.15
Au ^{III}	1.833	1.81	2.17	Nb ^V	1.911	1.87	2.27
B ^{III}	1.371	1.31	1.74	Nd ^{III}	2.117	2.008	2.492
Ba ^{II}	2.29	2.19	2.69	Ni ^{II}	1.654	1.599	2.02
Be ^{II}	1.381	1.28	1.76	Os ^{IV}	1.811	1.72	2.19
Bi ^{III}	2.09	1.99	2.48	P ^V	1.604	1.521	1.99
Bi ^V	2.06	1.97	2.44	Pb ^{II}	2.112	2.03	2.53
Bk ^{III}	2.08	1.96	2.46	Pb ^{IV}	2.042	1.94	2.43
Br ^{VII}	1.81	1.72	2.19	Pd ^{II}	1.792	1.74	2.05
Cl ^{IV}	1.39	1.32	1.76	Pr ^{III}	2.135	2.022	2.50
Ca ^{II}	1.967	1.842	2.37	Pt ^{II}	1.768	1.68	2.05
Cd ^{II}	1.904	1.811	2.23	Pt ^{IV}	1.879	1.759	2.17
Ce ^{III}	2.151	2.036	2.52	Pu ^{III}	2.11	2.00	2.48
Ce ^{IV}	2.028	1.995	2.41	Rb ^I	2.26	2.16	2.65
Cr ^{III}	2.07	1.95	2.45	Re ^{VII}	1.97	1.86	2.23
Cl ^{VII}	1.632	1.55	2.00	Rh ^{III}	1.791	1.71	2.17
Cm ^{III}	2.23	2.12	2.62	Ru ^{IV}	1.834	1.74	2.21
Co ^{II}	1.692	1.64	2.01	S ^{IV}	1.644	1.60	2.02
Co ^{III}	1.70	1.62	2.05	S ^{VI}	1.624	1.56	2.03
Cr ^I	1.73	1.67	2.09	Sb ^{III}	1.973	1.90	2.35
Cr ^{III}	1.724	1.64	2.08	Sb ^V	1.942	1.80	2.30
Cr ^V	1.794	1.74	2.12	Sc ^{III}	1.849	1.76	2.23
Cs ^I	2.42	2.33	2.79	Se ^{IV}	1.811	1.73	2.22
Cu ^I	1.593	1.6	1.85	Se ^{VI}	1.788	1.69	2.16
Cu ^{II}	1.679	1.60	2.00	Si ^{IV}	1.624	1.58	2.03

Cation	O	F	Cl	Cation	O	F	Cl
Dy ^{III}	2.036	1.922	2.41	Sm ^{III}	2.088	1.977	2.466
Er ^{III}	2.010	1.906	2.39	Sn ^{II}	1.984	1.925	2.36
Eu ^{II}	2.147	2.04	2.53	Sn ^{IV}	1.905	1.84	2.28
Eu ^{III}	2.076	1.961	2.455	Sr ^{II}	2.118	2.019	2.51
Fe ^{II}	1.734	1.65	2.06	Ta ^V	1.920	1.88	2.30
Fe ^{III}	1.759	1.67	2.09	Tb ^{III}	2.049	1.936	2.427
Ga ^{III}	1.730	1.62	2.07	Te ^{IV}	1.977	1.87	2.37
Gd ^{III}	2.065	1.95	2.445	Te ^{VI}	1.917	1.82	2.30
Ge ^{IV}	1.748	1.66	2.14	Th ^{IV}	2.167	2.07	2.55
H ^I	0.95	0.92	1.28	Ti ^{III}	1.791	1.723	2.17
Hf ^{IV}	1.923	1.85	2.30	Ti ^{IV}	1.815	1.76	2.19
Hg ^I	1.90	1.81	2.28	Tl ^I	2.172	2.15	2.56
Hg ^{II}	1.93	1.90	2.25	Tl ^{III}	2.003	1.88	2.32
Ho ^{III}	2.023	1.908	2.401	Tm ^{III}	2.000	1.842	2.38
I ^V	2.00	1.90	2.38	U ^{IV}	2.112	2.034	2.48
I ^{VII}	1.93	1.83	2.31	U ^{VI}	2.075	1.966	2.46
In ^{III}	1.902	1.79	2.28	V ^{III}	1.743	1.702	2.19
Ir ^V	1.916	1.82	2.30	V ^{IV}	1.784	1.70	2.16
K ^I	2.13	1.99	2.52	V ^V	1.803	1.71	2.16
La ^{III}	2.172	2.057	2.545	W ^{VI}	1.921	1.83	2.27
Li ^I	1.466	1.360	1.91	Y ^{III}	2.014	1.904	2.40
Lu ^{III}	1.971	1.876	2.361	Yb ^{III}	1.985	1.875	2.371
Mg ^{II}	1.693	1.581	2.08	Zn ^{II}	1.704	1.62	2.01
Mn ^{II}	1.790	1.698	2.13	Zr ^{IV}	1.937	1.854	2.33
Mn ^{III}	1.760	1.66	2.14				

Cu^{III}-O, R=1.724; Co^{IV}-O, R=1.75; Fe^{IV}-O, R=1.78;

Ni^{III}-O, R=1.68; Ni^{IV}-O, R=1.72

N.E.Brese and M.O'Keefe. Bond-Valence Parameters for Solids. Acta Cryst., B47, 192-197(1991)



资料, 请勿外传

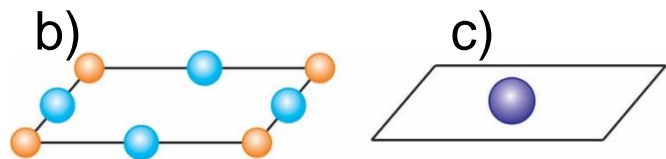
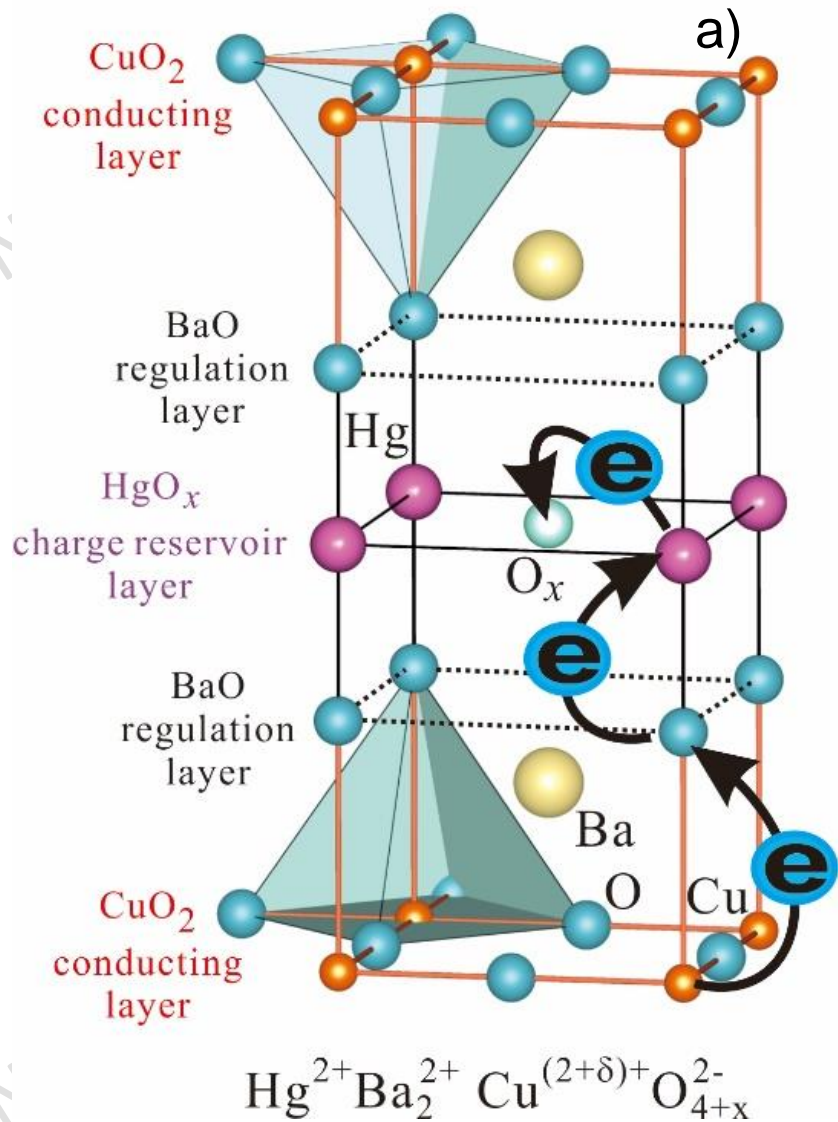
Oxygen dependence of the crystal structure of $\text{HgBa}_2\text{CuO}_{4+\delta}$ and its relation to superconductivity

CSNS中子学校内部资料, 请勿外传

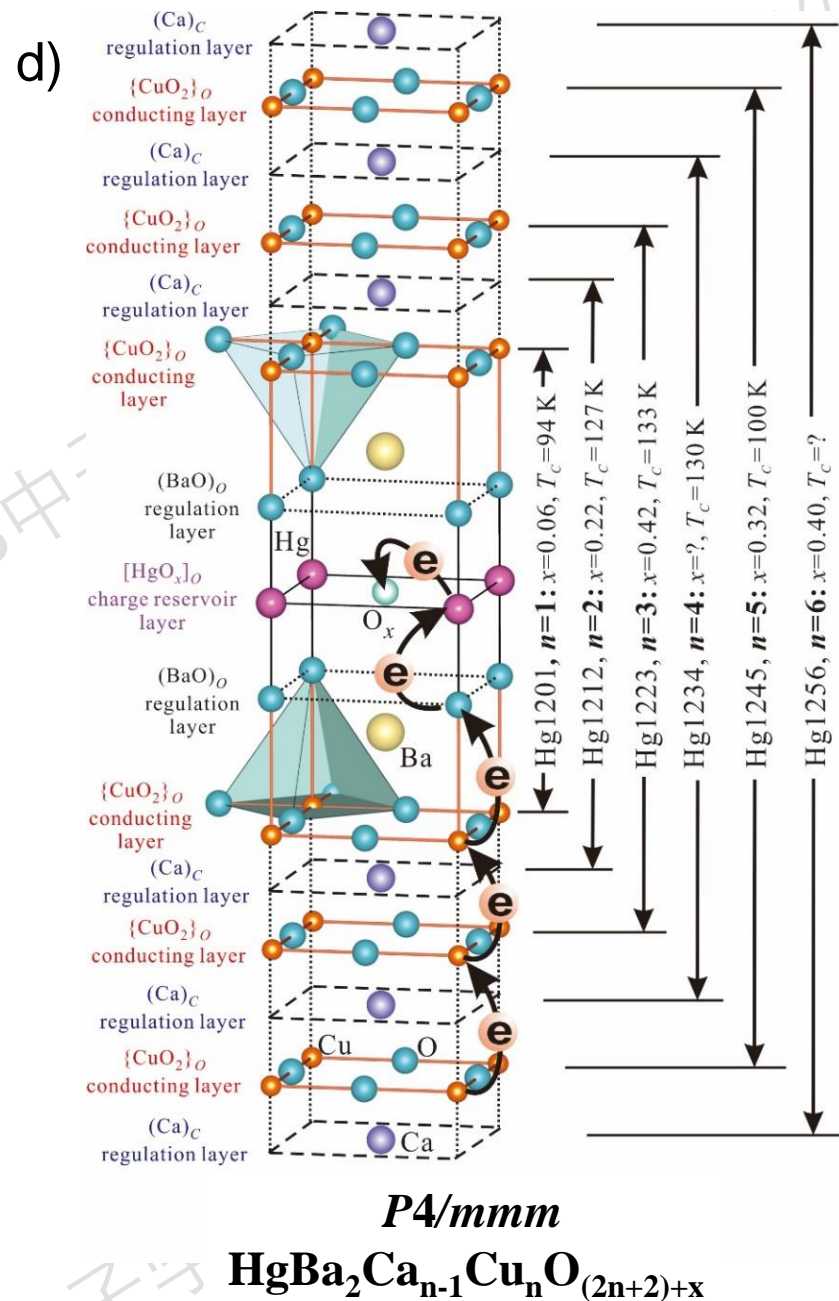
Q. HUANG, J. W. LYNN, Q. XIONG, AND C. W. CHU, Oxygen dependence of the crystal structure of $\text{HgBa}_2\text{CuO}_{4+z}$ and its relation to superconductivity. PRB, VOL. 52, NUMBER 11, JULY 1995-I

中子学校内部资料, 请勿外传

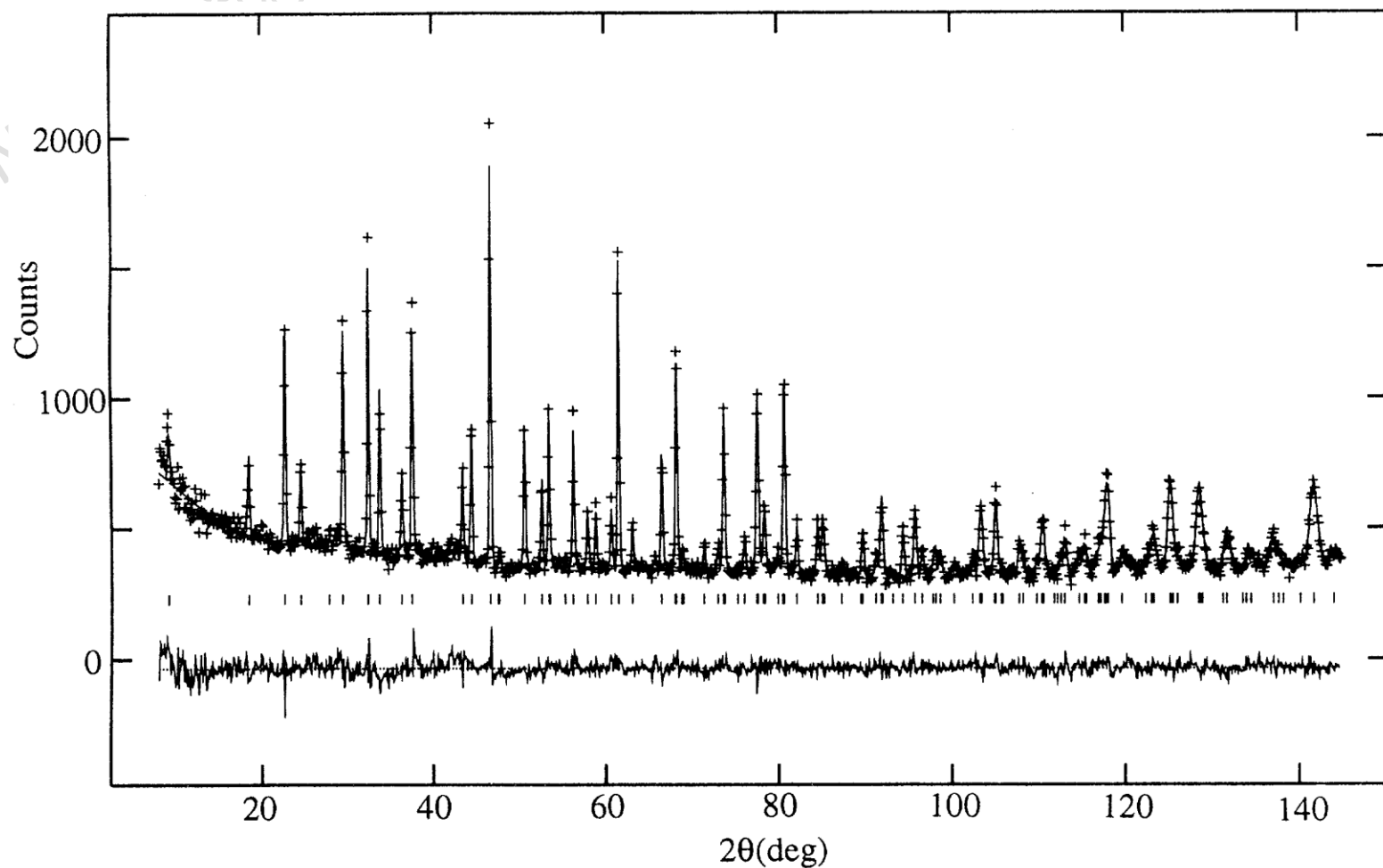
层状汞系铜基高温超导体 $\text{HgBa}_2\text{Ca}_{n-1}\text{Cu}_n\text{O}_{(2n+2)+x}$ 结构分析



以图a的 $\text{HgBa}_2\text{CuO}_{4+x}$ 的结构为基础，在c轴方向叠加图b和c的平面构成图d的三维层状结构层状汞系铜基高温超导体系列化合物 $\text{HgBa}_2\text{Ca}_{n-1}\text{Cu}_n\text{O}_{(2n+2)+x}$ 是。最高超导温度达到133 K。如图所示的电荷转移示意。当 HgO_x 层的 $x=0$ 时，化合物不超导。因此，准确测定 HgO_x 层中的O的占有率是研究化合物结构与性能的关系的关键。电荷转移可以用半经验方法计算Cu的价态得以估计。同时可以根据层间距估计化合物晶胞c轴的长度。

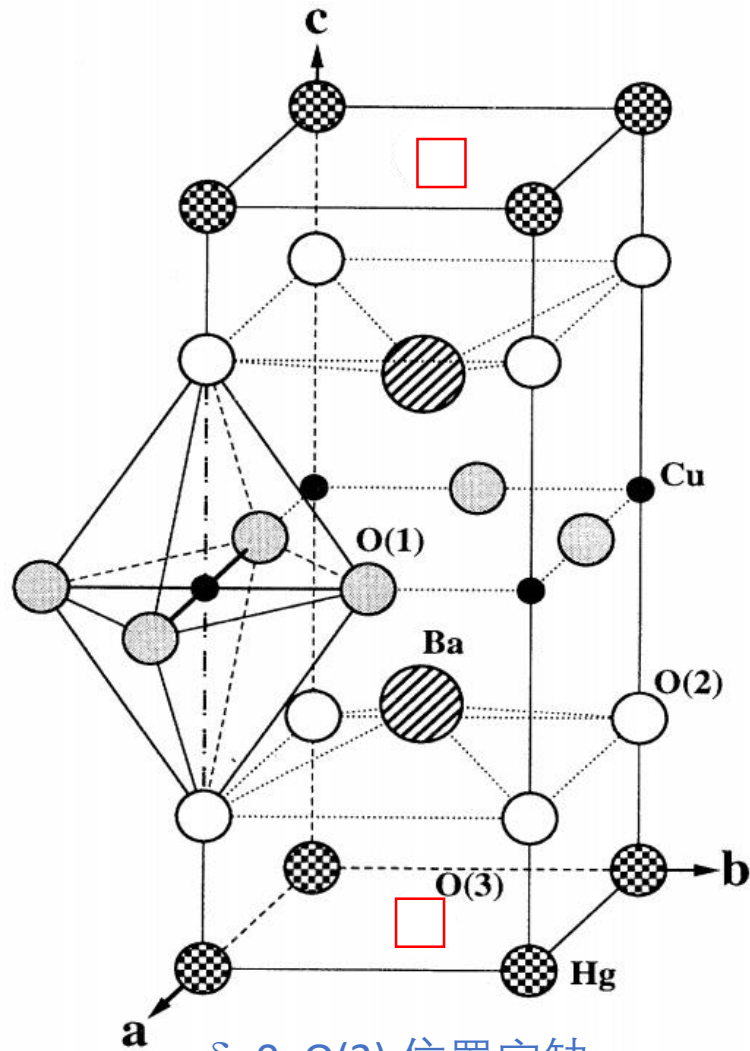


Oxygen dependence of the crystal structure of HgBa₂CuO_{4+δ} and its relation to superconductivity



Rietveld refinement profile of the neutron powder diffraction data obtained on the as-prepared sample of HgBa₂CuO_{4+δ} at 296 K. The observed data are represented by the points (+), and the refinement by the solid curve. The positions of the Bragg peaks are also indicated. The difference between the observations and the calculated profile is shown at the bottom of the figure.

Oxygen dependence of the crystal structure of HgBa₂CuO_{4+δ} and its relation to superconductivity



$\delta=0$, O(3) 位置空缺
化合物不超导

TABLE IV. (Top) Occupancy parameters of Hg (at room temperature). (Bottom) Anisotropic temperature factors of Ba ($B_{22}=B_{11}$) and Ba splitting into $[\frac{1}{2}, \frac{1}{2}, z(1)]$ and $[\frac{1}{2}, \frac{1}{2}, z(2)]$; constraints: $n_{\text{Ba}(2)} = n_{\text{O}(3)} = 1 - n_{\text{Ba}(1)}$, $B_{\text{Ba}(1)} = B_{\text{Ba}(2)}$ (at room temperature).

Sample label (δ)/[T_c (K)]	S1-AP (0.18)/(95)	S1-AV (0.08)/(53)	S1-RAO (0.18)/(94)	S1-RAV (0.04)/(0)	S2-PHP (0.21)/(80)	S1-RAOHP (0.23)/(30)
n	0.99(1)	0.97(1)	0.92(1)	0.95(1)	0.90(1)	0.91(1)
B (\AA^2)	1.37(5)	1.15(7)	1.13(6)	1.07(6)	0.84(8)	1.12(6)
Sample label (δ)/[T_c (K)]	S1-RAV (0.04)/(0)	S1-AV (0.08)/(53)	S1-AP (0.18)/(95)	S2-PHP (0.21)/(80)	S1-RAOHP (0.23)/(30)	
B_{11} (\AA^2)	0.84(7)	0.50(8)	0.51(7)	0.5(1)	0.72(6)	
B_{33} (\AA^2)	0.8(1)	0.9(1)	1.2(1)	1.0(1)	1.3(2)	
B_{33}/B_{11}	0.95	1.8	2.4	2.0	1.8	
		Ba splitting				
Ba(1)	$z(1)$	0.3045(5)	0.3025(7)	0.3001(10)	0.3006(7)	
	n	0.92(1)	0.83(1)	0.79(1)	0.77(1)	
Ba(2)	$z(2)$	0.276(7)	0.275(3)	0.284(4)	0.281(2)	
	n	0.08(1)	0.17(1)	0.21(1)	0.23(1)	
	B (\AA^2)	0.61(8)	0.59(6)	0.64(9)	0.57(6)	

Oxygen dependence of the crystal structure of HgBa₂CuO_{4+δ} and its relation to superconductivity

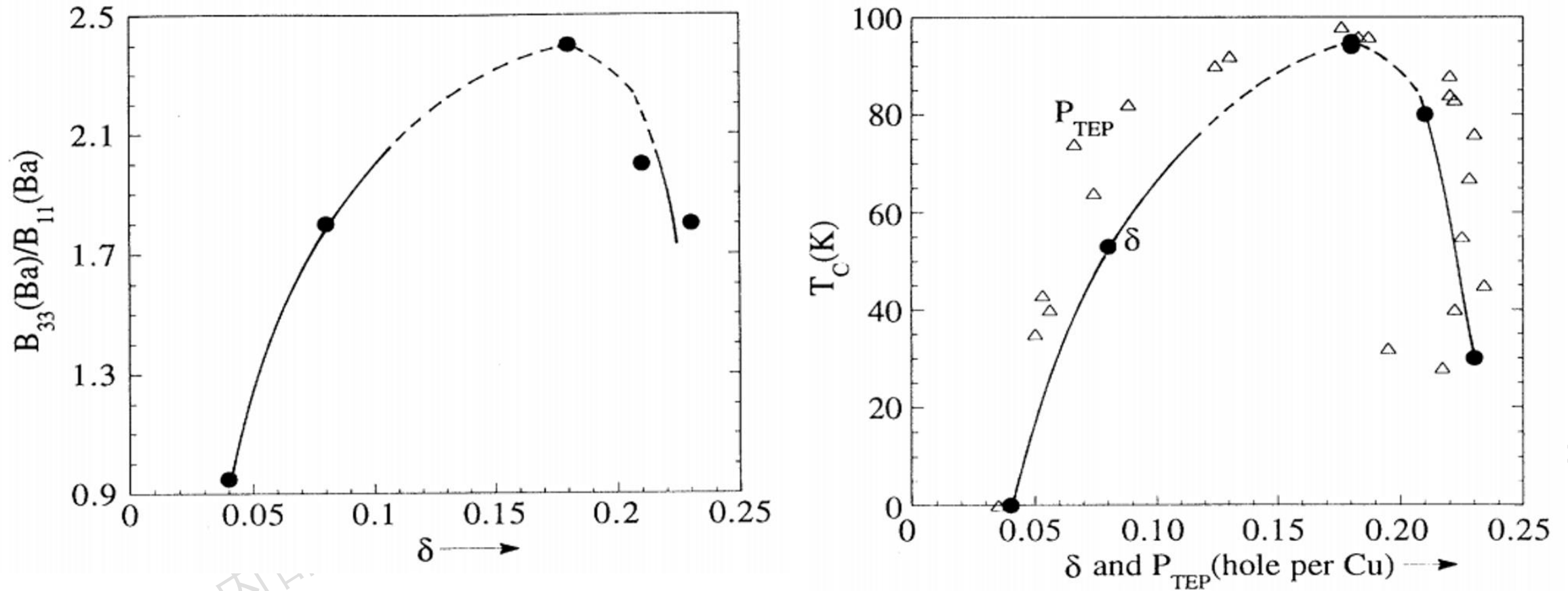
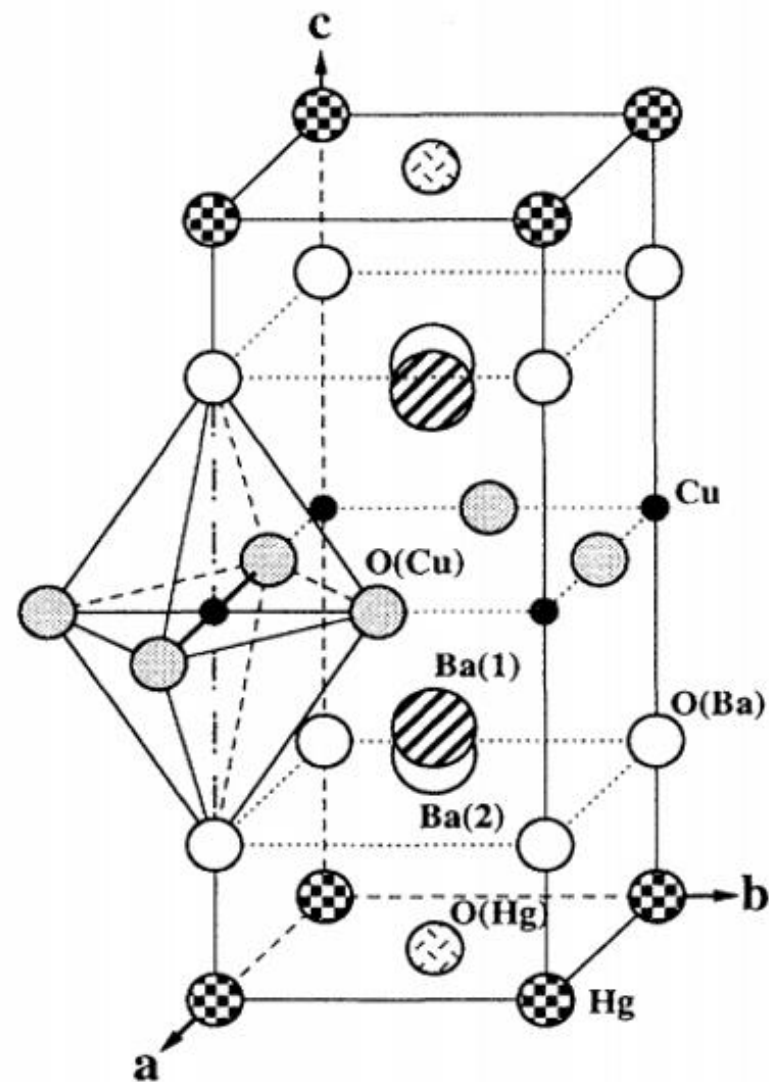
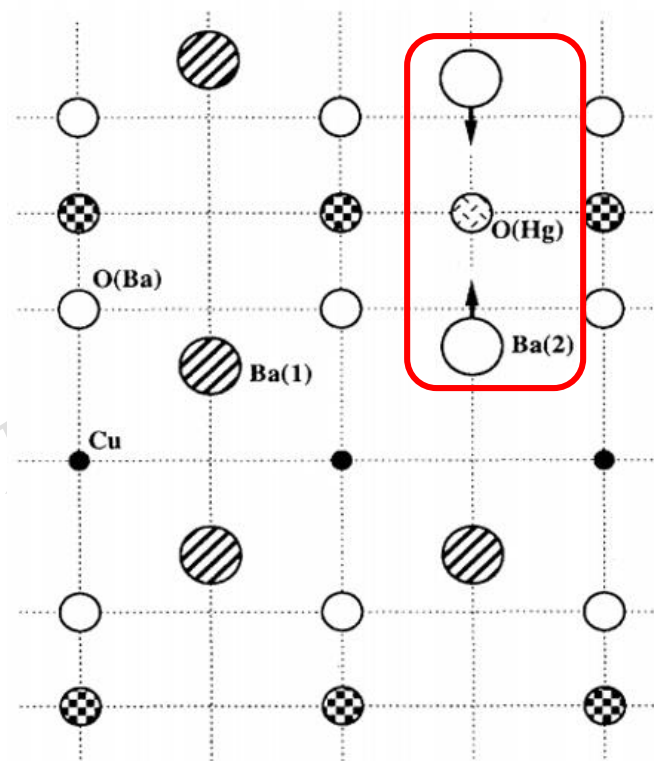
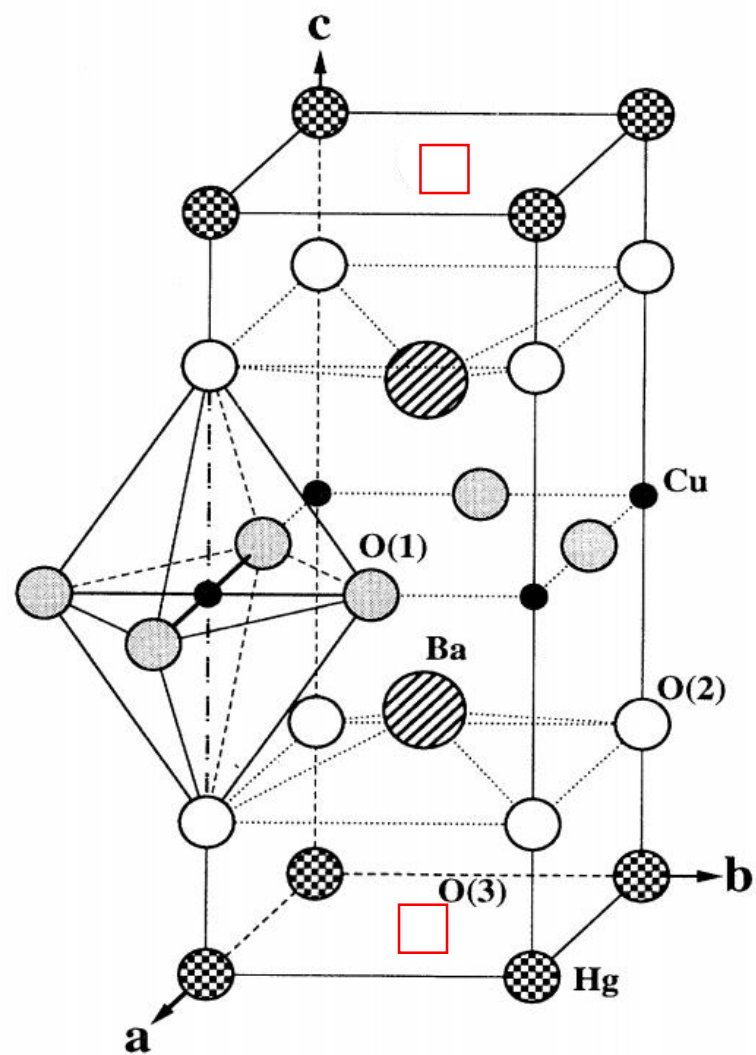
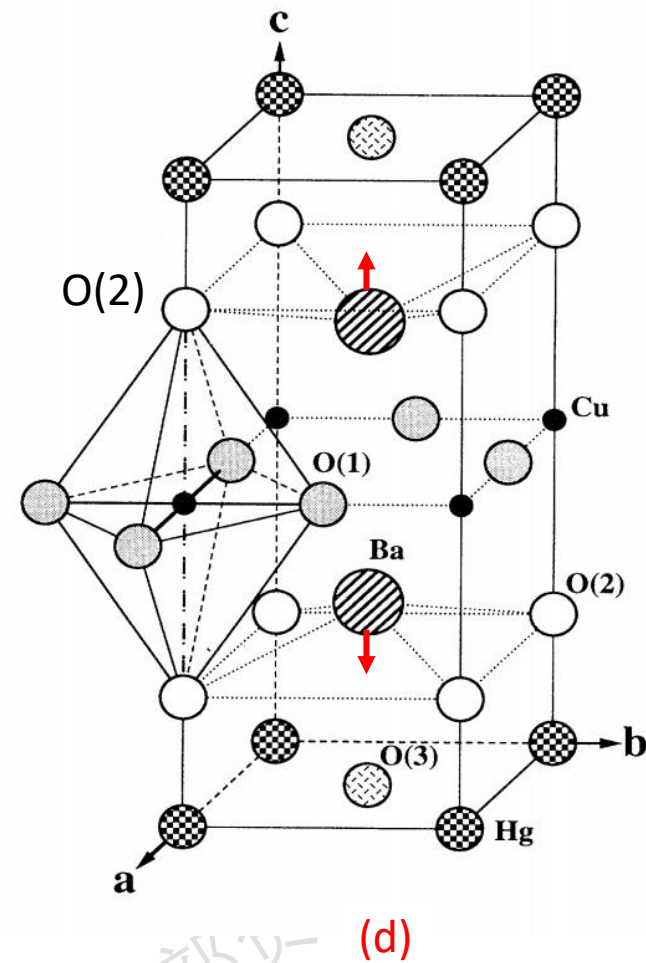
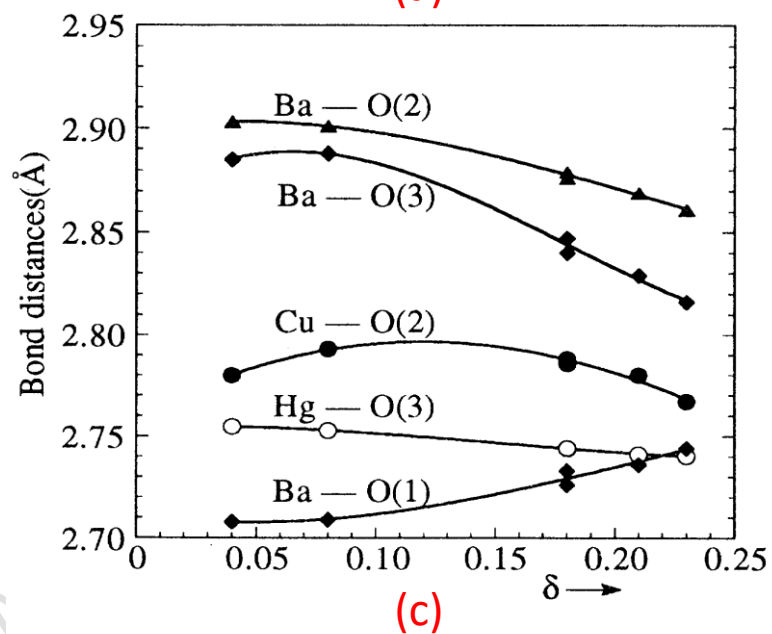
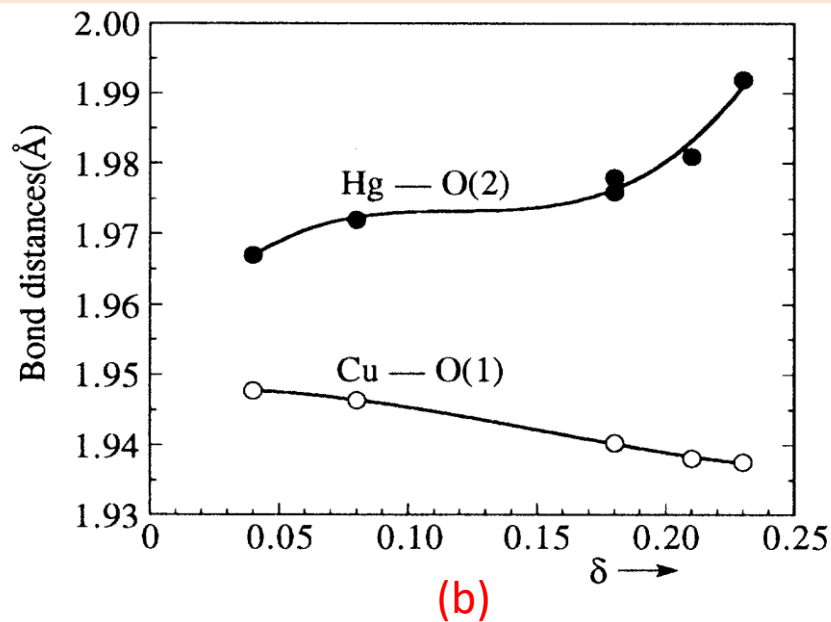
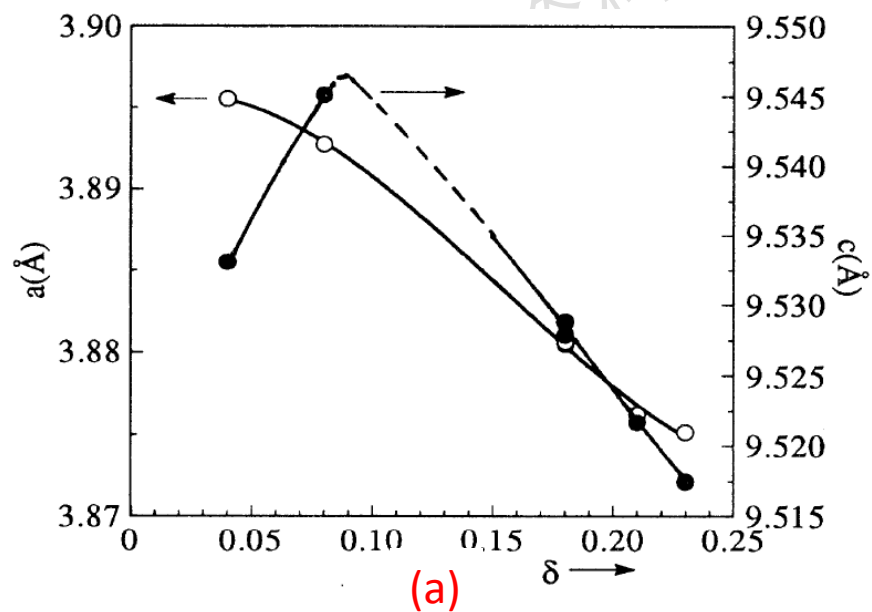


FIG. 8. (a) The ratio $B_{33}(\text{Ba})/B_{11}(\text{Ba})$ plotted versus oxygen content δ . (b) The behavior of T_c vs δ and T_c vs hole concentration. The solid points are from this study, while the triangles are taken from Ref. 15. The solid curves are simply a guide to the eye for the neutron data; parabolic fits to the T_c vs hole concentration are given in Ref. 15.

Oxygen dependence of the crystal structure of $\text{HgBa}_2\text{CuO}_{4+\delta}$ and its relation to superconductivity

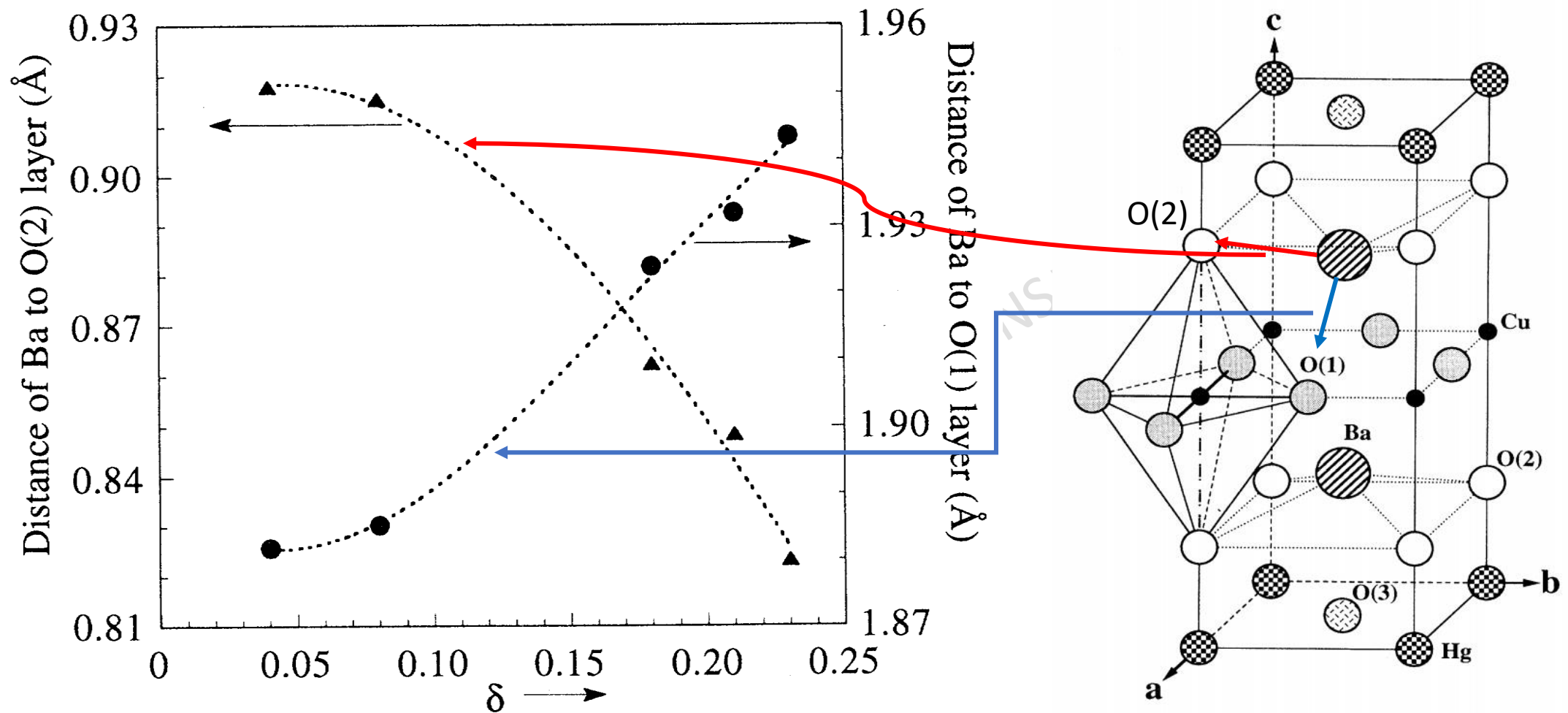


Oxygen dependence of the crystal structure of HgBa₂CuO_{4+δ} and its relation to superconductivity



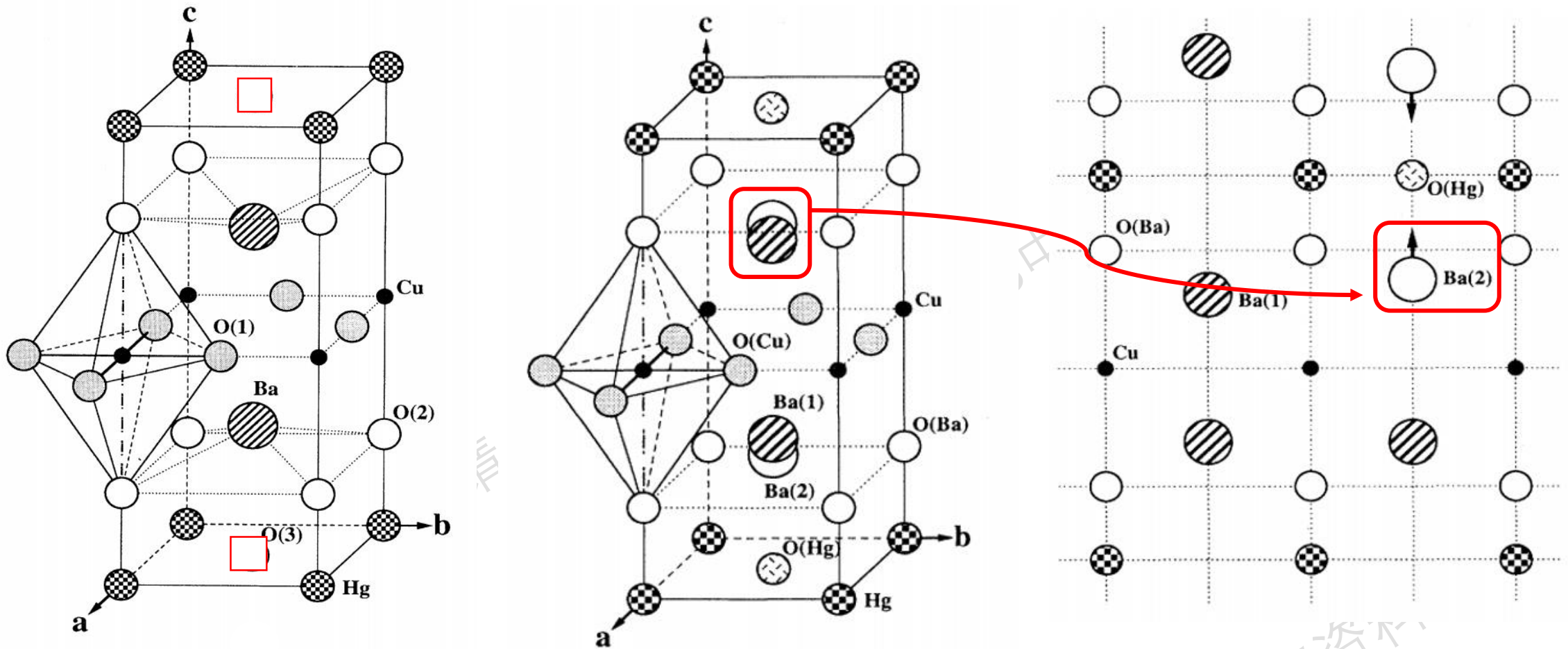
(a) Lattice parameters.
 (b), (c) selected bond distances,
 versus the oxygen content δ .
 (d) Structure of which shows
 that the Ba atoms shift toward
 the O(3) as δ increases.

Oxygen dependence of the crystal structure of $\text{HgBa}_2\text{CuO}_{4+\delta}$ and its relation to superconductivity



Distance of the Ba ion (z coordinate) from the O(1) and O(2) planes of oxygen ions, showing that the Ba ion moves towards the Q(3) site with increasing δ .

Oxygen dependence of the crystal structure of $\text{HgBa}_2\text{CuO}_{4+\delta}$ and its relation to superconductivity



(a) The parent compound structure $\text{HgBa}_2\text{CuO}_4$ where no oxygen on the Hg layer. (b) Two-site model [Ba(1) and Ba(2)] for the Ba ions in $\text{HgBa}_2\text{CuO}_{4+\delta}$, (c) the section of $a=1/2$ (or $b=1/2$). When the O(3) site is occupied, the Ba moves away from the Cu layer and towards the Hg layer.

Oxygen dependence of the crystal structure of HgBa₂CuO_{4+δ} and its relation to superconductivity

TABLE V. Valences (u.v.–unit valence) calculated from cation-oxygen bonds.

Sample label (δ)/[T_c (K)]	S1-RAV (0.04)/(0)	S1-AV (0.08)/(53)	S1-AP (0.18)/(95)	S2-PHP (0.21)/(80)	S1-RAOHP (0.23)/(30)
ΣV_{Hg} (u.v.)	1.827	1.824	1.836	1.836	1.795
ΣV_{Ba} (u.v.)	2.064	2.072	2.075	2.085	2.083
ΣV_{Cu} (u.v.)	2.036	2.040	2.075	2.088	2.094
$\Sigma V_{\text{O}(1)}$ (u.v.)	-2.259	-2.259	-2.195	-2.193	-2.166
$\Sigma V_{\text{O}(2)}$ (u.v.)	-1.719	-1.710	-1.749	-1.758	-1.755
$\Sigma V_{\text{O}(3)}$ (u.v.)	-0.831	-0.830	-0.896	-0.913	-0.931

$$V_i = \sum_j v_{ij}, \quad v_{ij} = \exp[(R_{ij} - d_{ij})/b], \quad e = 2.71828, \quad b = 0.37, \quad d_{ij} \text{ is bond distance}$$

6.1. Survey of computer programs for powder diffraction

C. J. Gilmore, J. A. Kaduk and H. Schenk

Table 6.1.1

Software for powder diffraction

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
ACerS–NIST Phase Equilibria Diagrams Database	Phase-equilibria diagrams	Database	ACerS–NIST Phase Equilibria Diagrams Database (2013)		Commercial
ADM	Software suite including device control, pattern evaluation, qualitative and quantitative phase analysis, indexing, lattice-parameter refinement, crystal-size evaluation, microstress analysis, profile analysis and pattern simulation	Phase analysis, indexing, lattice-parameter refinement, crystal size, stress, profile analysis, pattern simulation		http://www.RMSKempton.de/	Commercial
American Mineralogist Crystal Structure Database (AMCSD)	Online database that includes every structure published in <i>The American Mineralogist</i> , <i>The Canadian Mineralogist</i> , <i>European Journal of Mineralogy</i> and <i>Physics and Chemistry of Minerals</i> , as well as selected data sets from other journals	Minerals, database	Downs & Hall-Wallace (2003)	http://rruff.geo.arizona.edu/AMSamcsd.php	Free to use
ANAELU	Computer-based tools for inferring single-crystal structures and fibre textures from two-dimensional diffraction diagrams	Modelling, glasses	Fuentes-Montero <i>et al.</i> (2011)	http://www.esrf.eu/computing/scientific/ANAELU/Anelu_Page.htm	Free
ARITVE	Modelling of glass structures using the Rietveld method	Modelling, glasses	Le Bail (1995, 2000)	http://sdpd.univ-lemans.fr/aritve.html	Free
ATOMS	Structure visualization. Part of the <i>SHAPE</i> software package	Structure visualization	Shape Software, 521 Hidden Valley Road, Kingsport, TN 37663, USA	http://www.shapesoftware.com/	Commercial
AUTOFP	GUI for highly automated Rietveld refinement using an expert system algorithm based on <i>FULLPROF</i>	Rietveld refinement	Cui <i>et al.</i> (2015)	http://pmedia.shu.edu.cn/autofp	Free
AUTOX	Autoindexing multiphase samples; included with <i>VMRIA</i>	Indexing	Zlokazov (1995); Bergmann <i>et al.</i> (2004)	http://www.ccp14.ac.uk/ccp/web-mirrors/vmria	Free
AXES	Program for X-ray powder diffraction data evaluation, specially designed for peak-shape analysis and data preparation for Rietveld refinement in connection with <i>FULLPROF</i>	Data conversion, peak-shape analysis	Mändar <i>et al.</i> (1996)	http://www.ccp14.ac.uk/ccp/web-mirrors/axes/~hugo/axes/	Free for academic use

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>AXES-20B</i>	Estimation of crystal size and shape. Links to a wide range of programs and includes a range of data-processing and display functions	Phase identification, data conversion, structure visualization, peak location, peak profiling, indexing	Mändar & Vajakas (1998); Mändar <i>et al.</i> (1999)	http://www.ccp14.ac.uk/ccp/web-mirrors/axes/	Free
<i>Balls&Sticks</i>	Structure visualization and animation	Structure visualization	Ozawa & Kang (2004)	http://www.toycrate.org/	Free
<i>BALSAC</i>	Construction, visualization and interactive analysis of crystal lattices	Structure visualization	K. Hermann, Fritz-Haber-Institut der MPG, Berlin, Germany	http://www.fhi-berlin.mpg.de/~hermann/Balsac/	Free
<i>BEARTEX</i>	Texture analysis in polycrystalline materials	Texture analysis	Wenk <i>et al.</i> (1998)	http://eps.berkeley.edu/~wenk/TexturePage/beartex.htm	Free
<i>BGMN</i>	Fundamental parameters, Rietveld refinement and quantitative-analysis software	Fundamental parameters, Rietveld refinement, quantitative analysis	Bergmann <i>et al.</i> (1998)	http://www.bgm.de/methods.html	Free
Bilbao Crystallographic Server	Variety of symmetry tools including space-group retrieval tools, Wyckoff positions of space groups and group-subgroup relations	Symmetry	Aroyo <i>et al.</i> (2011)	http://www.cryst.ehu.es/	Free
B-IncStrDB	Database of incommensurate structures	Database	Kroumova <i>et al.</i> (2005)	http://www.cryst.ehu.es/icsdb/	Free
<i>Bond Valence Wizard</i>	Predicts interatomic distances in crystals by the bond-valence method	Bond-length prediction	Orlov <i>et al.</i> (1998); Brown (1977)	http://orlov.ch/bondval/	Free
<i>BRASS 2</i>	Rietveld refinement, Fourier-map calculation and visualization software	Rietveld refinement, visualization	Birkenstock <i>et al.</i> (2003)	http://www.brass.uni-bremen.de/	Free
<i>BREADTH</i>	Analysis of diffraction line broadening	Line broadening	Balzar (1995)	http://www.du.edu/~balzar/breadth.htm	Free
Cambridge Structural Database	Database (including structures derived from powder data) with associated software and software suites; organics and organometallics	Database	Groom <i>et al.</i> (2016)	http://www.ccdc.cam.ac.uk/	Commercial
<i>CaRine</i>	Visualization of lattices, surfaces, interfaces and diffraction patterns	Visualization	C. Boudias & D. Monceau, 1989–2004, CaRine Crystallography	http://carine.crystallography.pagesperso-orange.fr/	Commercial

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>CASTEP</i>	Calculates the properties of materials from first principles	Physical properties	Segall <i>et al.</i> (2002)	http://www.castep.org/	Free for European academics
CCP14	Web page with links to powder diffraction software	Software database	Stephenson <i>et al.</i> (2006)	http://www.ccp14.ac.uk/	Free
<i>cctbx</i>	A set of basic programming tools for X-ray crystallography	Programming	Grosse-Kunstleve <i>et al.</i> (2002)	https://github.com/cctbx/cctbx_project	Free
<i>CDASH</i>	Runs <i>DASH</i> over a network of computers	Structure solution	Spillman <i>et al.</i> (2015)	https://www.ccdc.cam.ac.uk/Solutions/PowderDiffraction/Pages/DASH.aspx	Commercial
<i>Celref</i>	Program for refining the unit cell	Unit-cell refinement	Laugier & Bochu (2004)	http://www.ccp14.ac.uk/ccp/web-mirrors/lmgp-laugier-bochu/	Free
<i>CHECKCELL</i>	Indexing and space-group determination, links into <i>Crysfire</i>	Indexing, space-group determination		http://www.ccp14.ac.uk/tutorial/lmgp/chekcellb.htm	Free
<i>checkCIF</i>	Cif checking service	CIF		http://journals.iucr.org/services/cif/checking/checkfull.html	Free
Chemical Database Service (CDS)	EPSRC-funded access to a large number of scientific databases, including all the main crystallographic databases, to UK-based academics and students	Database	Fletcher <i>et al.</i> (1996)	http://cds.dl.ac.uk/	Free within UK
<i>CIFTOOLS</i>	Includes two programs: <i>pdCIFplot</i> to plot Rietveld results from pdCIF files, and <i>CIFEDIT</i> to create, browse through and edit CIF files	CIF		ftp://ftp.ncnr.nist.gov/pub/cryst/cif/	Free
<i>Cinema:Debye-Scherrer</i>	Interactive visualization of multi-data-set Rietveld analyses	Data visualization	Vogel <i>et al.</i> (2018)	https://github.com/cinemascience/cinema_debye_scherrer	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>Crystal Maker X</i>	Visualization, structure design, simulation and 3D printing package	Visualization, simulation	CrystalMaker Software Ltd, 5 Begbroke Science Park, Sandy Lane, Yarnton, OX5 1PF, UK	http://www.crystallmaker.com/	Commercial
<i>Crystal Studio</i>	Visualization including surfaces, interfaces, defects and simulation of diffraction data	Visualization, defects, data simulation	Crystalsoftcorp, PO Box 7006, Wattle Park, VIC 3128, Australia	http://www.crystalsoftcorp.com, http://www.crystal0studio.com/	Commercial
<i>Crystallographica</i>	A software toolkit for crystallography	Visualization, pattern simulation	Oxford Cryosystems (1997)	http://www.oxcryo.com/cg/	Free
Crystallography Open Database (COD)	Open-access database including predicted structures, covering organics, inorganics, metal–organics and minerals	Database	Grazulis <i>et al.</i> (2009)	http://www.crystallography.net/	Free
CRYSTMET	Database of metals, alloys and intermetallics	Database	White <i>et al.</i> (2002); Toth Information Systems, Inc., 2045 Quincy Avenue, Ottawa, Ontario, K1J 6B2, Canada	https://cds.dl.ac.uk/cds/datasets/crys/mdf/llmdf.html	Free to UK academic users
<i>DAJUST</i>	Intensity extraction and space-group determination. Part of the <i>XLENS</i> package	Space-group determination, cell-parameter refinement, profile fitting, Le Bail method	Vallcorba <i>et al.</i> (2012)	https://departments.icmab.es/crystallography/news/42-pdsoftware	Free to non-commercial users
<i>DANSE</i>	General software for neutron scattering	Neutron scattering	Cummings <i>et al.</i> (2002)	http://wiki.cacr.caltech.edu/danse/index.php/Main_Page	Free
<i>DASH</i>	Software for solving crystal structures from X-ray powder diffraction patterns	Structure solution	David <i>et al.</i> (2006)	http://www.ccdc.cam.ac.uk/products/powder_diffraction/dash/	Commercial
Database of Zeolite Structures	Provides structural information on all of the zeolite framework types that have been approved by the Structure Commission of the International Zeolite Association (IZA-SC)	Database	Baerlocher <i>et al.</i> (2003)	http://www.iza-structure.org/databases/	Free
<i>DataScan</i>	Diffractometer automation hardware and software	Instrumentation, diffractometer control	Materials Data, Inc., 2551 Second Street, Livermore, CA 94550, USA	http://www.materialsdata.com/ds.htm	Commercial

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>CMPR</i>	Displays diffraction data, manual and auto-indexing, and peak fitting	Data visualization, indexing, peak fitting	Toby (2005)	http://www.ncnr.nist.gov/xtal/software/cmpr/	Free
<i>CMWP-fit</i>	Determination of microstructural parameters from diffraction profiles	Profile fitting	Ribárik <i>et al.</i> (2001)	http://csendes.elte.hu/cmwp/	Free
<i>CONEX</i>	Angular calibration and averaging of two-dimensional powder diffraction patterns	Two-dimensional data	Gommes & Goderis (2010)	Available from the authors	Free
<i>Conograph</i>	Integrated system with a graphical user interface for indexing	Indexing	Esmaeili <i>et al.</i> (2017)	https://osdn.net/projects/conograph/releases/	Free
<i>ConQuest</i>	Software for searching the Cambridge Structural Database	Database searching	Bruno <i>et al.</i> (2002)	http://www.ccdc.cam.ac.uk/	Commercial
<i>ConvX</i>	Data-conversion program that can convert multiple files at once. Also possible to select a particular angular range to convert	Data conversion	Bowden (2000)	http://www.ccp14.ac.uk/ccp/web-mirrors/convx/	Free
<i>Corina</i>	Structure generator for drug-like molecules	Structure prediction	Molecular Networks GmbH, Nuremberg, Germany; Sadowski <i>et al.</i> (1994); Schwab (2010)	https://www.mn-am.com/products/corina	Commercial
<i>CRISP</i>	Uses crystallographic image processing to extract information from electron micrographs	Electron diffraction	Hovmöller (1992)	http://www.calidris-em.com/crisp.php	Commercial
<i>CRYSCA</i>	Crystal-structure calculations for flexible molecules	Structure prediction	Schmidt & Kalkhof (1997)	Available from the authors	
<i>Cryscon</i>	Allows conversion between several popular crystallographic file formats	Data conversion	Eric Dowry, Shape Software, 521 Hidden Valley Road, Kingsport, TN 37663, USA	http://www.shapesoftware.com/	Shareware
<i>Crysfire</i>	Automatic powder-pattern indexing	Indexing	Shirley (2002)	http://www.ccp14.ac.uk/tutorial/crysf/ , http://www.ccp14.ac.uk/ccp/web-mirrors/crysf-r-shirley/	Free
<i>CrystalDiffract</i>	Data processing and visualization	Peak fitting	CrystalMaker Software Ltd, 5 Begbroke Science Park, Sandy Lane, Yarnton, OX5 1PF, UK	http://crystalmaker.com/crystaldiffract/index.html	Commercial

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>Datasqueeze</i>	Graphical interface for analysing data from two-dimensional X-ray diffraction detectors	Two-dimensional data analysis	Heiney (2002)	http://www.physics.upenn.edu/~heiney/datasqueeze/index.html	Free
<i>DDM</i>	Refinement using the Rietveld method or derivative difference minimization	Rietveld refinement, derivative difference refinement	Solovyov (2004)	https://sites.google.com/site/ddmsuite/	Free
<i>Debussy 2.0</i>	Analysis of powder diffraction data from nanocrystalline, defective and/or nonperiodic materials	Nanocrystalline and non-periodic materials, defects	Cervellino <i>et al.</i> (2015)	http://debussy.sourceforge.net/debussy/debussy.html	Free
<i>DEBVIN</i>	Rietveld refinement with the use of generalized coordinates	Rietveld refinement, polymers	Brückner & Immirzi (1997)	https://vdocuments.site/debvin-a-program-for-Rietveld-refinement-with-generalized-coordinates.html	Free
<i>decryst</i>	Software suite for structure determination from powder diffraction using the direct-space method	Structure solution	Liu (2018)	https://gitlab.com/CasperVector/decryst/	Free
<i>Diamond</i>	Crystal and molecular structure visualization and exploration	Structure visualization	Crystal Impact – K. Brandenburg and H. Putz GbR, Postfach 1251, D-53002 Bonn, Germany	http://www.crystalimpact.com/	Commercial
<i>DICVOL</i>	Automatic indexing	Indexing	Boultif & Louër (2004)	http://www.ccp14.ac.uk/ccp/web-mirrors/dicvol/	Free
<i>DIFFaX</i>	Simulation of powder diffraction patterns for faulted materials	Stacking faults, simulation	Treacy <i>et al.</i> (1991)	http://www.public.asu.edu/~mtreacy/DIFFaX.html	Free
<i>DIFFaX+</i>	A modified version of <i>DIFFaX</i> for the refinement of disordered stack materials	Refinement, disordered stack materials	Leoni <i>et al.</i> (2004)	Available from the author (matteo.leoni@unitn.it) on request	Free
<i>Diffpy-CMI</i>	Python software for diffraction analysis and the study of the atomic structure of materials	Modelling	See https://www.diffpy.org/publications.html	http://diffpy.org	Free
<i>DIFFRAC.EVA</i>	Data processing, phase identification and quantitative analysis. Cluster analysis allows high-throughput screening	Profile fitting, cluster analysis, quantitative phase analysis	Bruker AXS GmbH, Oestliche Rheinbrueckenstr. 49, D-76187 Karlsruhe, Germany	https://www.bruker.com/products/x-ray-diffraction-and-elemental-analysis/x-ray-diffraction/xrd-software/eva/overview.html	Commercial
<i>DIFFRAC.TOPAS</i>	Profile and structure analysis (see <i>TOPAS</i>)	Profile analysis, Rietveld refinement	Bruker AXS GmbH, Oestliche Rheinbrueckenstr. 49, D-76187 Karlsruhe, Germany	https://www.bruker.com/products/x-ray-diffraction-and-elemental-analysis/x-ray-diffraction/xrd-software/topas.html	Commercial

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>DISCUS</i>	Simulates defect structures and diffuse scattering, plus the simulation of perfect structures of non-crystalline materials	Defect structures, simulation	Proffen & Neder (1997, 1999)	https://www.xray.cz/ecm-cd/soft/xray/general/discus/discus/disfeat.html	Free
<i>DLS-76</i>	Structure refinement with distance constraints; part of <i>FOCUS</i>	Rietveld refinement	Baerlocher <i>et al.</i> (1976, 1977)	https://github.com/stefsmeeets/focus_package , http://www.crystal.mat.ethz.ch/Software/index.html	Free
<i>DRAWxtl</i>	Open-source program for producing crystal-structure drawings	Visualization	Finger <i>et al.</i> (2007)	http://www.lwfinger.net/drawxtl/	Free
<i>DRXWin</i>	Diffraction data manipulation and qualitative analysis	Visualization, qualitative phase analysis, peak search	Vicent Primo Martín, El Instituto de Ciencia de los Materiales, Universitat de Valencia, Avda. Blasco Ibáñez, 13, 46010 Valencia, Spain	http://www.ccp14.ac.uk/ccp/web-mirrors/drxwin/drxwin/	Free
<i>Dysnomia</i>	Calculates electron and nuclear densities from observed structure factors by the maximum-entropy method	Electron and nuclear density calculations	Momma <i>et al.</i> (2013)	http://jp-minerals.org/dysnomia/en/	Free
<i>Eager</i>	Genetic algorithm for crystal structure solution from powder diffraction data	Structure solution	Harris <i>et al.</i> (2004)	Contact K. D. M. Harris	
<i>EFLECH/INDEX</i>	Whole-pattern indexing	Indexing	Bergmann (2007)	http://www.bgm.de/indexpgm.html	Free
<i>ELD</i>	Quantitative information from electron-diffraction patterns	Electron diffraction	Calidris, Sweden	http://www.calidris-em.com/eld.php	Commercial
<i>enCIFer</i>	Viewing and editing CIF files	CIF editing	Allen <i>et al.</i> (2004)	https://www.ccdc.cam.ac.uk/Community/csd-community/encifer/	Free
<i>Endeavour</i>	Direct-space method for <i>ab initio</i> solution of structures from powder diffraction data by combined global optimization of pattern difference and potential energy	Structure solution	Putz <i>et al.</i> (1999)	http://www.crystalimpact.com/endeavour/	Commercial
<i>EPCryst</i>	Structure solution from powder diffraction data using models generated by an equivalent-position combination algorithm based on unit-cell content and space-group information	Structure solution	Deng & Dong (2011)	http://www.epcryst.com	Free
<i>ESPOIR</i>	Structure solution by Monte Carlo analysis of powder diffraction data	Structure solution	Le Bail (2001)	http://www.cristal.org/sdpd/espoir/	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>EXPGUI</i>	Graphical interface for <i>GSAS</i>	Rietveld refinement, visualization	Toby (2001)	http://www.ncnr.nist.gov/programs/crystallography/software/expgui/	Free
<i>EXPO2014</i>	Structure-solution toolkit for phasing crystal structures from powder data	Structure solution	Altomare <i>et al.</i> (2013)	http://www.ba.ic.cnr.it/softwareic/expo/expo2014-installation/	Free for non-profit users
<i>ExtSym</i>	Space-group assignment. Bundled with <i>DASH</i> but can be obtained separately from Anders Markvardsen	Space-group determination	Markvardsen <i>et al.</i> (2001)	http://www.markvardsen.net/projects/ExtSym/main.html	Free
<i>FABLE</i>	Reads many X-ray detector image formats	Data processing	Schmidt <i>et al.</i> (2009)	http://sourceforge.net/projects/fable/	Free
<i>FAULTS</i>	Refinement of structures with extended defects, based on <i>DIFFaX</i>	Defect structures, simulation	Casas-Cabanas <i>et al.</i> (2016)		Free
<i>FIT</i>	Decomposition of powder diffraction patterns and profile analysis of pair correlation functions	Profile analysis, PDF analysis	Petkov & Bakaltchev (1990)	http://web.pa.msu.edu/people/petkov/software.html	Free for academic use
<i>FIT2D</i>	Calibration and correction of detector distortions, integration of two-dimensional data to a variety of one-dimensional scans, and one- and two-dimensional model fitting	Data analysis, visualization	Hammersley (1998, 2016); Hammersley <i>et al.</i> (1996)	http://www.esrf.fr/computing/scientific/FIT2D/	Free
<i>FitAllb</i>	Fitting centre-of-mass grain positions, orientation and elastic strain tensors	Strain	Oddershede <i>et al.</i> (2010)	https://sourceforge.net/p/fable/wiki/FitAllB/	Free
<i>FitGISAXS</i>	Modelling and analysis of grazing-incidence small-angle X-ray scattering data	GISAXS	Babonneau (2010)	https://www.pprime.fr/?q=en/node/1013;contact D. Babonneau, david.babonneau@univ-poitiers.fr	Free
<i>FlexCryst</i>	Program suite including crystal-structure prediction, structure determination and clustering	Structure prediction, structure determination	Hofmann & Apostolakis (2003)	http://www.flexcryst.com/html/home.pdf	Commercial
<i>FOCUS</i>	Automated structure determination of zeolites. Can use electron-diffraction data	Structure determination, zeolites	Grosse-Kunstleve <i>et al.</i> (1997)	https://github.com/stefsmets/focus_package	Free
<i>FOX</i>	<i>Ab initio</i> structure determination using global-optimization algorithms, suitable for any type of crystal structure (organic, inorganic including automatic correction of special positions); can combine several data sets	Structure solution	Favre-Nicolin & Cerný (2002)	http://objcryst.sourceforge.net/Fox/	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>FOXGrid</i>	Structure solution using <i>FOX</i> with distributed computing	Structure solution	Rohliček <i>et al.</i> (2007)	https://fox.vincetn.net/Manual/Fox.Grid	Free
<i>FULLPAT</i>	Full-pattern quantitative phase analysis merging the advantages of existing full-pattern fitting methods with the traditional reference intensity ratio method. Coded into Excel	Quantitative phase analysis	Chipera & Bish (2002)	http://www.ccp14.ac.uk/ccp/web-mirrors/fullpat/	Free
<i>FULLPROF</i>	Rietveld refinement and pattern matching	Rietveld refinement, pattern matching	Rodriguez-Carvajal (1990)	https://www.ill.eu/sites/fullprof/	Free
<i>GDASH</i>	Structure solution using grid computing	Structure solution	Griffin <i>et al.</i> (2009)	Contact Tom Griffin, tom.griffin@stfc.ac.uk	
<i>GEST</i>	Structure solution using a genetic algorithm	Structure solution	Feng & Dong (2007)	Contact Zhen Jie Feng, fengzhenjie@gmail.com	Free
<i>GETSPEC</i>	Calculates the symmetry operators and special positions for any setting of any space group based on the Hall space-group symbol	Symmetry	Altermatt & Brown (1987)	http://www.ccp14.ac.uk/ccp/web-mirrors/i_d_brown/getspec/	Free
<i>Gnu Xtal System</i>	Set of portable crystallographic routines	Software development		http://xtal.sourceforge.net/	Free
<i>GrainSpotter</i>	Indexing grains in monophase crystalline materials	Indexing	Schmidt (2014)	https://sourceforge.net/p/fable/wiki/grainspotter/	Free
<i>Gretep</i>	Displacement-ellipsoid plot structure viewing and analysis. Part of the <i>LMGP</i> suite	Visualization	Laugier & Bochu (2004)	http://www.ccp14.ac.uk/tutorial/lmgp/#gretep , http://www.ccp14.ac.uk/ccp/web-mirrors/lmgp-laugier-bochu/	Free
<i>GRINSP</i>	Inorganic structure prediction	Structure prediction	Le Bail (2005)	http://www.cristal.org/grinsp/	Free
<i>GSAS</i>	Comprehensive system for the refinement of structural models to both X-ray and neutron diffraction data. Superseded by <i>GSAS-II</i>	Pattern fitting, structure solution, Rietveld refinement, visualization	Larson & Von Dreele (2004)	http://subversion.xray.aps.anl.gov/trac/EXPGUI	Free
<i>GSAS-II</i>	Open-source Python project that addresses all types of crystallographic studies, from simple materials through macromolecules, using both powder and single-crystal diffraction and with both X-ray and neutron probes	Pattern fitting, structure solution, Rietveld refinement, visualization	Toby & Von Dreele (2013)	https://subversion.xray.aps.anl.gov/trac/pyGSAS	Free
<i>gsaslanguage</i>	A language that provides input to and processes output from the <i>GSAS</i> package	Scripting	Vogel (2011)	http://code.google.com/p/gsaslanguage/	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>GSASIIscriptable</i>	Scripting interface for <i>GSAS-II</i>	Scripting	O'Donnell <i>et al.</i> (2018)	https://gsas-ii.readthedocs.io/en/latest/GSASIIscriptable.html	Free
<i>GudrunN</i> and <i>GudrunX</i>	Programs for correcting raw neutron and X-ray total-scattering data to differential cross sections	Data processing, neutron diffraction, total scattering	Soper & Barney (2011)	In-house software at ISIS	
<i>HELIX</i>	Simulation of helical diffraction	Powder-pattern calculation	Knupp & Squire (2004)	https://www.diamond.ac.uk/Instruments/Soft-Condensed-Matter/small-angle/SAXS-Software/CCP13.html	Free
Hypertext book of crystallographic space-group diagrams and tables	Space-group diagrams	Space-group information	Cockcroft (1999)	http://img.chem.ucl.ac.uk/sgp/	Commercial
ICDD Powder Diffraction File	Comprehensive powder database: PDF-4+, PDF-4 organics and PDF-2, plus derivatives	Database, phase identification	International Centre for Diffraction Data, 12 Campus Boulevard, Newtown Square, PA, 19073–3273, USA	http://www.icdd.com/	Commercial
<i>IGOR Pro</i>	Scientific data-analysis software	Data analysis	Kline (2006)	https://www.wavemetrics.com/products/igorpro	Commercial
<i>ImageD11</i>	Python codes for processing diffraction data from 2D detectors at the ID11 beamline at the ESRF	Data processing, neutron diffraction	Wright (2011)	http://sourceforge.net/projects/fable/files/ImageD11	Free
Inorganic Crystal Structure Database (ICSD)	Database of inorganic structures	Database	Belsky <i>et al.</i> (2002)	http://www2.fiz-karlsruhe.de/icsd_home.html	Commercial

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>input4MAUD</i>	Input of two-dimensional data into <i>MAUD</i>	Data conversion	Raue (2014)	http://kristall.uni-mki.gwdg.de/Docs/raue/input4MAUD/main.html	Free
<i>International Tables Online</i>	Online editions of all volumes of <i>International Tables for Crystallography</i> , including Volume A (<i>Space-Group Symmetry</i>), and an extensive symmetry database	Space-group information, database	<i>International Tables Online</i> (2019)	https://it.iucr.org	Commercial
<i>iotbx.cif</i>	Input, output and validation of CIFs	CIF	Gildea <i>et al.</i> (2011)	https://cci.lbl.gov/cctbx_docs/iotbx/iotbx.cif.html	Free
<i>ISAACS</i>	Interactive structure analysis of amorphous and crystalline systems	Structure analysis, amorphous materials	Le Roux & Petkov (2010)	http://isaacs.sourceforge.net	Free
<i>IsoDEC</i>	Calculation of diffraction elastic constants	Strain	Gnäupel-Herold (2012)	https://www.ncnr.nist.gov/xtal/software/isodec	Free
<i>IsoStar</i>	Information on intermolecular interactions derived from small-molecule crystal structures in the Cambridge Structural Database and from protein–ligand interactions observed in the Protein Data Bank (PDB)	Database	Bruno <i>et al.</i> (1997)	https://www.ccdc.cam.ac.uk/solutions/csd-system/components/isostar/	Commercial
<i>ISOTROPY</i>	Applies group-theoretical methods to the analysis of phase transitions in crystalline solids	Space-group determination	Stokes & Hatch (2002)	http://stokes.byu.edu/isotropy.html	Free web service
<i>ITO</i>	Powder-pattern indexing. Incorporated into many other packages, both free and commercial	Indexing	Visser (1969)	http://sdpd.univ-lemans.fr/ftp/ito13.zip	Free/commercial
<i>JADE</i>	Comprehensive powder diffraction data-processing suite. <i>JADE Standard</i> is available as perpetual license and <i>JADE Pro</i> as an annual subscription	Cluster analysis, data conversion, indexing, Le Bail fitting, Pawley fitting, peak location, peak profiling, phase identification, powder-pattern viewing, quantitative phase analysis, Rietveld refinement, size–strain analysis	Materials Data, Inc. (2016)	http://www.materialsdata.com/products.htm ; http://www.icdd.com/	Commercial

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>Jana2006</i>	Analysis of incommensurate, modulated and composite structures, as well as standard Rietveld refinement	Le Bail fitting, Rietveld refinement, Fourier-map calculation and visualization, structure solution	Petríček <i>et al.</i> (2000); Dusek <i>et al.</i> (2001)	http://jana.fzu.cz/	Free
<i>JEMS</i>	Simulation of high-resolution TEM (HRTEM), CBED, PED and SAED patterns, and simulation of powder diffraction rings	Electron diffraction, simulation	Stadelmann (2003)	http://www.jems-saas.ch/Home/jemsWebSite/jems.html	Free for students
<i>J-ICE</i>	Visualization of crystallographic and electronic properties using <i>Jmol</i>	Visualization, data conversion	Canepa <i>et al.</i> (2011)	http://j-ice.sourceforge.net	Free
<i>Jpowder</i>	JAVA data viewer	Data visualization	Markvardsen <i>et al.</i> (2010)	https://sourceforge.net/projects/jpowder/	Free
<i>KoalaRiet</i>	Powder-pattern viewing, peak profiling and Rietveld refinement, fundamental parameters approach	Pattern visualization, peak fitting, Rietveld refinement	Cheary & Coelho (1996)	http://www.ccp14.ac.uk/	Free
<i>LABOTEX</i>	Crystallographic texture analysis of rocks, semiconductors and superconductors, ceramics and composites, polymers, metals and alloys	Texture analysis	Pawlik & Özga (1999)	http://www.labosoft.com.pl/	Commercial
<i>LAPOD</i>	Refinement of lattice parameters using optimal regression	Unit-cell refinement	Dong & Langford (2000)	http://www.ccp14.ac.uk/ccp/web-mirrors/powderx/lapod/	Free
<i>LauePt</i>	GUI for Laue-pattern simulation and analysis	Data analysis, simulation	Huang (2010)	http://www.ccp14.ac.uk/ccp/web-mirrors/xianrong-huang/	Free
<i>MacDiff</i>	Analysis and display of powder data on Macintosh platforms	Database, phase identification, data conversion	Petschick (2001)	http://www.geol-pal.uni-frankfurt.de/Staff/Homepages/Petschick/MacDiff/MacDiffInfoE.html	Free
<i>MATCH!</i>	Compares the diffraction pattern of the unknown to a database containing reference patterns; additional knowledge such as known phases, elements or density can also be used	Phase identification, quantitative phase analysis	Crystal Impact – K. Brandenburg and H. Putz GbR, Postfach 1251, D-53002 Bonn, Germany	http://www.crystalimpact.com/match/	Commercial
<i>Materials Studio</i>	Multipurpose suite	Indexing, model building, structure solution, visualization		http://accelrys.com/products/collaborative-science/biovia-materials-studio/	Commercial

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>MAUD</i>	Rietveld-based program for combined analysis. Can be used to fit diffraction, fluorescence and reflectivity data collected using X-rays, neutrons or electrons	Data integration, indexing, space-group assignment, texture analysis, size-strain analysis, Rietveld refinement, quantitative phase analysis	Lutterotti <i>et al.</i> (1999)	http://maud.radiographema.eu/	Free
<i>MCE-Marching Cubes</i>	Interactive visualization of electron and other density maps, optimized for small molecules	Fourier-map calculation and visualization	Husák & Kratochvíl (2003)	https://sourceforge.net/projects/mce-marching-cube-eld/	Free
<i>McMaille</i>	Indexing using Monte Carlo methods	Indexing	Le Bail (2004)	http://www.cristal.org/McMaille	Free
<i>Mercury</i>	Statistical analysis of data from the Cambridge Structural Database, structure visualization and simulation of powder patterns	Structure visualization	Sykes <i>et al.</i> (2011)	http://www.ccdc.cam.ac.uk/products/mercury/	Free; commercial version has more functions
MINCRYST	Contains crystal-structure data for minerals	Database	Chichagov <i>et al.</i> (2001)	http://database.iem.ac.ru/mincryst/	Free
The Mineralogy Database	Includes information on the crystal structures, powder diffraction patterns, chemical composition, and physical and optical properties of minerals	Database		http://webmineral.com	Free
<i>Mogul</i>	Validation of bond lengths, angles and torsions against the Cambridge Structural Database	Structure validation	Bruno <i>et al.</i> (2004)	http://www.ccdc.cam.ac.uk/products/csd_system/mogul/	Commercial
<i>MOPAC2016</i>	Semi-empirical quantum chemistry calculations. <i>Mercury</i> has an interface to <i>MOPAC</i>	Structure validation, structure building	Stewart (2012)	http://openmopac.net	Free for academic use
<i>Mstruct</i>	Microstructure analysis using powder diffraction	Microstructure analysis	Matěj <i>et al.</i> (2014)	http://www.xray.cz/mstruct/	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>Multifit/Polydefit</i>	Open-source IDL software package for data analysis of synchrotron-based plasticity, rheology or other time-dependent experiments	Texture analysis	Merkel & Hilaret (2015)	http://merkel.texture.rocks/Multifit-Polydefix/	Free
<i>MultiRef</i>	<i>MATLAB</i> framework for <i>GSAS</i> ; Rietveld refinement of multiple powder diffractograms from <i>in situ</i> , scanning or diffraction tomography experiments	Rietveld refinement	Frølich & Birkedal (2015)	http://chem.au.dk/en/research/research-areas/inorganicchemistrymaterialschemistry/biological-and-bioinspired-materials/multiref/downloads/	Free
<i>NanoPDF64</i>	Software package for analysis of pair-distribution functions of nanocrystals, also capable of calculating theoretical powder diffraction patterns and pair-distribution functions for atomistic models of nanocrystals	PDF analysis	Skrobas <i>et al.</i> (2017)	http://www.unipress.waw.pl/nanopdf	Free
<i>NEWMOD II</i>	Modelling of interstratified layer structures	Powder-pattern calculation	Yuan & Bish (2010)	https://newmod-for-clays.com	Commercial
Nickel–Nichols Mineral Database	Mineral database	Phase identification	The Nickel–Nichols Mineral Database, Materials Data, Inc., 1224 Concannon Blvd, Livermore, CA 94550, USA; Nickel & Nichols (1991)	http://www.materialsdata.com/MINERALS.htm	Free
<i>Nika</i>	Open-source <i>Igor Pro</i> -based program for reduction of two-dimensional scattering data from area-detector small- and wide-angle scattering instruments. Can be used in transmission and grazing-incidence geometries	SAXS, WAXS	Ilavsky (2012)	https://usaxs.xray.anl.gov/software/nika	Free
<i>NXS</i>	Library for neutron cross-section calculations	Neutron diffraction	Boin (2012)	https://www.helmholtz-berlin.de/people/boin/nxs_en.html	Free
<i>OpenBabel</i>	Open-source chemistry toolbox, allowing users to search, convert, analyse or store data from molecular modelling, chemistry, solid-state materials, biochemistry or related areas	File conversion	O'Boyle <i>et al.</i> (2011)	http://openbabel.sourceforge.net/	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>OpenGenie</i>	Neutron-data analysis	Powder-pattern visualization	Akeroyd <i>et al.</i> (1999)	http://www.opengenie.org/Main_Page	Free
<i>PASCal</i>	Online tool for determining principal coefficients of thermal expansion and compressibilities using variable-pressure and variable-temperature lattice-parameter data	Strain	Cliffe & Goodwin (2012)	http://pascal.chem.ox.ac.uk	Free web access
<i>PATE</i>	GSAS powder-pattern extractor	Data conversion		http://www.ccp14.ac.uk/ccp/web-mirrors/scott-belmonte-software/pate/	Free
<i>PCED3</i>	Simulation of polycrystalline electron diffraction and phase identification	Electron diffraction, phase identification		http://www.unl.edu/ncmn-cfem/xzli/download	Free
PCOD	Open database of structures of inorganic compounds predicted or enumerated mainly using <i>ZEFSA II</i> or <i>GRINSP</i>	Phase identification, database		http://www.crystallography.net/pcod/	Free
<i>PDXL 2</i>	Provides various analysis tools such as automatic phase identification, quantitative analysis, crystallite-size analysis, lattice-constants refinement, Rietveld analysis, <i>ab initio</i> structure determination and data clustering. Peak shapes can be modelled using a fundamental-parameters approach	Rietveld refinement, quantitative analysis, phase identification, structure determination	Rigaku (2014)	https://www.rigaku.com/en/service/software/pdxl	Commercial
<i>PeckCryst</i>	Molecular crystal structure determination from powder diffraction data using a particle swarm optimization algorithm	Structure solution	Feng <i>et al.</i> (2009)	Contact J. C. Zhang, jczhang@shu.edu.cn	Free
<i>PLATON</i>	Implements a large variety of standard geometrical calculations and tests (<i>e.g.</i> for missing symmetry, voids in the lattice <i>etc.</i>), and includes utilities for cell transformation and graphics. Used in <i>checkCIF</i>	Data conversion, space-group assignment, structure refinement, H-atom placement, structure validation, powder-pattern calculation, structure visualization	Spek (1998, 2003)	http://www.cryst.chem.uu.nl/platon/	Free
<i>PM2K</i>	Full-pattern fitting allowing microstructural investigation of nanocrystalline materials. Includes analysis of size broadening due to an analytical or to a generic (unconstrained) distribution of crystallites, the presence of dislocations, stacking faults and antiphase boundaries	Whole-powder-pattern modelling	Leoni <i>et al.</i> (2006)	Contact M. Leoni, matteo.leoni@unitn.it	Free for academic use

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>PolySNAP3</i>	Clustering powder diffraction, XRF and spectroscopic data. Semi-quantitative analysis	Cluster analysis, phase analysis	Barr <i>et al.</i> (2009)	http://www.bruker.com/	Commercial
<i>popLA</i>	Pole-figure evaluation using the harmonic and WIMV methods	Texture analysis	Kocks <i>et al.</i> (1998)	http://lansce.lanl.gov/facilities/lujan/instruments/hippo/popLA.php	Free
<i>Powder Cell</i>	Represents the crystal structure and the corresponding powder pattern simultaneously. Translations and rotations of a preselected part of the asymmetric unit allow manipulation of the structure	Data conversion, powder-pattern calculation, powder-pattern viewing, Le Bail fitting, size-strain analysis, Rietveld refinement, quantitative phase analysis, structure visualization	Kraus & Nolze (1996)	http://powdercell-for-windows.software.informer.com/2.4/	Free
<i>Powder3D</i>	Multiple-pattern data reduction and graphical presentation. Part of <i>Materials Studio</i>	Data integration, data conversion, powder-pattern viewing, peak location	Hinrichsen <i>et al.</i> (2008)	https://www.fkf.mpg.de/4708688/Powder-3D-Software	Free
<i>PowderX</i>	Uses include plotting X-ray patterns, data smoothing, background subtraction, α_2 -peak elimination, peak search, indexing, zero-angle error correction and data-format conversions	Data conversion, powder-pattern viewing, peak location, indexing	Dong (1999)	http://www.ccp14.ac.uk/ccp/web-mirrors/powderx/Powder/	Free
<i>PowDLL</i>	.NET dynamic link library used for interconversion between variable formats of X-ray powder diffraction files	Data conversion	Kourkoumelis (2005)	http://users.uoi.gr/nkourkou/powdll.htm	Free
<i>PRODD</i>	Refinement of crystal and magnetic structures from powder diffraction data, can be used with time-of-flight and constant-wavelength neutron data as well as synchrotron X-ray data	Pawley fitting, Rietveld refinement, proteins	Wright (2004); Wright & Forsyth (2000); Wright <i>et al.</i> (2007)	http://www.ccp14.ac.uk/ccp/web-mirrors/prodd/~jpw22/	Free
<i>PROFEX</i>	A platform-independent open-source graphical user interface for the Rietveld refinement program <i>BGMN</i> . Reads and converts a variety of proprietary raw data formats	Rietveld refinement, data conversion	Doebelin & Kleeberg (2015)	http://profex.doebelin.org/	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>Profil</i>	Rietveld refinement and utilities. Can be used for organics and organometallics requiring restrained refinement	Unit-cell refinement, Rietveld refinement, Fourier-map calculation, visualization, data conversion	Cockcroft (1994)	http://img.chem.ucl.ac.uk/www/cockcroft/profil.htm	Free for academic use
<i>pubCIF</i>	Editing, validating and formatting CIF files	CIF editing	Westrip (2010)	http://journals.iucr.org/services/cif/pubcif/	Free
<i>PyCifRW</i>	Support for reading and writing of CIF files using Python, with optional validation against DDL dictionaries	CIF editing	Hester (2006)	https://pypi.org/project/PyCifRW/4.3/	Free
<i>PyMOL</i>	A comprehensive software package for rendering and animating 3D structures	Structure visualization		https://pymol.org/2/	Commercial
<i>Quanto</i>	Allows automatic estimation of the weight fraction of each crystalline phase in a mixture. The amorphous content can be estimated using the internal-standard method, and corrections for preferred orientation and microabsorption effects are available	Quantitative phase analysis	Altomare <i>et al.</i> (2001)	http://www.ic.cnr.it/	Free
<i>RIETAN-FP</i>	A package of programs for analysing crystal structures from X-ray and neutron powder diffraction data and visualizing the results in three dimensions, along with electronic-state calculations	Rietveld refinement, data conversion	Izumi & Momma (2007)	http://fujioizumi.verse.jp/download/download_Eng.html	Free
<i>Rietica</i>	Aids the creation and updating of Rietveld input files for and provides point-and-click control of the Rietveld program <i>LHPM</i>	Le Bail fitting, Rietveld refinement, visualization	Hunter (1998)	http://www.rietica.org/	Free
<i>RMC++</i>	Reverse Monte Carlo modelling of the structure of disordered materials, mainly using diffraction data, but additional information such as EXAFS data and different constraints can also be used	PDF analysis	Evrard & Pusztai (2005)	http://www.szfk.hu/~nphys/rmc++/opening.html	Free
<i>RMCAW95</i>	Uses the reverse Monte Carlo method for modelling both lattice and magnetic disorder in powder crystalline materials by direct calculation of the structure factor	PDF analysis	Møllergård & McGreevy (1999)	http://www.cristal.org/glasses/rmca/rmcaw95.html	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>RMCprofile</i>	Built from the original <i>RMCA</i> code of McGreevy & Puztai to determine the local structure of crystalline materials while still being capable of analysing disordered systems. Can be used to fit many data types (including total scattering, EXAFS, diffuse scattering) simultaneously	PDF analysis	Tucker <i>et al.</i> (2007)	http://www.rmcpfile.org	Free
<i>ROCKJOCK</i>	Quantitative analysis of mineralogical samples with or without using an internal standard by automatically fitting the sum of stored X-ray diffraction patterns of pure standard minerals to the measured pattern	Quantitative phase analysis, minerals	Eberl (2003)	http://pubs.er.usgs.gov/publication/ofr200378	Free
<i>Siroquant</i>	Comes with a comprehensive database of over 1800 phases. Phases from public sources such as the ICSD and AMCSD can also be added	Phase identification	Sietronics (2012)	http://www.siroquant.com	Commercial
<i>SPEC</i>	Software for the control of data acquisition. Includes <i>C-PLOT</i> for data display and analysis	Instrumentation	SPEC, Certified Scientific Software, PO Box 390640, Cambridge, MA, 02139-0007, USA	http://www.certif.com/	Commercial
<i>STEREOPOLE</i>	Analysis of X-ray diffraction pole figures, allowing experimental data to be compared graphically with simulated results, and epitaxial relationships between up to five different crystal layers to be evaluated	Texture analysis	Salzmann & Resel (2004)	http://www.if.tugraz.at/amd/stereopole/	Free
<i>StruVir</i>	Converts <i>STRUPLO</i> data files into Virtual Reality Modeling Language (VRML) for 3D visualization	Visualization	Le Bail (1996); Fischer (1985)	http://www.cristal.org/vrml/struvir.html	Free
<i>SUPERFLIP</i>	Structure solution from diffraction data using charge flipping. Can solve periodic structures, incommensurately modulated structures and the structures of quasicrystals from X-ray and neutron diffraction data	Structure solution	Palatinus & Chapuis (2007)	http://superflip.fzu.cz	Free

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>TOPAS</i> and <i>TOPAS-Academic</i>	Nonlinear least-squares optimization programs written in C++, allowing users to easily modify program functionality to suit the problem at hand	Peak fitting, Le Bail fitting, Pawley fitting, Rietveld refinement, PDF analysis, indexing, structure solution, charge flipping, magnetic structures, stacking-fault analysis	Coelho (2018)	https://www.bruker.com/products/x-ray-diffraction-and-elemental-analysis/x-ray-diffraction/xrd-software/topas.html , http://www.topas-academic.net	Free
<i>VALENCE</i>	Calculates bond valences from bond lengths and <i>vice versa</i> . Can also calculate bond-valence sums and average bond lengths, and can determine bond-valence parameters from the bonding environments of various cations	Structure validation	Brown (1996)	http://netlib.ccp14.ac.uk/ccp/web-mirrors/valence	Free
<i>VMRIA</i>	Full-profile analysis of neutron-diffraction time-of-flight data for multiphase samples. Includes <i>Autox</i>	Rietveld refinement, indexing	Zlokazov & Chernyshev (1992); Zlokazov (2003)	http://www.ccp14.ac.uk/ccp/web-mirrors/vmria/	Free
<i>WinPLOTR</i>	A graphical user interface for displaying powder patterns; part of <i>FULLPROF</i>	Powder-pattern viewing		http://www.cdifx.univ-rennes1.fr/winplotr/readme.htm	Free
<i>WinPSSP</i>	Uses direct-space methods to solve the crystal structures of small-molecule organic materials from X-ray powder diffraction data	Structure solution	Pagola <i>et al.</i> (2017)	http://users.uoi.gr/nkourkou/winpssp/	Free
<i>WinXPow</i>	Comprehensive package for data collection, profile fitting, indexing, and size-strain and crystallinity analysis	Phase identification, profile fitting, indexing, size-strain analysis, crystallinity	STOE & Cie GmbH, Hilpertstr. 10, D-64295 Darmstadt, Germany	https://www.stoe.com/product/software-powder-xrd/	Commercial
<i>XDrawChem</i>	Two-dimensional molecule drawing program for Unix operating systems	Model building	Herger (2002)	http://xdrawchem.sourceforge.net/	Free
<i>xINTERPDF</i>	Analysis of intermolecular pair-distribution functions of organic compounds from X-ray total-scattering data	PDF analysis	Shi (2018)	https://github.com/curieshicy/xINTERPDF	Free
<i>XLENS</i>	Uses Patterson-function direct methods instead of the modulus function. Well suited for handling powder diffraction data for organic compounds	Structure solution	Rius (2011, 2013)	https://departments.icmab.es/crystallography/software	Free for academic use

Table 6.1.1 Software for powder diffraction. Continue

Program	Description	Function(s)	Reference(s)	URL or other source	Availability
<i>xPDFsuite</i>	Suite of algorithms and software for pair-distribution-function analysis	PDF analysis	Juhás <i>et al.</i> (2013)	https://www.diffpy.org/products/xPDFsuite.html	Free for academic use
<i>X'Pert HighScore Plus</i>	Integrated powder-processing suite including profile fitting, structure solution, Rietveld refinement, quantitative analysis and cluster analysis	Cluster analysis, data conversion, indexing, Le Bail fitting, Pawley fitting, peak location, peak profiling, phase identification, powder-pattern viewing, quantitative phase analysis, Rietveld refinement, size-strain analysis	Malvern Panalytical BV, De Schakel 18, Kamer 50, Eindhoven, 5651 GH, Netherlands	https://www.malvernpanalytical.com/products/category/software/x-ray-diffraction-software/highscore-with-plus-option	Commercial
<i>Xp powder</i>	Identification, quantification and characterization of the crystalline components of solids, including natural minerals, artificial compounds and biological crystals	Phase identification	Quetzal Com S. L., La Carrera 5, 18110 HIJAR– Las Gabias, Granada, Spain	http://www.xpowder.com/	Commercial
<i>XRD2DScan</i>	Display and analysis of two-dimensional X-ray diffraction patterns collected with an area detector	2D data	Rodriguez-Navarro (2006)	http://www.ugr.es/~anava/xrd2dscan.htm	Free
<i>XRDU A</i>	Reconstruction of crystalline phase distribution maps from two-dimensional diffraction patterns collected by scanning and tomographic X-ray powder diffraction	Data processing, phase distributions	de Nolf (2014)	http://xrdua.ua.ac.be	Free
<i>ZEFS A II</i>	Zeolite structure solution using a biased Monte Carlo scheme	Structure solution, zeolites	Falcioni & Deem (1999)	http://www.mwdeem.rice.edu/zefsaII/	Free