

Solving and refining crystal structures using symmetry modes: FullProf + AMPLIMODES.

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- Overview of the *Symmetry Analysis* in phase transitions
- How is implemented the use of symmetry modes in *FullProf*
- Detailed examples:
Jahn-Teller transition in LaMnO_3

Symmetry and Phase Transitions

In a displacive phase transition the symmetry-breaking distortion (with respect to the high symmetry phase) is mainly caused by the freezing of the *primary mode*, associated with the order parameter.

In general, *secondary modes* are also triggered at the transition and can have non-zero amplitudes in the distorted structure.

The symmetry-mode analysis of a structural phase transition consists on the calculation of the amplitudes of the *symmetry modes* frozen in the distortion characterized by the *eigenvectors* of both primary and secondary modes present in the distortion.

Symmetry and Phase Transitions

Modes are collective correlated atomic displacements fulfilling certain symmetry properties. Structural distortions can be decomposed into contributions from different modes with symmetries given by irreducible representations of the parent space group.

In general, the use of symmetry-adapted modes in the description of distorted structures introduces a natural physical hierarchy among the structural parameters. This can be useful not only for investigating the physical mechanisms that stabilize these phases, but also for pure crystallographic purposes.

Mode Crystallography

The team of the Bilbao Crystallographic Server has developed the computer program: **AMPLIMODES**, that allows an easy calculation of the decomposition in modes of a distorted crystal structure with respect to a (virtual) high symmetry structure.

The originality of this approach with respect to more classical ones (e.g. BasIreps, MODY, Sarah, ...) is that **the polarization vectors are referred to the basis of the low symmetry phase**, allowing to use conventional crystallographic approaches (asymmetric unit and space group operators) to the crystal structure analysis.

AMPLIMODES: Symmetry mode analysis on the Bilbao Crystallographic Server, D. Orobengoa, C.Capillas, M.I. Aroyo and J.M. Perez-Mato, JApplCryst. (in press)

Distorted structures in terms of modes

Let $\mathbf{r}(\mu)$ be the positions of the atoms μ ($\mu = 1, \dots, s$) within an asymmetric unit of the parent structure with space group \mathbf{H} . The asymmetric unit of the observed distorted structure with lower space group \mathbf{L} , subgroup of \mathbf{H} , will in general have a larger number of atoms due to the splitting of the Wyckoff orbits in \mathbf{H} .

$$\mathbf{r}(\mu, i) = \mathbf{r}_0(\mu, i) + \mathbf{u}(\mu, i) \quad \mu = 1, 2, \dots, s, \quad i = 1, 2, \dots, n_\mu$$

$$\mathbf{u}(\mu, i) = \sum_{\tau, m} A_{\tau, m} \boldsymbol{\varepsilon}(\tau, m | \mu, i)$$

The indices τ and m label all possible distinct allowed symmetry-adapted distortion modes. τ stands for the possible different mode symmetries, while m ($m = 1, \dots, n_\tau$) enumerates the possible different independent modes of a given symmetry.

Distorted structures in terms of modes

$$\mathbf{u}(\mu, i) = \sum_{\tau, m} A_{\tau, m} \boldsymbol{\varepsilon}(\tau, m | \mu, i) \quad \mu = 1, 2, \dots, s, \quad i = 1, 2, \dots, n_{\mu}$$

The mode (τ, m) is defined by the polarisation vectors:

$$\boldsymbol{\varepsilon}(\tau, m | \mu, i)$$

One can refer to the global polarization vector $\boldsymbol{\varepsilon}(\tau, m)$, taking all atoms simultaneously, of the mode (τ, m)

The displacements of an atom (μ', i') related by the symmetry operator $\{\mathbf{R}|\mathbf{t}\}$ to the atom (μ, i) are given directly by:

$$\mathbf{R} \boldsymbol{\varepsilon}(\tau, m | \mu, i)$$

Distorted structures in terms of modes

The normalization of the polarisation vectors is chosen to verify:

$$\sum_{\mu,i} mult_{\mu,i} |\boldsymbol{\varepsilon}(\tau, m | \mu, i)|^2 = 1$$

" $mult_{\mu,i}$ " represents the multiplicity in a primitive cell of the space group L for the Wyckoff position (μ, i) .

The following orthogonality relation is verified by the polarization vectors:

$$\sum_{\mu,i} mult_{\mu,i} \boldsymbol{\varepsilon}(\tau, m | \mu, i) \boldsymbol{\varepsilon}(\tau', m' | \mu, i) = \delta_{\tau\tau'} \delta_{mm'}$$

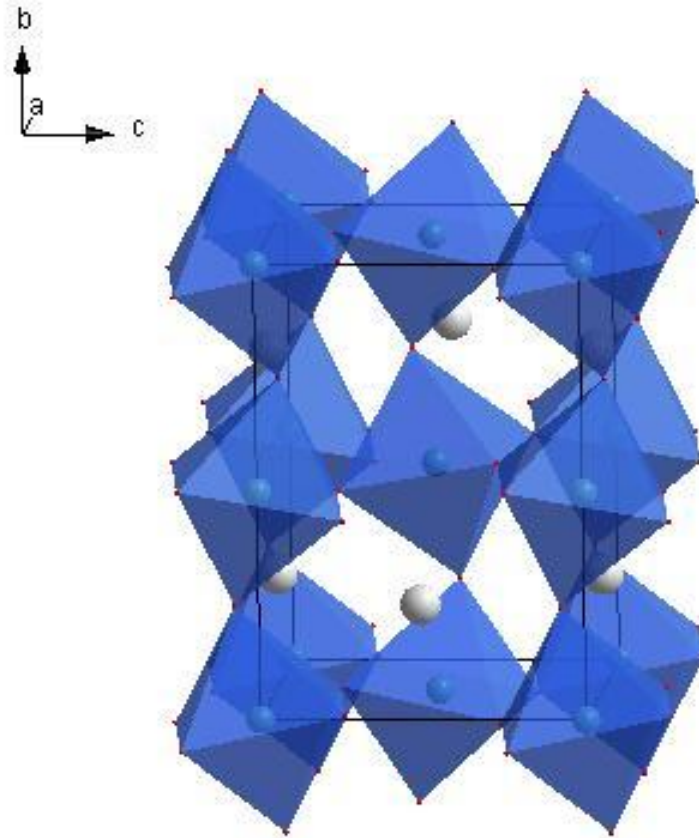
Distorted structures in terms of modes

The distortion modes of the phase with group H having **isotropy group** equal to L can be called **primary**, while those with **isotropy groups** given by **subgroups** of H which are distinct **supergroups** of L , are usually termed **secondary**.

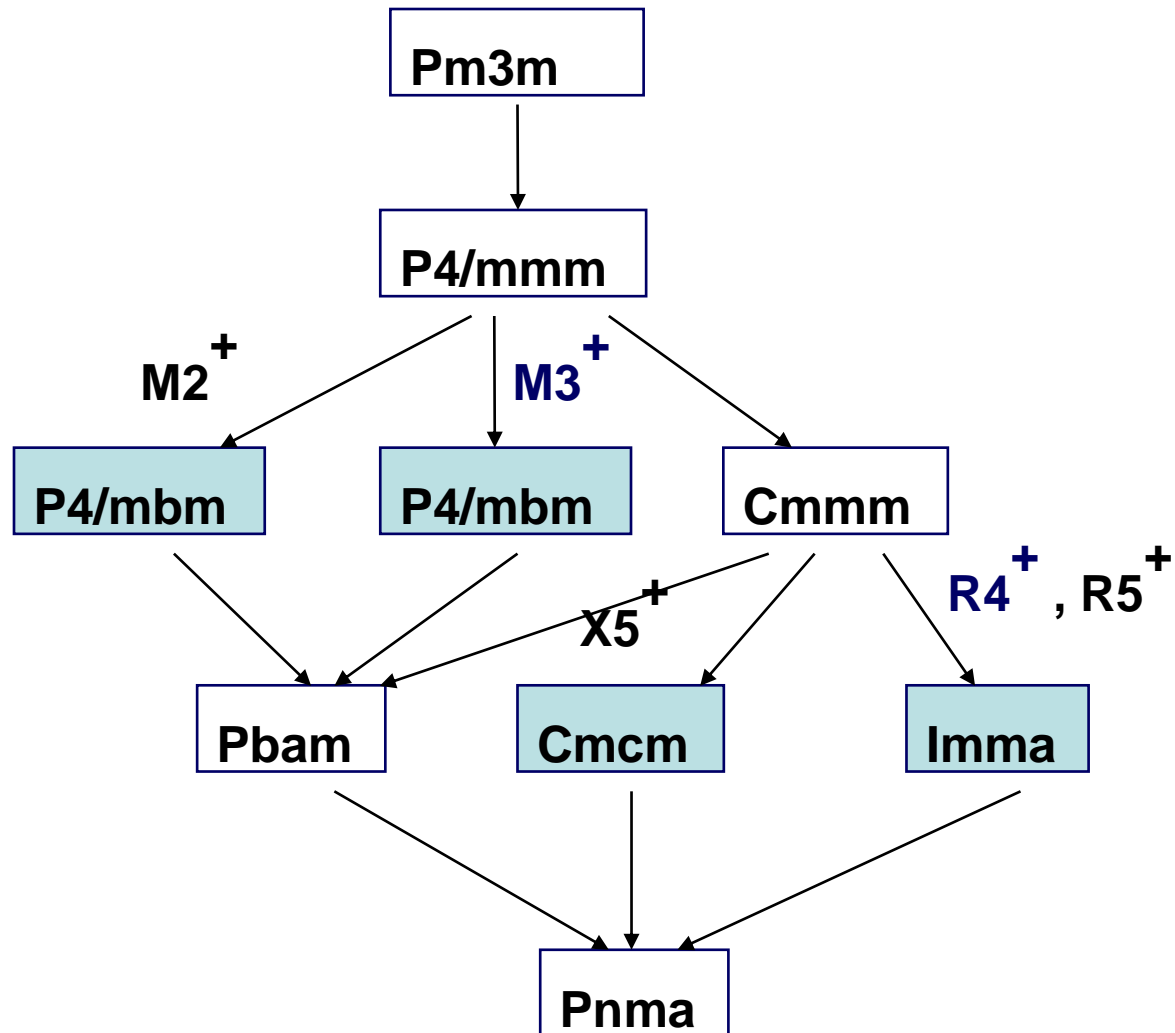
A **primary distortion mode** is sufficient to produce the observed symmetry breaking between the **parent** and the **observed** structure, while **secondary distortion modes** alone would yield a **higher symmetry**.

Distorted perovskite: structure type $GdFeO_3$

Space group: $Pnma$, parent structure $Pm3m$



Group-subgroup chains relating $Pm3m$ and $Pnma$



Distorted structures in terms of modes

It is also in general very convenient to express the global distortion in terms of the **different symmetry components** (this is done in AMPLIMODES):

$$\mathbf{u}(\mu, i) = \sum_{\tau, m} A_{\tau, m} \boldsymbol{\varepsilon}(\tau, m | \mu, i) = \sum_{\tau} A_{\tau} \mathbf{e}(\tau | \mu, i)$$

$$A_{\tau} = \left(\sum_m A_{\tau, m}^2 \right)^{1/2}$$

$$\mathbf{e}(\tau | \mu, i) = \sum_m a_{\tau, m} \boldsymbol{\varepsilon}(\tau, m | \mu, i); \quad a_{\tau, m} = \frac{A_{\tau, m}}{\left(\sum_m A_{\tau, m}^2 \right)^{1/2}}$$

- Overview of the *Symmetry Analysis* in phase transitions
- How is implemented the use of symmetry modes in *FullProf*
- Detailed examples:
Jahn-Teller transition in LaMnO_3

Refinement of crystal structures using amplitudes of symmetry modes instead of atom positions in *FullProf*

In **FullProf** the refinement of a crystal structure can be done in terms of symmetry adapted modes.

<http://www.ill.eu/sites/fullprof/>

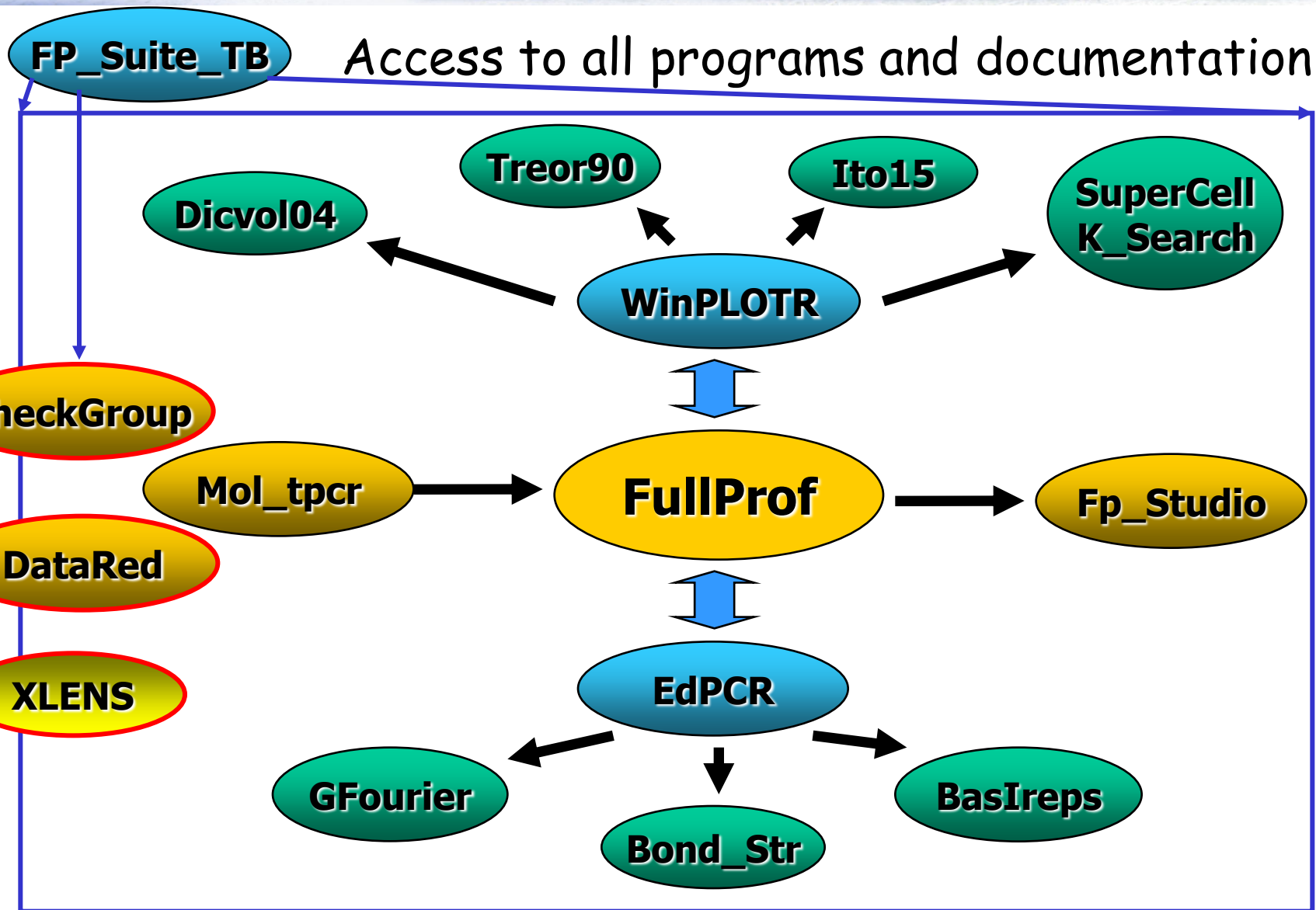
FullProf uses the polarisation vectors obtained from the output of the program **AMPLIMODES** from the Bilbao Crystallographic Server

<http://www.cryst.ehu.es/cryst/amplimodes.html>

A low symmetry (LS) crystal structure (Space Group L) is supposed to derive (from a phase transition) from a high symmetry (HS) structure (Space Group H) with $L \subset H$.

The free parameters, instead of atom positions, are the amplitudes of a combination of allowed symmetry modes.

FullProf Suite



Magnetic structures

Magnetic moment of each atom: Fourier series

$$\mathbf{m}_{ljs} = \sum_{\mathbf{k}} \mathbf{S}_{\mathbf{k}js} \exp \left\{ 2\pi i \mathbf{k} \mathbf{R}_l \right\}$$

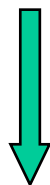
The program **Fp_Studio** performs the above sum and represents graphically the magnetic structure. This program can help to learn about this formalism because the user can write manually the Fourier coefficients and see what is the corresponding magnetic structure immediately.

Web site: <http://www.ill.eu/sites/fullprof/>

Group Theory: Symmetry Analysis

Fourier coefficients as linear combinations of the basis functions of the irreducible representation of the propagation vector group $G_{\mathbf{k}}$

$$\mathbf{S}_{\mathbf{k}js} = \sum_{n\lambda} C_{n\lambda}^{\nu} \mathbf{S}_{n\lambda}^{\mathbf{k}\nu} \quad js$$



$$\mathbf{M}(\mathbf{h}) = p \sum_{j=1}^n O_j f_j(\mathbf{h}) T_j \sum_{n\lambda} C_{n\lambda}^{\nu} \sum_s \mathbf{S}_{n\lambda}^{\mathbf{k}\nu} \quad js \exp 2\pi i \mathbf{h}_s \mathbf{r}_j$$

Competing multi-q magnetic structures in HoGe_3 (I & II)

P Schöbinger-Papamantellos, J Rodríguez-Carvajal, LD Tung, C Ritter and KHJ Buschow

J. Physics: Condensed Matter **20** (2008) 195201 (12pp)
195202(13pp)

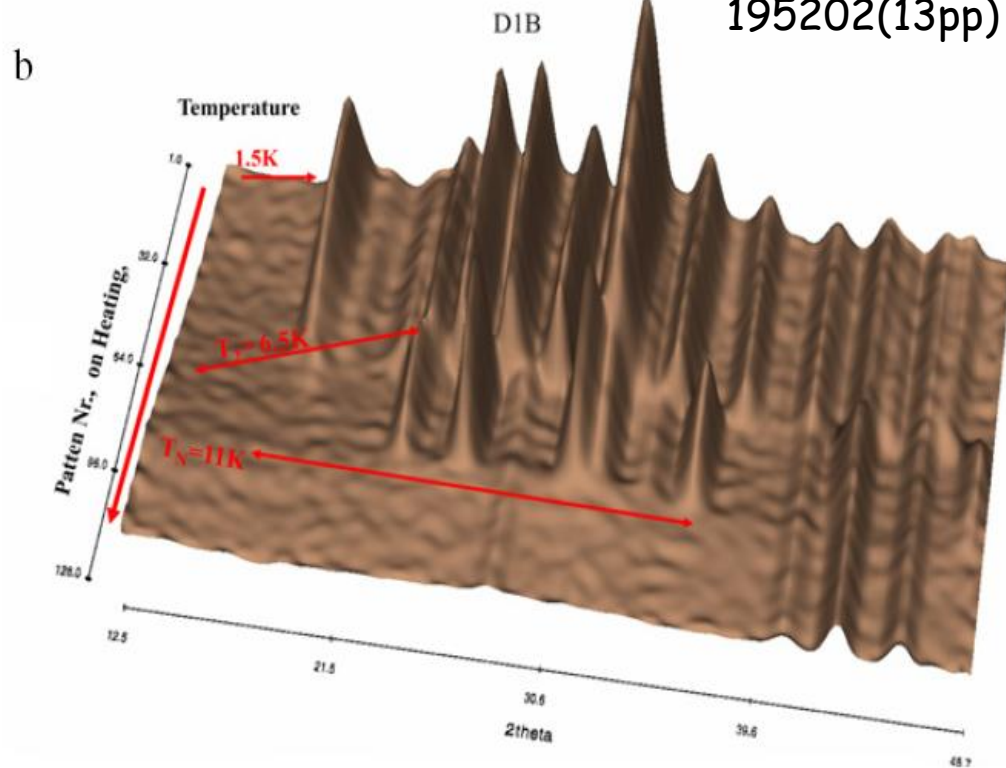
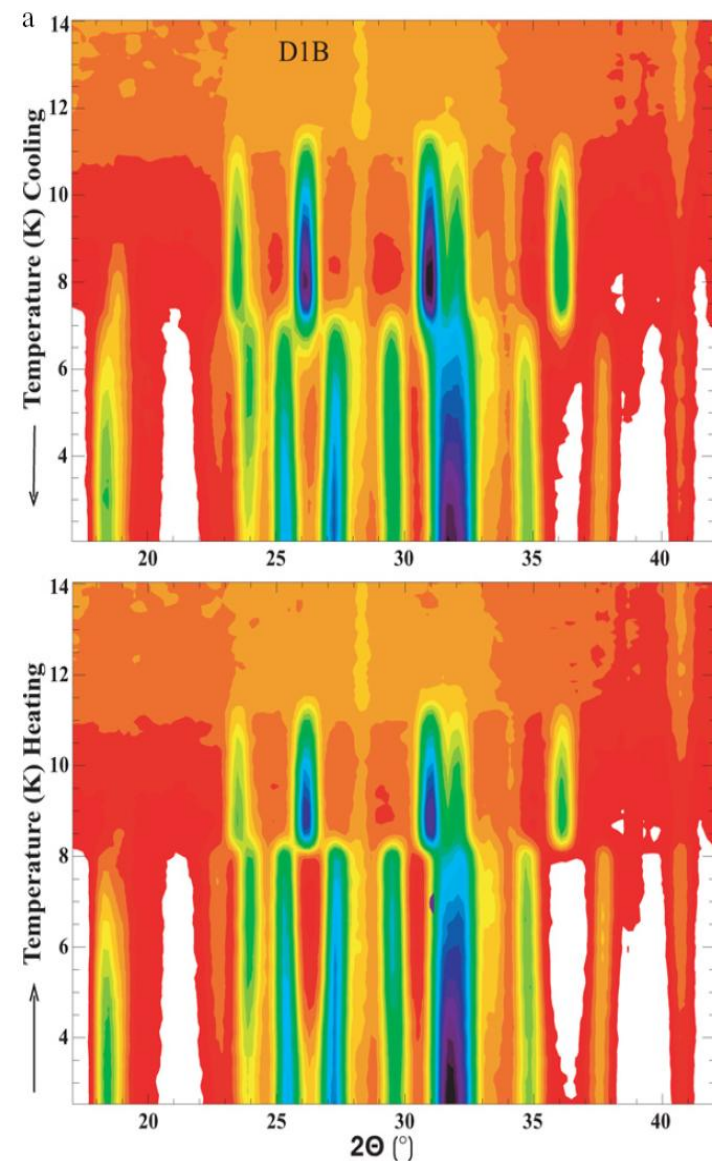


Figure 6. Thermodiffractogram of HoGe_3 : (a) in a 2D projection on heating and cooling showing the succession of magnetic phase transitions below $T_N = 11$ K at $T_2^H = 8.1$ K and $T_3^H = 4.8$ K (temperatures given on heating) and (b) in a 3D view on cooling.

The atoms position are calculated from the following formula:

$$\mathbf{r}_j^{LS} = \mathbf{r}_j^{HS} + \sum_m c_m Q_m \boldsymbol{\varepsilon}(m | j)$$

Where j runs over the atoms in the asymmetric unit of the LS phase
 The index m runs over all contributing modes. It may content modes corresponding to different representations and wave vectors of the H space group (Isotropy subgroups) that are compatible with the L space group.

The polarisation vectors $\boldsymbol{\varepsilon}(m | j)$ have normalized components referred to the conventional cell of the LS phase and are provided by **AMPLIMODES**. The refined parameters are the amplitudes Q_m . c_m are normalisation coefficients.

A representation of the modes using arrows and the HS phase can be visualised using **FullProf Studio**

Example of PCR file for FullProf corresponding to the compound LaMnO_3

LaMnO3

!

Symmetry modes option

!Nat Dis Ang Pr1 Pr2 Pr3 **Jbt** Irf Isy Str Furth ATZ Nvk Npr More
4 0 0 0.0 0.0 1.0 6 0 0 0 7 967.370 0 7 1

P b n m <--Space group symbol

!Atom	Typ	X	Y	Z	Biso	Occ
La	LA	0.00000	0.50000	0.25000	0.35050	0.50000
		0.00	0.00	0.00	251.00	0.00
Mn	MN	0.00000	0.00000	0.00000	0.21228	0.50000
		0.00	0.00	0.00	261.00	0.00
O1	O	0.75000	0.25000	0.00000	0.43965	1.00000
		0.00	0.00	0.00	271.00	0.00
O2	O	0.50000	0.50000	0.75000	0.50234	0.50000
		0.00	0.00	0.00	281.00	0.00

Normalisation coefficients

keyword

Number of polarisation vector modes

! Polarisation vectors of symmetry modes for each atom

V_MODES	12					
! Nm	Atm	Irrep	Vx	Vy	Vz	Coeff
1	O1	R4+	0.000000	0.000000	0.031721	1.000000
1	O2	R4+	0.063442	0.000000	0.000000	1.000000
2	La	R5+	-0.089721	0.000000	0.000000	1.000000
3	O1	R5+	0.000000	0.000000	-0.031721	1.000000
...
7	O2	M3+	0.000000	0.000000	0.000000	1.000000

Indices of the modes

! Amplitudes of Symmetry Modes

Example of PCR file for *FullProf* corresponding to the compound LaMnO_3

Indices of the modes

Symbols of the Irreducible representations

```

! Polarisation Vectors of Symmetry Modes for each atom
V_MODES      12
! Nm Atm      Irrep      Vx      Vy      Vz      Coeff
1   O1        R4+        0.000000  0.000000  0.031721  1.000000
1   O2        R4+        0.063442  0.000000  0.000000  1.000000
2   La        R5+       -0.089721  0.000000  0.000000  1.000000
3   O1        R5+        0.000000  0.000000 -0.031721  1.000000
.   .   .   .   .   .   .   .
7   O2        M3+        0.000000  0.000000  0.000000  1.000000
! Amplitudes of Symmetry Modes
A_MODES      7      1 1 1 1 1 1 1
Q1_R4+      -1.189680  181.0000
Q2_R5+      -0.086467  191.0000
Q3_R5+       0.018171  201.0000
Q4_X5+     -0.546082  211.0000
Q5_X5+     -0.139910  221.0000
Q6_M2+      0.355652  231.0000
Q7_M3+      0.901264  241.0000
!-----> Profile Parameters for Pattern # 1
!  Scale      Shape1      Bov      Str1      Str2      Str3      Strain-Model
0.86919E-01  0.00000  0.00000  0.0000  0.0000  0.0000  0
. . . . .

```

Polarisation vectors components

Keyword, # of modes, output for FST

Names of amplitudes, values and refinement codes (allowing constraints)

- Overview of the *Symmetry Analysis* in phase transitions
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- Detailed example:
Jahn-Teller transition in LaMnO_3

Example: The Jahn-Teller transition in LaMnO_3

RAPID COMMUNICATIONS

PHYSICAL REVIEW B

VOLUME 57, NUMBER 6

1 FEBRUARY 1998-II

Neutron-diffraction study of the Jahn-Teller transition in stoichiometric LaMnO_3

J. Rodríguez-Carvajal,* M. Hennion, F. Moussa, and A. H. Moudden

Laboratoire Léon Brillouin (CEA-CNRS), Centre d'Etudes de Saclay, 91191 Gif sur Yvette Cedex, France

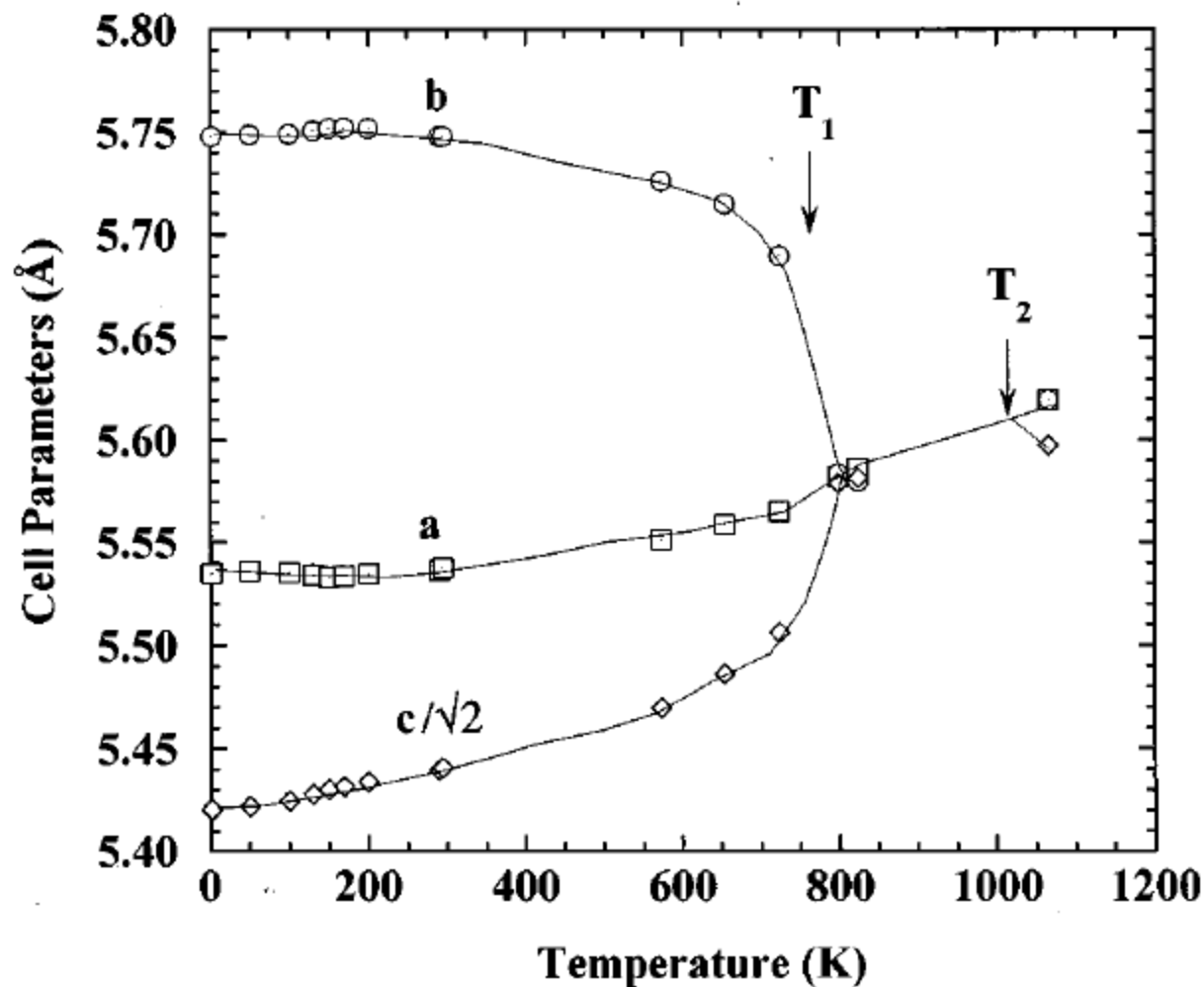
L. Pinsard and A. Revcolevschi

Laboratoire de Chimie des Solides, Université Paris Sud, 91405 Orsay Cedex, France

(Received 2 September 1997)

The parent compound of the giant magnetoresistance Mn-perovskite, LaMnO_3 , has been studied by thermal analysis and high-resolution neutron-powder diffraction. The orthorhombic $Pbnm$ structure at room temperature is characterized by an antiferrodistorsive orbital ordering due to the Jahn-Teller effect. This ordering is evidenced by the spatial distribution of the observed Mn-O bond lengths. LaMnO_3 undergoes a structural phase transition at $T_{JT} \approx 750$ K, above which the orbital ordering disappears. There is no change in symmetry although the lattice becomes metrically cubic on the high-temperature side. The MnO_6 octahedra become nearly regular above T_{JT} and the thermal parameter of oxygen atoms increases significantly. The observed average cubic lattice is probably the result of dynamic spatial fluctuations of the underlying orthorhombic distortion. [S0163-1829(98)51706-7]

Example: The Jahn-Teller transition in LaMnO_3



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bilbao crystallographic server

[The crystallographic site at the Condensed Matter Physics Dept. of the University of the Basque Country]

[Space Groups] [Layer Groups] [Rod Groups] [Frieze Groups] [Wyckoff Sets]



CrystallographyOnline:
International School on
the Use and Applications
of the Bilbao
Crystallographic
Server

REGISTRATION OPEN

Space Groups Retrieval Tools

GENPOS	Generators and General Positions of Space Groups
WYCKPOS	Wyckoff Positions of Space Groups
HKLCOND	Reflection conditions of Space Groups
MAXSUB	Maximal Subgroups of Space Groups
SERIES	Series of Maximal Isomorphic Subgroups of Space Groups
WYCKSETS	Equivalent Sets of Wyckoff Positions
NORMALIZER	Normalizers of Space Groups
KVEC	The k-vector types and Brillouin zones of Space Groups

PhD Studentship

Sections

- [Retrieval Tools](#)
- [Group-Subgroup](#)
- [Representations](#)
- [Solid State](#)
- [Structure Utilities](#)
- [Subperiodic](#)
- [ICDD](#)

Group - Subgroup Relations of Space Groups

SUBGROUPGRAPH	Lattice of Maximal Subgroups
HERMANN	Distribution of subgroups in conjugated classes
COSETS	Coset decomposition for a group-subgroup pair
WYCKSPLIT	The splitting of the Wyckoff Positions
MINSUP	Minimal Supergroups of Space Groups
SUPERGROUPS	Supergroups of Space Groups

Done

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modes.

- **TRANSTRU**
5-2008: Transform structures to lower symmetry Space Group basis.
- **AMPLIMODES**
2-2008: Symmetry Mode Analysis of Structural Phase Transitions.
- **CELLSUPER**
10-2007: New version of program CELLSUPER.
- **TRANPATH**
7-2007: Minor update and fixes.
- **SUPERGROUPS**
6-2007: Added link to Wyckoff Positions splitting.
- **SERIES**
1-2007: New version of series of maximal isomorphic subgroups for a given maximum index.
- **SIMPLE RETRIEVAL TOOLS**
1-2007: Minor upgrade for GENPOS, WYCKPOS, MAXSUB and SERIES programs.
- **HERMANN**
1-2007: New version of program HERMANN.
- **SETSTRU**
1-2007: CIF input data, JMOL visualization and minor bugs fixed

POINT	Point Group Tables
SITESYM	Site-symmetry induced representations of Space Groups

Solid State Theory Applications

SAM	Spectral Active Modes (IR and RAMAN Selection Rules)
NEUTRON	Neutron Scattering Selection Rules
SYMMODES	Primary and Secondary Modes for a Group - Subgroup pair
AMPLIMODES	Symmetry Mode Analysis
PSEUDO	Pseudosymmetry Search in a Structure
DOPE	Degree of Pseudosymmetry Estimation
BPLOT	Pseudosymmetry Search with KPLOT
TRANPATH	Transition Paths (Group not subgroup relations)

Structure Utilities

CELLTRAN	Transform Unit Cells
STRAIN	Strain Tensor Calculation
WPASSIGN	Assignment of Wyckoff Positions
TRANSTRU	Transform structures to lower symmetry Space Group basis.
SETSTRU	Alternative Settings for a given Crystal Structure
EQUIVSTRU	Equivalent Descriptions for a given Crystal Structure

Subperiodic Groups: Layer, Rod and Frieze Groups Retrieval Tools

http://www.cryst.ehu.es/cryst/amplimodes.html

AMPLIMODES for FullProf - Mozilla Firefox

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http://www.cryst.ehu.es/html/cryst/full-modes.html

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Symmetry modes for FullProf

AMPLIMODES carries out a symmetry-mode analysis of a displacive phase transition. Starting from the experimental structures of the high- and low symmetry phases, the program determines the global structural distortion that relates the two phases. The symmetry modes compatible with the symmetry break are then calculated. Their orthogonality permits the decomposition of the global distortion obtaining the amplitude of each symmetry mode as well as the corresponding eigenvectors.

The input of the program consists of:

- The information about the structures of the high- and low symmetry phases: Space group number, lattice parameters and relative atomic coordinates of the asymmetric unit.
- The transformation matrix that relates the basis of the two space groups.

Structure data

HINT: [Upload the structure as a CIF file (default), or as a text in the window below]

```
# Space Group ITA number
221
# Lattice parameters
3.8 3.8 3.8 90 90 90
# Number of independent atoms in the asymmetric unit
3
# [atom type] [number] [WP] [x] [y] [z]
Ca 1 1b 0.5 0.5 0.5
Ti 1 1a 0.0 0.0 0.0
O 1 3d 0.5 0.0 0.0
```

High Symmetry Structure

For the low symmetry structure, only the space group and the lattice parameters are necessary, the structure is optional, if given, the program will return the amplitudes of the modes.

HINT: [Upload the structure as a CIF file (default), or as a text in the window below]

```
# Space Group ITA number
62
# Lattice parameters
5.440791 7.644515 5.379452 90 90 90
# Number of independent atoms in the asymmetric unit
4
# [atom type] [number] [WP] [x] [y] [z]
Ca 1 4c 0.03539 0.25000 -0.00483
Ti 1 4b 0.50000 0.00000 0.00000
O 1 4c 0.48475 0.25000 0.07094
O 2 8d 0.28945 0.03690 0.70986
```

Low Symmetry Structure

Maximum Δ [Maximum distance allowed]

Transformation

Done

AMPLIMODES for FullProf - Mozilla Firefox

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http://www.cryst.ehu.es/html/cryst/full-modes.html

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corresponding eigenvectors.

The input of the program consists of:

- The information about the structures of the high- and low symmetry phases: Space group number, lattice parameters and relative atomic coordinates of the asymmetric unit.
- The transformation matrix that relates the basis of the two space groups.

For the low symmetry structure, only the space group and the lattice parameters are necessary, the structure is optional, if given, the program will return the amplitudes of the modes.

HINT: [Upload the structure as a CIF file (default), or as a text in the window below]

```
# Space Group ITA number
62
# Lattice parameters
5.374012 7.600000 5.374012 90.0 90.0 90.0
```

Low Symmetry Structure

Maximum Δ [Maximum distance allowed]

Transformation Matrix

In matrix form:

	Rotational part			Origin Shift
<input type="text" value="1"/>	<input type="text" value="0"/>	<input type="text" value="-1"/>	<input type="text" value="0.5"/>	
<input type="text" value="0"/>	<input type="text" value="2"/>	<input type="text" value="0"/>	<input type="text" value="0"/>	
<input type="text" value="1"/>	<input type="text" value="0"/>	<input type="text" value="1"/>	<input type="text" value="0.5"/>	

[Bilbao Crystallographic Server Main Menu]

Done

Symmetry modes for FullProf - Mozilla Firefox

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http://www.cryst.ehu.es/cgi-bin/cryst/programs/nph-full-modes

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Transformed high symmetry structure in the subgroup basis

```

062
5.374012 7.600000 5.374012 90.000000 90.000000 90.000000
4
Ca 1 4c 0.000000 0.250000 0.000000
Ti 1 4b 0.500000 0.000000 0.000000
O 1 8d 0.750000 0.000000 0.750000
O 1_2 4c 0.500000 0.250000 0.000000
  
```

Symmetry Modes Summary

Atoms	WP	Modes	Show Modes
O1	3d	R4+(1) R5+(1) X5+(1) M2+(1) M3+(1)	<input type="button" value="Show"/>
Ca1	1b	R5+(1) X5+(1)	<input type="button" value="Show"/>

You can copy and paste the following text on your .pcr file

```

062
!Atom Typ X Y Z Basis Occ In Fin N_t Spc /Codes
Ca1 CA 0.000000 0.250000 0.000000 0.500000 0.500000 0 0 0 1
0.00 0.00 0.00 0.00 0.00
Ti1 TI 0.500000 0.000000 0.000000 0.500000 0.500000 0 0 0 1
0.00 0.00 0.00 0.00 0.00
O1 O 0.750000 0.000000 0.750000 0.500000 1.000000 0 0 0 1
0.00 0.00 0.00 0.00 0.00
O1_2 O 0.500000 0.250000 0.000000 0.500000 0.500000 0 0 0 1
0.00 0.00 0.00 0.00 0.00
! Polarisation Vectors of Symmetry Modes for each atom
V_MODES 12
! Nm Atom Irrep Vx Vy Vz Coeff
1 O1 R4+ 0.000000 0.032895 -0.000000 1.00
1 O1_2 R4+ 0.000000 0.000000 0.065789 1.00
2 Ca1 R5+ 0.000000 0.000000 -0.093040 1.00
3 O1 R5+ 0.000000 -0.032895 0.000000 1.00
3 O1_2 R5+ 0.000000 0.000000 0.065789 1.00
4 Ca1 X5+ -0.093040 0.000000 0.000000 1.00
5 O1 X5+ 0.000000 0.000000 0.000000 1.00
5 O1_2 X5+ 0.093040 -0.000000 -0.000000 1.00
6 O1 M2+ -0.046520 0.000000 0.046520 1.00
6 O1_2 M2+ 0.000000 0.000000 0.000000 1.00
7 O1 M3+ -0.046520 0.000000 -0.046520 1.00
  
```

Done

Example of *FullProf* running a refinement of LaMnO3 using symmetry modes

```

FullProf Program
Load Edit PCR Mode Run Exit
=> Bragg R-factor: 4.651
=> RF-factor : 3.019

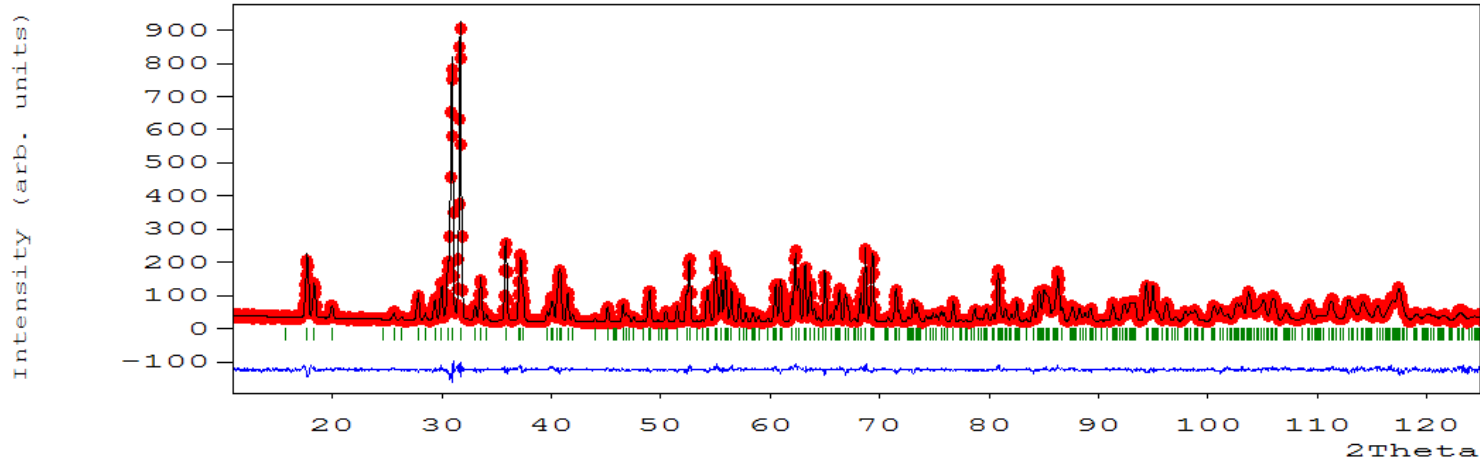
=> Convergence reached at this CYCLE !!!!: CYCLE No. 15
=> R-Factors: 4.94 5.79 Chi2: 2.11 DW-Stat.: 1.0292 Patt#: 1
=> Expected : 3.98 1.8768
=> Conventional Rietveld R-factors for Pattern: 1
=> Rp: 9.07 Rwp: 8.89 Rexp: 6.12 Chi2: 2.11
=> Global user-weighted Chi2 (Bragg contrib.): 2.150
=> -----> Pattern# 1
=> Phase: 1
=> Bragg R-factor: 4.651
=> RF-factor : 3.019
=> Normal end, final calculations and writing...

=> CPU Time: 9.453 seconds
=> 0.158 minutes

=> END Date:19/08/2008 Time => 01:44:25.133
  
```

Cycle: 15

lamn_3t2.dat



```

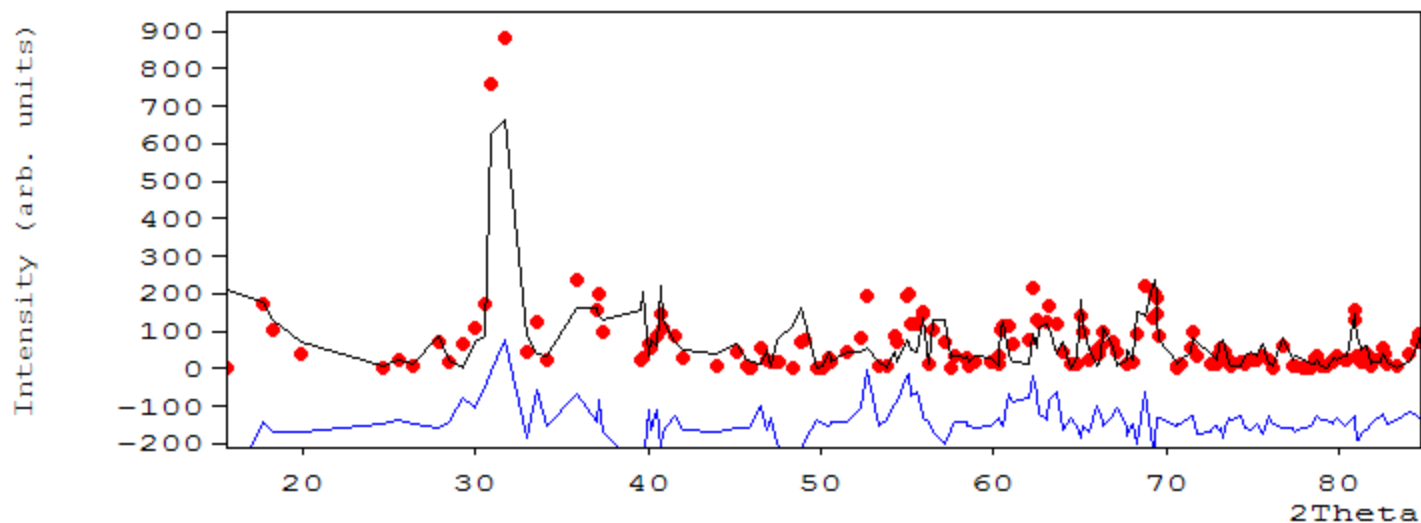
=>      Rietveld, Profile Matching & Integrated Intensity
=>      Refinement of X-ray and/or Neutron Data
=>      (Multi_Pattern: Windows-version)

=> START Date:20/09/2008  Time => 15:37:17.640
=> Reading control file *.PCR ...
=> End of preliminary calculations !

=> **** SIMULATED ANNEALING SEARCH FOR STARTING CONFIGURATION ****

=> Initial configuration cost:  107.15
=> Initial configuration state vector:
=>   Q1_R4+   Q2_R5+   Q3_R5+   Q4_X5+   Q5_X5+   Q6_M2+   Q7_M3+
=>         1         2         3         4         5         6         7
=>   0.0000   0.0000   0.0000   0.0000   0.0000   0.0000   0.0000
=> NT:  1 Temp:  6.00 (%Acc): 42.04 <Step>:  4.0000 <Rp-factor>: 85.4356
  
```

Function evaluations: 491 lamn_san.int



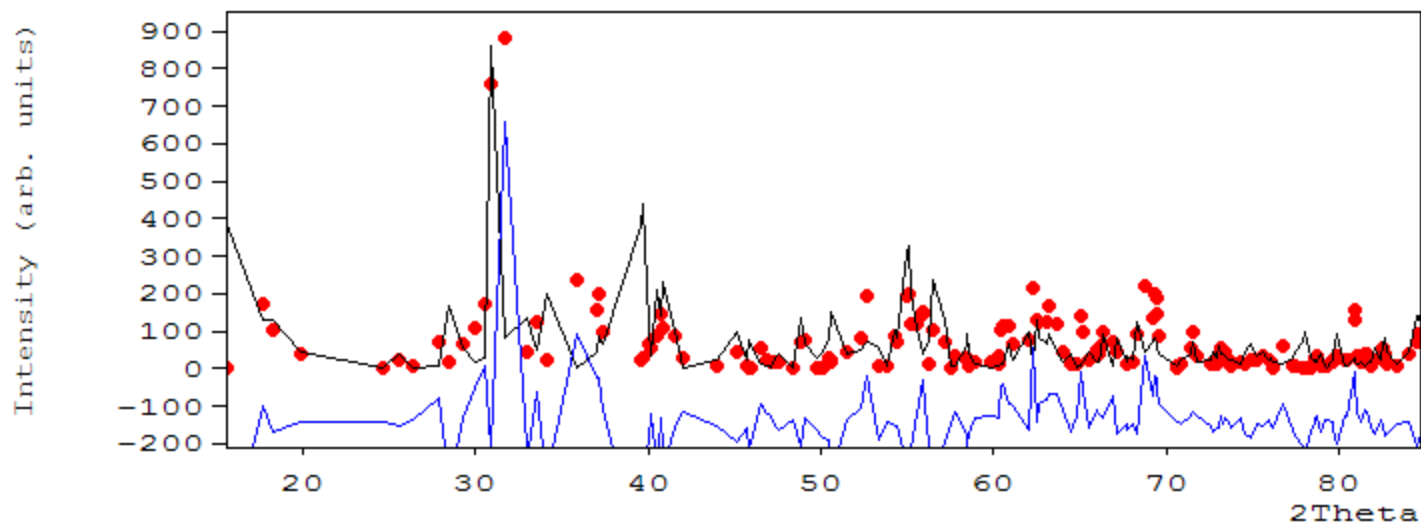
```

=> START Date:20/09/2008 Time => 15:33:20.281
=> Reading control file *.PCR ...
=> End of preliminary calculations !

=> **** SIMULATED ANNEALING SEARCH FOR STARTING CONFIGURATION ****

=> Initial configuration cost: 107.15
=> Initial configuration state vector:
=>   Q1_R4+   Q2_R5+   Q3_R5+   Q4_X5+   Q5_X5+   Q6_M2+   Q7_M3+
=>         1         2         3         4         5         6         7
=>   0.0000   0.0000   0.0000   0.0000   0.0000   0.0000   0.0000
=> NT:  1 Temp:  6.00 (%Acc): 42.04 <Step>:  4.0000 <Rp-factor>: 85.4356
=> NT:  2 Temp:  5.70 (%Acc): 56.12 <Step>:  3.7571 <Rp-factor>: 86.7607
=> NT:  3 Temp:  5.41 (%Acc): 24.49 <Step>:  3.8286 <Rp-factor>: 65.8903
=> NT:  4 Temp:  5.14 (%Acc): 32.04 <Step>:  2.2809 <Rp-factor>: 56.3059
  
```

Function evaluations: 1961 lamn_san.int



=> **** SIMULATED ANNEALING SEARCH FOR STARTING CONFIGURATION ****

=> Initial configuration cost: 107.15

=> Initial configuration state vector:

	Q1_R4+	Q2_R5+	Q3_R5+	Q4_X5+	Q5_X5+	Q6_M2+	Q7_M3+
=>	1	2	3	4	5	6	7
=>	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000

=> NT: 1 Temp: 6.00 (%Acc): 42.04 <Step>: 4.0000 <Rp-factor>: 85.4356

=> NT: 2 Temp: 5.70 (%Acc): 56.12 <Step>: 3.7571 <Rp-factor>: 86.7607

=> NT: 3 Temp: 5.41 (%Acc): 24.49 <Step>: 3.8286 <Rp-factor>: 65.8903

=> NT: 4 Temp: 5.14 (%Acc): 32.04 <Step>: 2.2809 <Rp-factor>: 56.3059

=> NT: 5 Temp: 4.89 (%Acc): 31.02 <Step>: 1.6271 <Rp-factor>: 49.8507

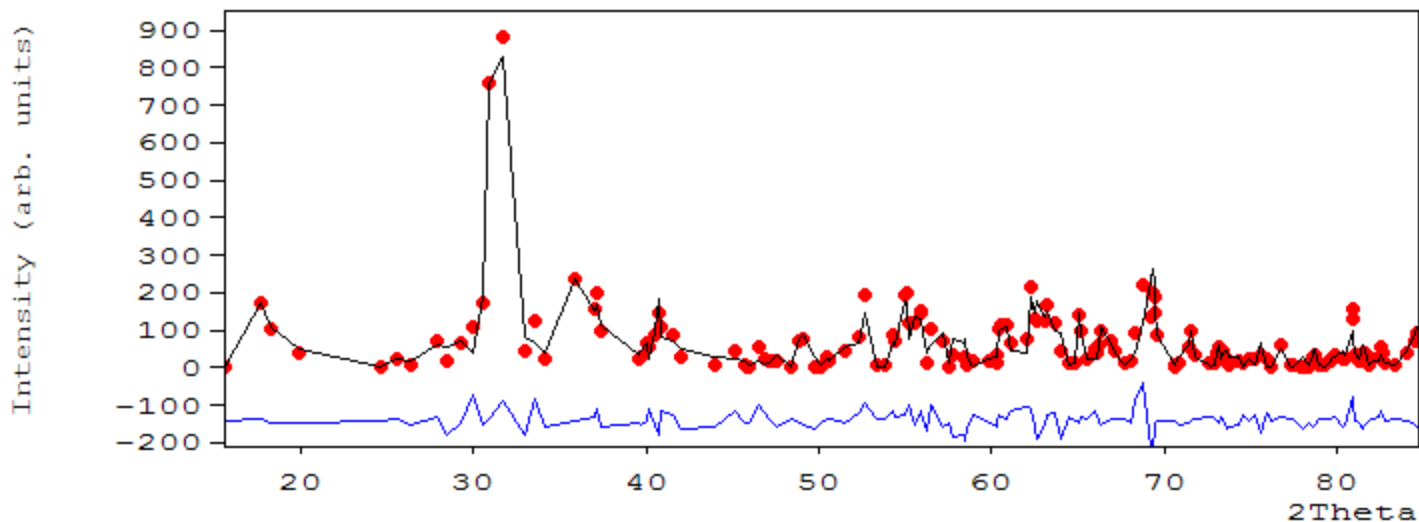
=> NT: 6 Temp: 4.64 (%Acc): 47.76 <Step>: 1.1245 <Rp-factor>: 48.1753

=> NT: 7 Temp: 4.41 (%Acc): 34.29 <Step>: 1.1456 <Rp-factor>: 33.1074

=> NT: 8 Temp: 4.19 (%Acc): 42.24 <Step>: 0.8090 <Rp-factor>: 35.1309

Function evaluations: 3921

lamn_san.int



=> **** SIMULATED ANNEALING SEARCH FOR STARTING CONFIGURATION ****

=> Initial configuration cost: 107.15

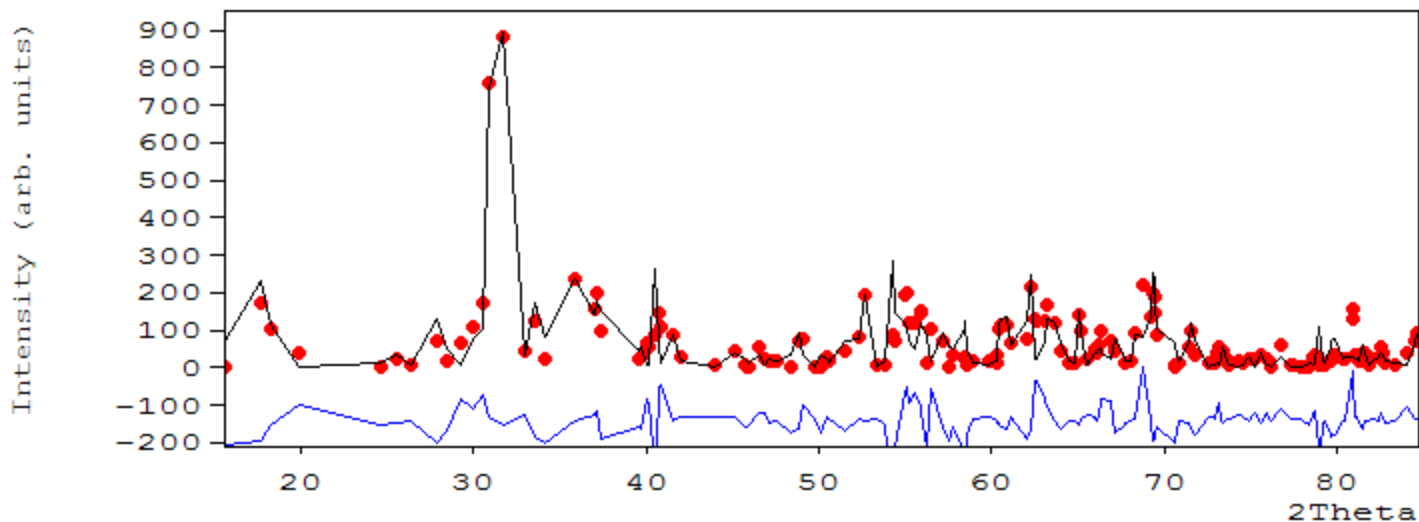
=> Initial configuration state vector:

	Q1_R4+	Q2_R5+	Q3_R5+	Q4_X5+	Q5_X5+	Q6_M2+	Q7_M3+
	1	2	3	4	5	6	7
	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000

NT	Temp	(%Acc)	<Step>	<Rp-factor>
1	6.00	42.04	4.0000	85.4356
2	5.70	56.12	3.7571	86.7607
3	5.41	24.49	3.8286	65.8903
4	5.14	32.04	2.2809	56.3059
5	4.89	31.02	1.6271	49.8507
6	4.64	47.76	1.1245	48.1753
7	4.41	34.29	1.1456	33.1074
8	4.19	42.24	0.8090	35.1309
9	3.98	42.04	0.7418	28.3311

Function evaluations: 4411

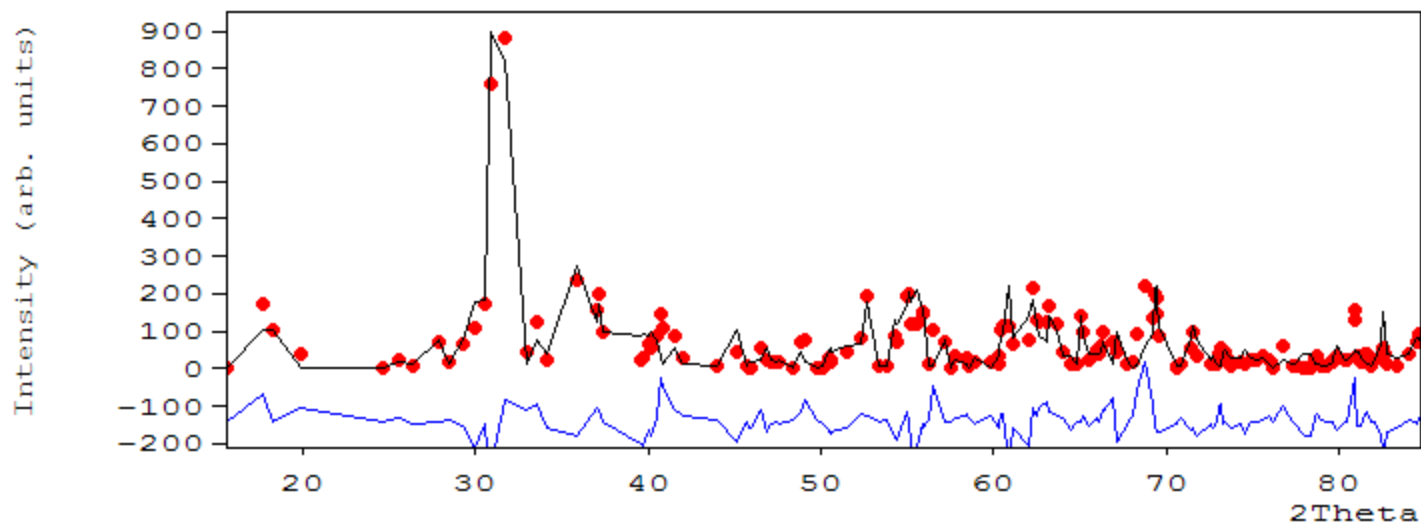
lamn_san.int



```

=> Initial configuration cost: 107.15
=> Initial configuration state vector:
=>   Q1_R4+   Q2_R5+   Q3_R5+   Q4_X5+   Q5_X5+   Q6_M2+   Q7_M3+
=>       1       2       3       4       5       6       7
=>   0.0000   0.0000   0.0000   0.0000   0.0000   0.0000   0.0000
=> NT:  1 Temp:  6.00 (%Acc): 42.04 <Step>:  4.0000 <Rp-factor>: 85.4356
=> NT:  2 Temp:  5.70 (%Acc): 56.12 <Step>:  3.7571 <Rp-factor>: 86.7607
=> NT:  3 Temp:  5.41 (%Acc): 24.49 <Step>:  3.8286 <Rp-factor>: 65.8903
=> NT:  4 Temp:  5.14 (%Acc): 32.04 <Step>:  2.2809 <Rp-factor>: 56.3059
=> NT:  5 Temp:  4.89 (%Acc): 31.02 <Step>:  1.6271 <Rp-factor>: 49.8507
=> NT:  6 Temp:  4.64 (%Acc): 47.76 <Step>:  1.1245 <Rp-factor>: 48.1753
=> NT:  7 Temp:  4.41 (%Acc): 34.29 <Step>:  1.1456 <Rp-factor>: 33.1074
=> NT:  8 Temp:  4.19 (%Acc): 42.24 <Step>:  0.8090 <Rp-factor>: 35.1309
=> NT:  9 Temp:  3.98 (%Acc): 42.04 <Step>:  0.7418 <Rp-factor>: 28.3311
=> NT: 10 Temp:  3.78 (%Acc): 49.39 <Step>:  0.6986 <Rp-factor>: 38.8114
  
```

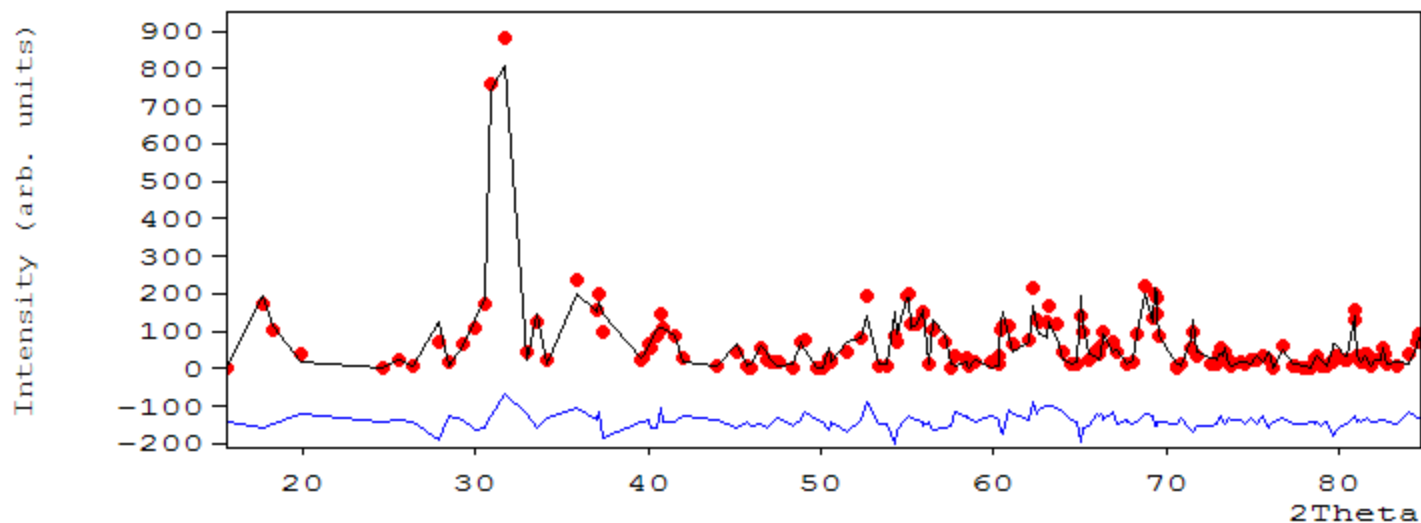
Function evaluations: 4901 lamn_san.int



```

=> NT: 6 Temp: 4.64 (%Acc): 47.76 <Step>: 1.1245 <Rp-factor>: 48.1753
=> NT: 7 Temp: 4.41 (%Acc): 34.29 <Step>: 1.1456 <Rp-factor>: 33.1074
=> NT: 8 Temp: 4.19 (%Acc): 42.24 <Step>: 0.8090 <Rp-factor>: 35.1309
=> NT: 9 Temp: 3.98 (%Acc): 42.04 <Step>: 0.7418 <Rp-factor>: 28.3311
=> NT: 10 Temp: 3.78 (%Acc): 49.39 <Step>: 0.6986 <Rp-factor>: 38.8114
=> NT: 11 Temp: 3.59 (%Acc): 43.88 <Step>: 0.6925 <Rp-factor>: 35.3227
=> NT: 12 Temp: 3.41 (%Acc): 42.24 <Step>: 0.6375 <Rp-factor>: 30.5381
=> NT: 13 Temp: 3.24 (%Acc): 50.41 <Step>: 0.5964 <Rp-factor>: 34.7674
=> NT: 14 Temp: 3.08 (%Acc): 57.14 <Step>: 0.5964 <Rp-factor>: 43.1870
=> NT: 15 Temp: 2.93 (%Acc): 33.67 <Step>: 0.6340 <Rp-factor>: 20.1178
=> NT: 16 Temp: 2.78 (%Acc): 54.08 <Step>: 0.4866 <Rp-factor>: 28.4385
=> NT: 17 Temp: 2.64 (%Acc): 42.24 <Step>: 0.5231 <Rp-factor>: 25.6775
=> NT: 18 Temp: 2.51 (%Acc): 41.22 <Step>: 0.4843 <Rp-factor>: 20.3788
=> NT: 19 Temp: 2.38 (%Acc): 50.00 <Step>: 0.4271 <Rp-factor>: 20.7542
=> NT: 20 Temp: 2.26 (%Acc): 40.61 <Step>: 0.4271 <Rp-factor>: 19.0894
=> NT: 21 Temp: 2.15 (%Acc): 40.00 <Step>: 0.3790 <Rp-factor>: 17.4597
  
```

Function evaluations: 10291 lamn_san.int

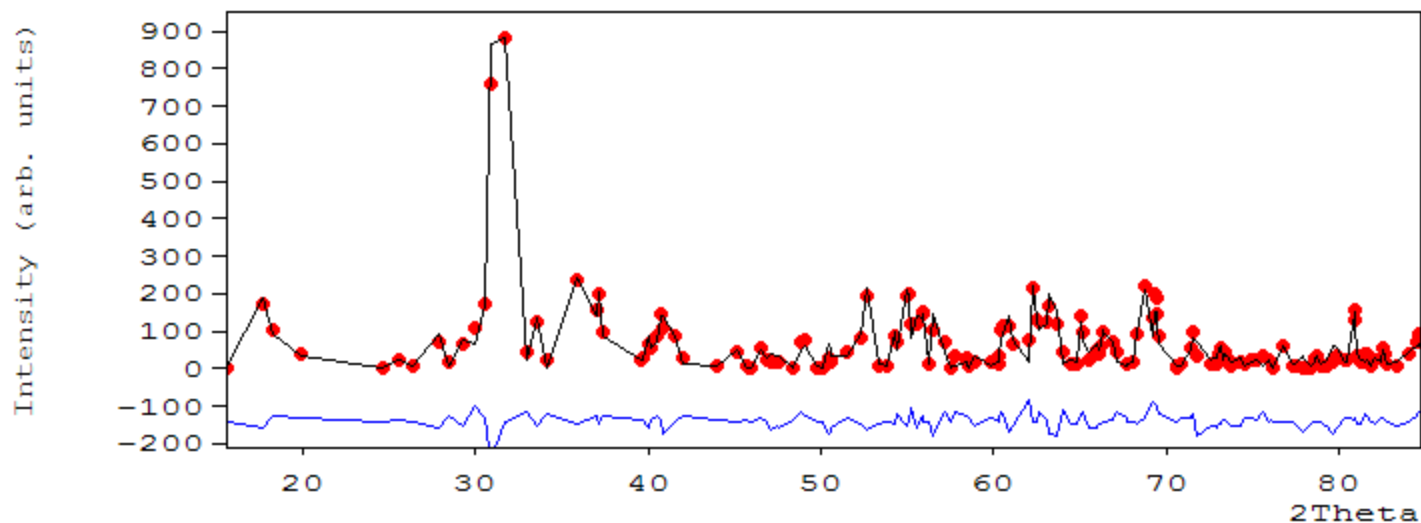


```

=> NT: 13 Temp: 3.24 (%Acc): 50.41 <Step>: 0.5964 <Rp-factor>: 34.7674
=> NT: 14 Temp: 3.08 (%Acc): 57.14 <Step>: 0.5964 <Rp-factor>: 43.1870
=> NT: 15 Temp: 2.93 (%Acc): 33.67 <Step>: 0.6340 <Rp-factor>: 20.1178
=> NT: 16 Temp: 2.78 (%Acc): 54.08 <Step>: 0.4866 <Rp-factor>: 28.4385
=> NT: 17 Temp: 2.64 (%Acc): 42.24 <Step>: 0.5231 <Rp-factor>: 25.6775
=> NT: 18 Temp: 2.51 (%Acc): 41.22 <Step>: 0.4843 <Rp-factor>: 20.3788
=> NT: 19 Temp: 2.38 (%Acc): 50.00 <Step>: 0.4271 <Rp-factor>: 20.7542
=> NT: 20 Temp: 2.26 (%Acc): 40.61 <Step>: 0.4271 <Rp-factor>: 19.0894
=> NT: 21 Temp: 2.15 (%Acc): 40.00 <Step>: 0.3790 <Rp-factor>: 17.4597
=> NT: 22 Temp: 2.04 (%Acc): 44.90 <Step>: 0.3499 <Rp-factor>: 16.5334
=> NT: 23 Temp: 1.94 (%Acc): 45.92 <Step>: 0.3413 <Rp-factor>: 17.2707
=> NT: 24 Temp: 1.84 (%Acc): 46.33 <Step>: 0.3162 <Rp-factor>: 15.8605
=> NT: 25 Temp: 1.75 (%Acc): 50.20 <Step>: 0.3022 <Rp-factor>: 17.9183
=> NT: 26 Temp: 1.66 (%Acc): 43.06 <Step>: 0.3022 <Rp-factor>: 13.5925
=> NT: 27 Temp: 1.58 (%Acc): 40.61 <Step>: 0.2821 <Rp-factor>: 14.2617
=> NT: 28 Temp: 1.50 (%Acc): 47.35 <Step>: 0.2455 <Rp-factor>: 12.0899
  
```

Function evaluations: 13231

lamn_san.int



```

=> NT: 81 Temp: 0.10 (%Acc): 45.92 <Step>: 0.0423 <Rp-factor>: 5.7588
=> NT: 82 Temp: 0.09 (%Acc): 49.39 <Step>: 0.0400 <Rp-factor>: 5.7702

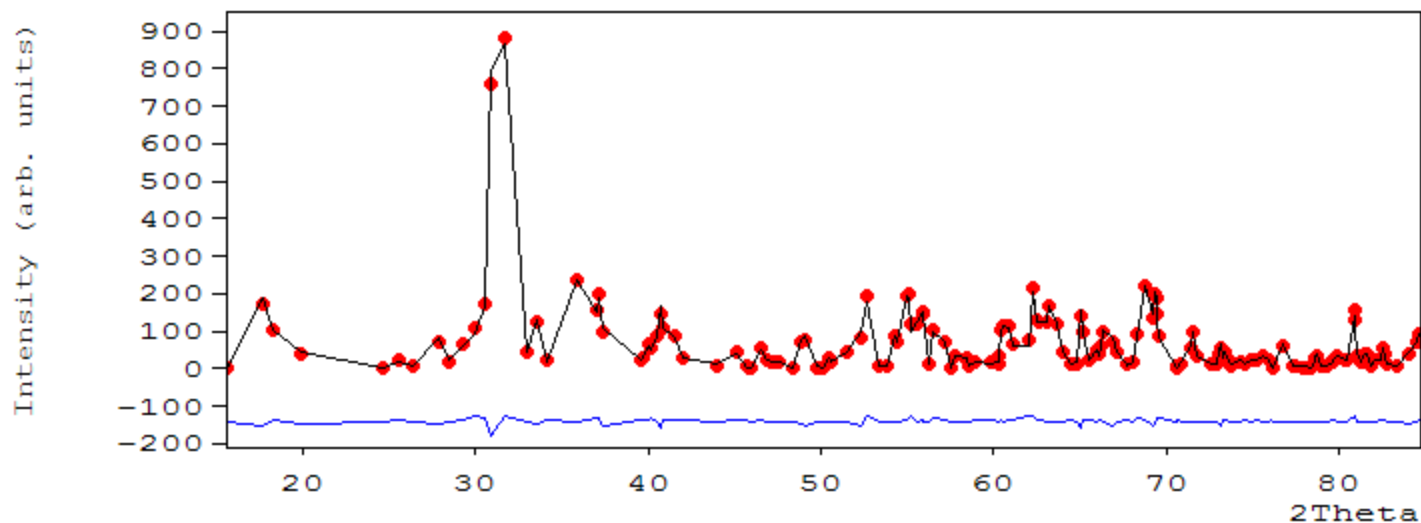
=>BEST CONFIGURATIONS FOUND BY Simulated Annealing FOR PHASE:      1
=> -> Configuration parameters ( 200 reflections):

=> Sol#: 1 RF2= 5.449 ::
=>   Q1_R4+   Q2_R5+   Q3_R5+   Q4_X5+   Q5_X5+   Q6_M2+   Q7_M3+
=>       1       2       3       4       5       6       7
=>   1.2033   0.0830  -0.0167   0.5630   0.1370   0.3626   0.9074

=>           CPU Time: 41.578 seconds
=>           0.693 minutes

=> END   Date:20/09/2008   Time => 15:30:28.109
  
```

Function evaluations: 40181 lamn_san.int



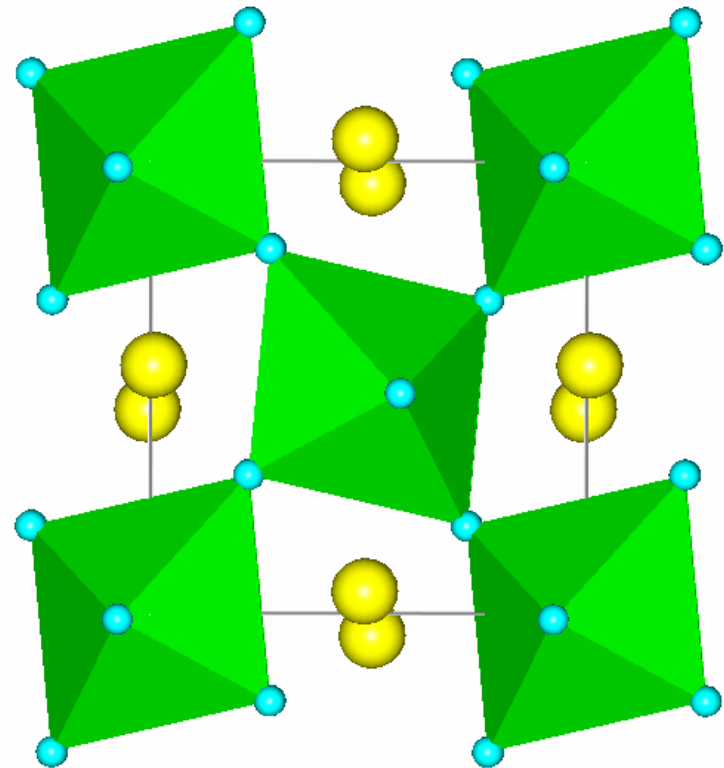
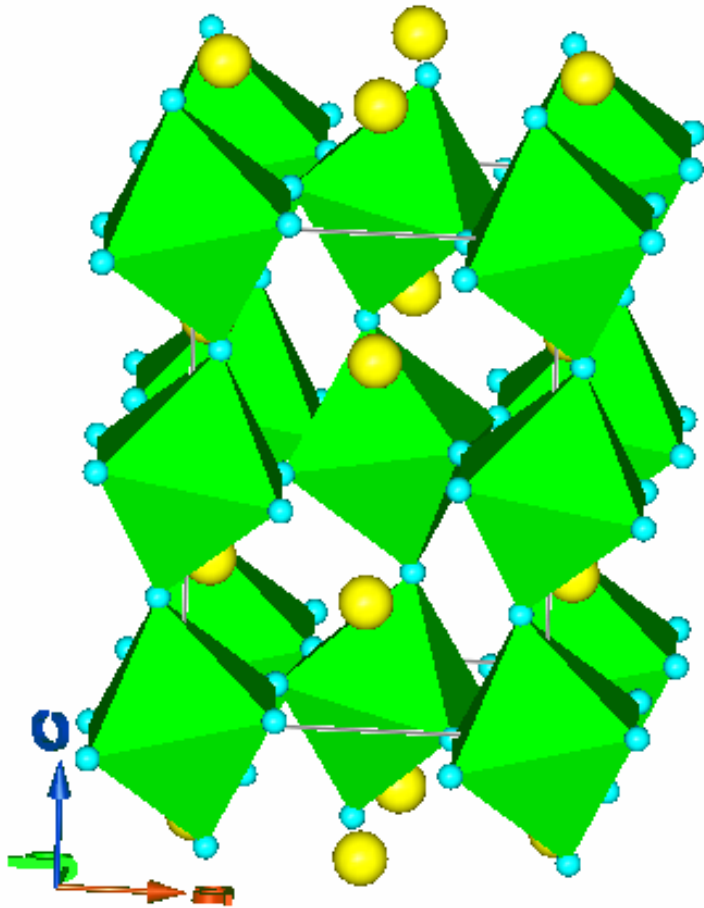
Visualisation of single modes using FullProf Studio

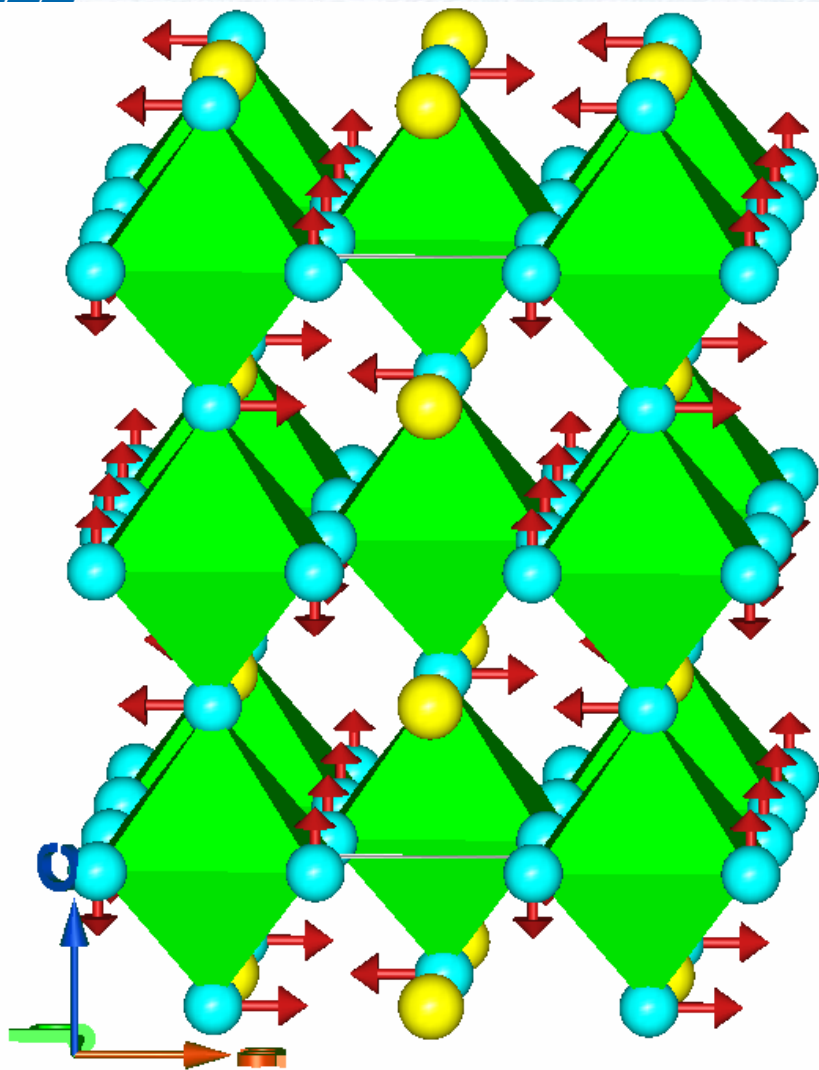
- A part from the normal FST file generated normally for the final crystal structure, **FullProf** outputs a series of FST files containing
- The “virtual structures” corresponding to single modes
(e.g. `A_MODES 7 0 0 0 0 0 0`)
- A representation of the high symmetry phase together with arrows indicating the displacement of atoms in the corresponding mode:
(e.g. `A_MODES 7 1 1 1 1 1 1`)
- Both kinds of representations depending on the mode
(e.g. `A_MODES 7 1 0 0 1 1 0 1`)

The items after the number of modes are:

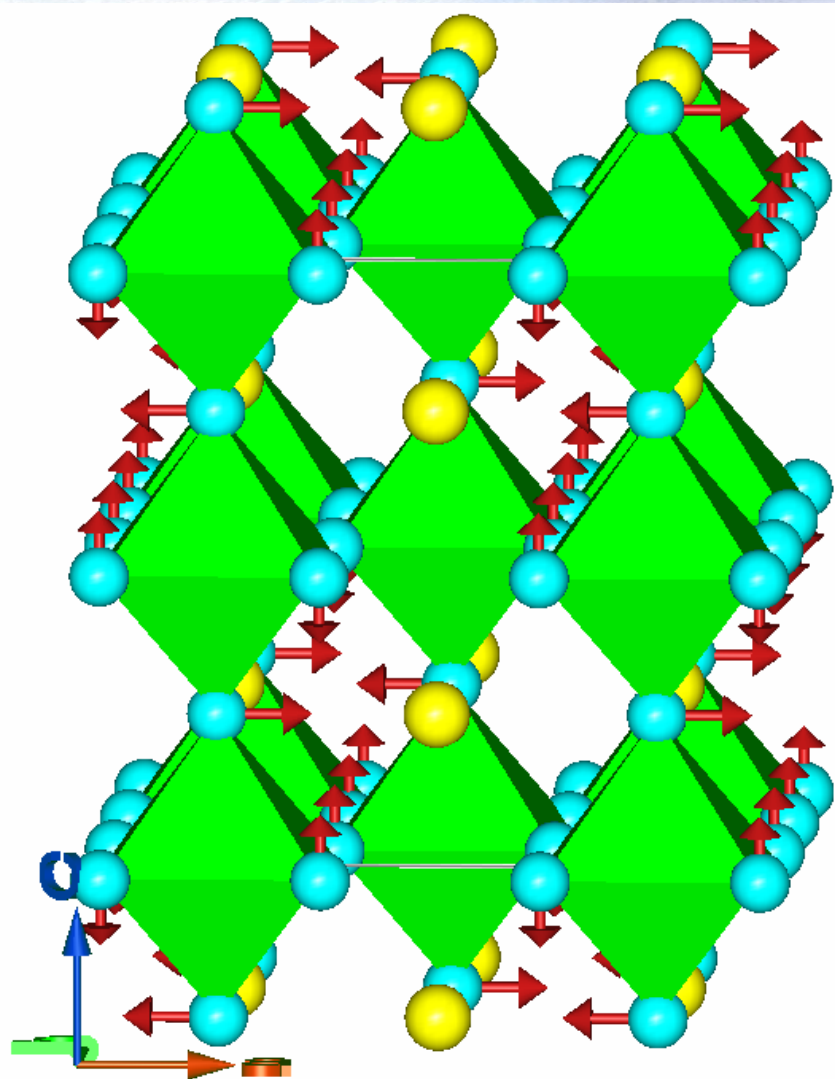
$$p_{\text{mode}}(i) \quad i=1, \dots, n_{\text{modes}}$$

Crystal Structure of LaMnO_3

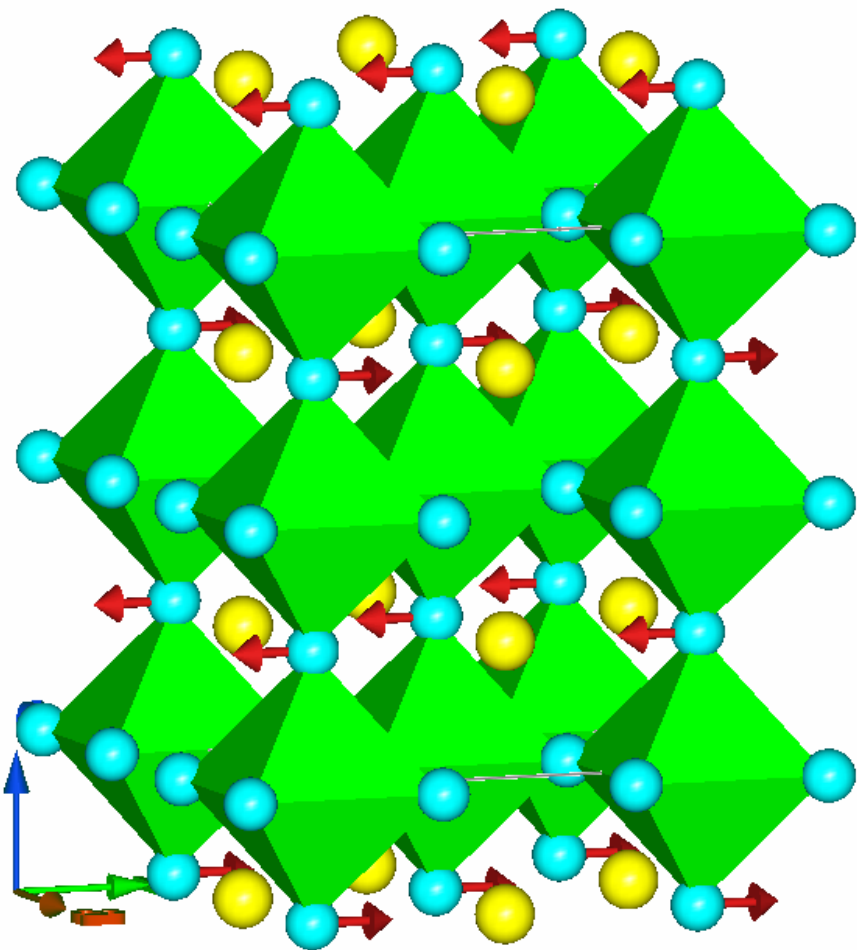




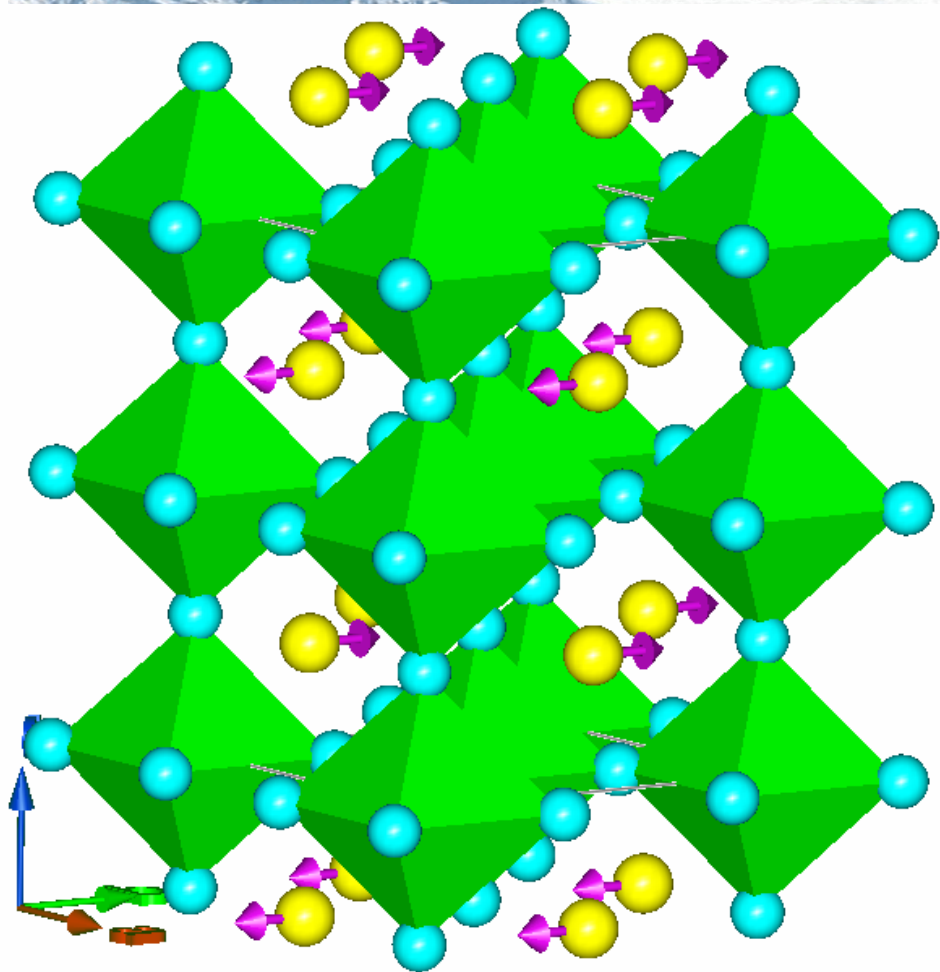
Mode 1, $Q1_{R4+} = -1.18968$
 O1 R4+ (0.0, 0.0, 0.031721)
 O2 R4+ (0.063442, 0.0, 0.0)



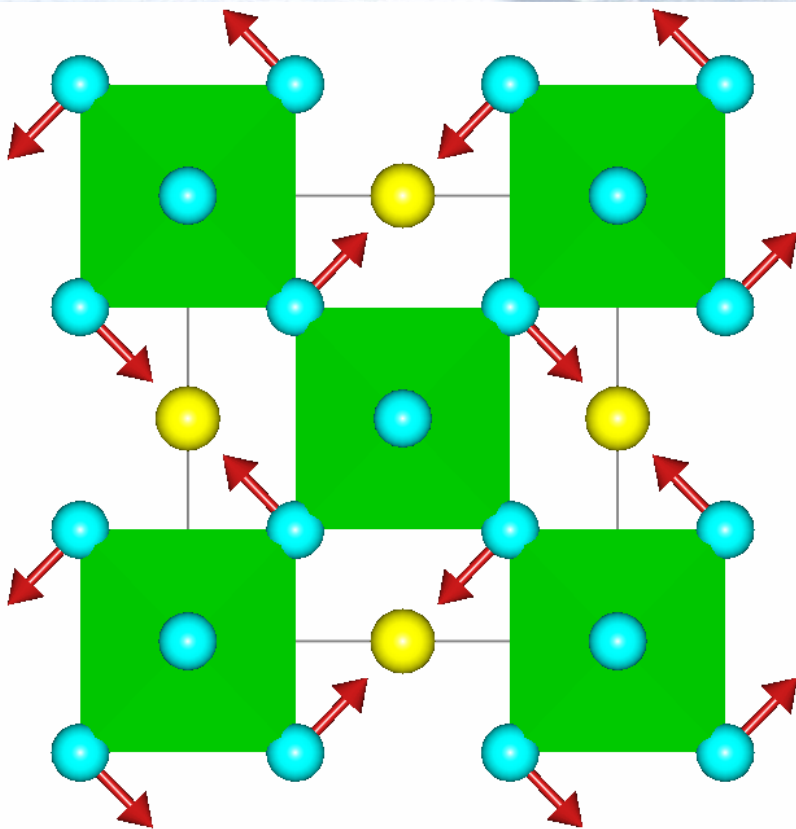
Mode 3, $Q3_{R5+} = 0.018171$
 O1 R5+ (0.0, 0.0, -0.031721)
 O2 R5+ (0.063442, 0.0, 0.0)



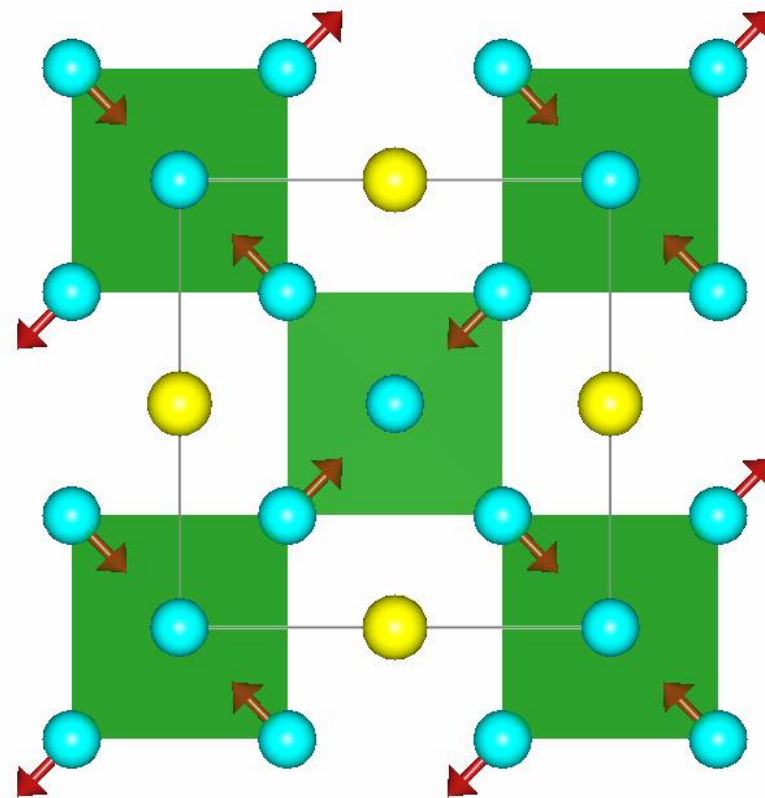
Mode 5, $Q5_X5+ = -0.139910$
 O2 X5+ (0.0, -0.089721, 0.0)



Mode 4, $Q4_X5+ = -0.546082$
 La X5+ (0.0, -0.089721, 0.0)



Mode 7, $Q7_M3+ = 0.901264$
 O1 M3+ $(-0.04486, -0.04486, 0.0)$



Mode 6, $Q6_M2+ = 0.355652$
 O1 M2+ $(0.04486, -0.04486, 0.0)$

Visualisation of Irreps modes using FullProf Studio

If the value of **p_mode(1)=2** (see note of 29 August 2008 in **fp2k.inf**) the other values are not needed.

The program interprets this value as an indication to output in the FST and OUT files the structures corresponding to single irreducible representations (Irreps).

All modes corresponding to a single Irrep are combined in the FST file.

Visualisation of modes using FullProf Studio: Summary

Examples:

A_MODES 7 7 → All the 7 independent modes are represented by displacement vectors (arrows)

A_MODES 7 -7 → All the 7 independent modes are represented by virtual distorted structures

A_MODES 7 -3 → No output of independent modes in FST files

Visualisation of single modes using FullProf Studio

Examples:

A_MODES 7 2 → Modes regrouped in an FST
file per irreducible representation (arrows)

A_MODES 7 -2 → Modes regrouped in an FST
file per irreducible representation (structures)

A_MODES 7 1 1 1 0 1 1 0 → Explicit output of
all modes (1: arrows, 0: distorted structure)

Visualisation of single modes using FullProf Studio

Examples:

A_MODES 7 4 1 3 -4 7 → Only the 4 modes 1,3,4 and 7 are output in FST files. All of them, except the mode 4, are represented by arrows.

A more complex
case:
 LiMn_2O_4

A complex structure determined and refined with FullProf

VOLUME 81, NUMBER 21

PHYSICAL REVIEW LETTERS

23 NOVEMBER 1998

Electronic Crystallization in a Lithium Battery Material: Columnar Ordering of Electrons and Holes in the Spinel LiMn_2O_4

J. Rodríguez-Carvajal,¹ G. Rousse,² C. Masquelier,² and M. Hervieu³

¹Laboratoire Léon Brillouin (CEA-CNRS), CEA/Saclay, 91191 Gif sur Yvette Cedex, France

²Laboratoire de Chimie des Solides, Université Paris-Sud, 91405 Orsay Cedex, France

³CRISMAT, ISMRA, 6 Boulevard du Maréchal Juin, 14050 Caen Cedex, France

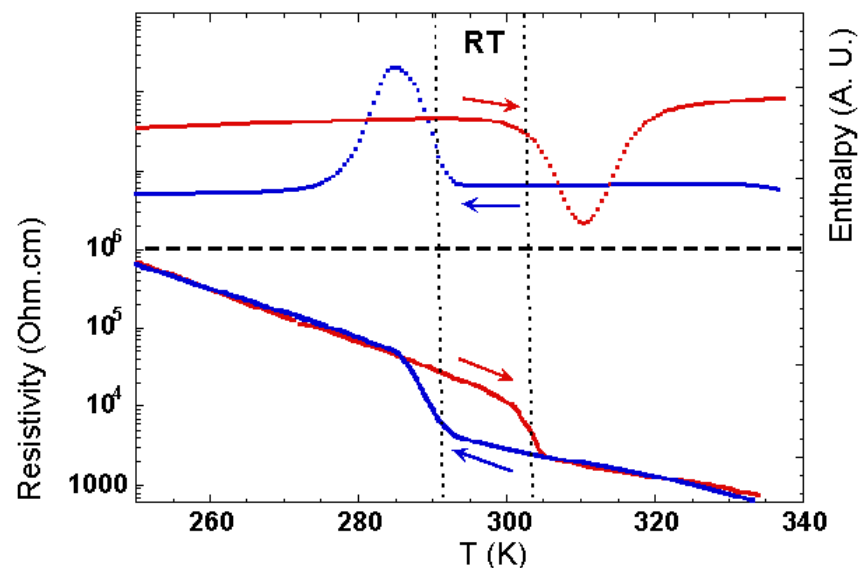
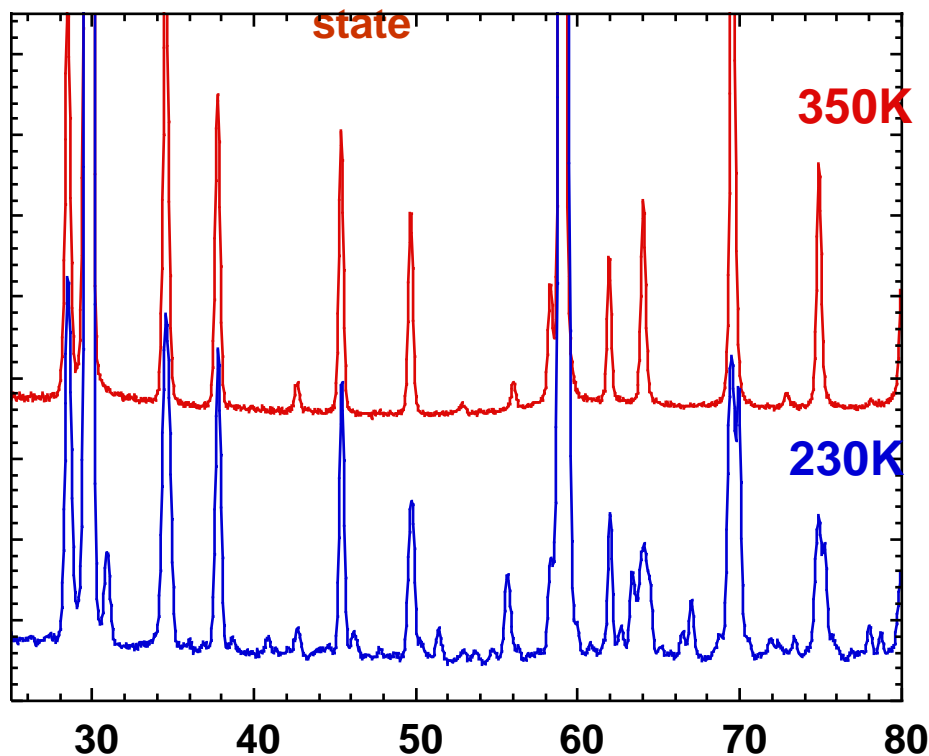
(Received 13 July 1998)

LiMn_2O_4 presents a first order structural transition at 290 K that was known to perturb the functioning as cathode in rechargeable Li batteries. We have solved the structure at 230 K and deciphered unambiguously the nature of this phase transition. The analysis of valence bond sums shows that the transition results from a partial charge ordering: two of the five Mn sites correspond to well-defined Mn^{4+} and the other three sites are close to Mn^{3+} ions. Charge ordering is accompanied by simultaneous orbital ordering due to the Jahn-Teller effect in Mn^{3+} ions. The microscopic details obtained from the structure are crucial for understanding the electron hopping persisting below the transition. [S0031-9007(98)07667-4]

Electronic crystallization in a Li battery material: columnar ordering of electron and holes in the spinel LiMn_2O_4

J. Rodríguez-Carvajal, G. Rousse, Ch. Masquelier and M. Hervieu
Physical Review Letters, **81**, 4660 (1998)

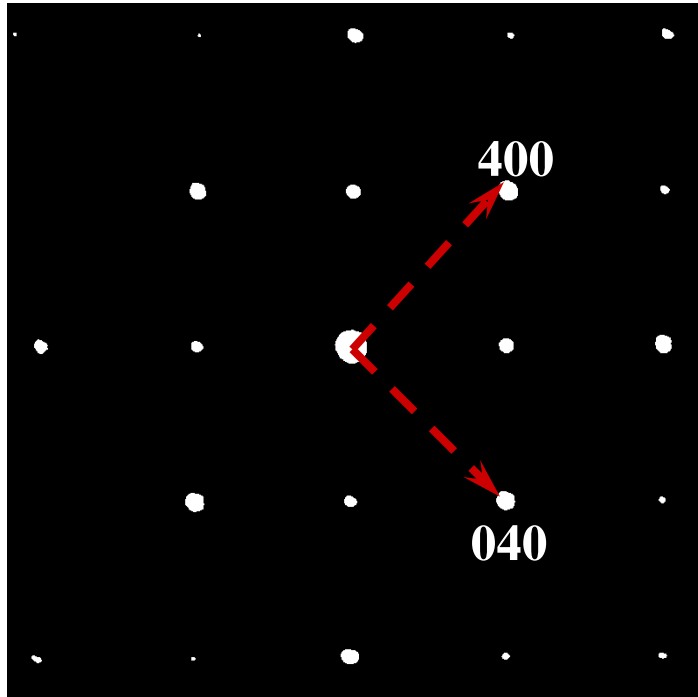
$\text{Li}_{\text{Td}} [\text{Mn}_2]_{\text{Oct.}} \text{O}_4$: $\text{Mn}^{3.5+}$
High temperature: mixed valence




Orthorhombic Distortion
Superstructure reflections

LiMn₂O₄: Electron Diffraction

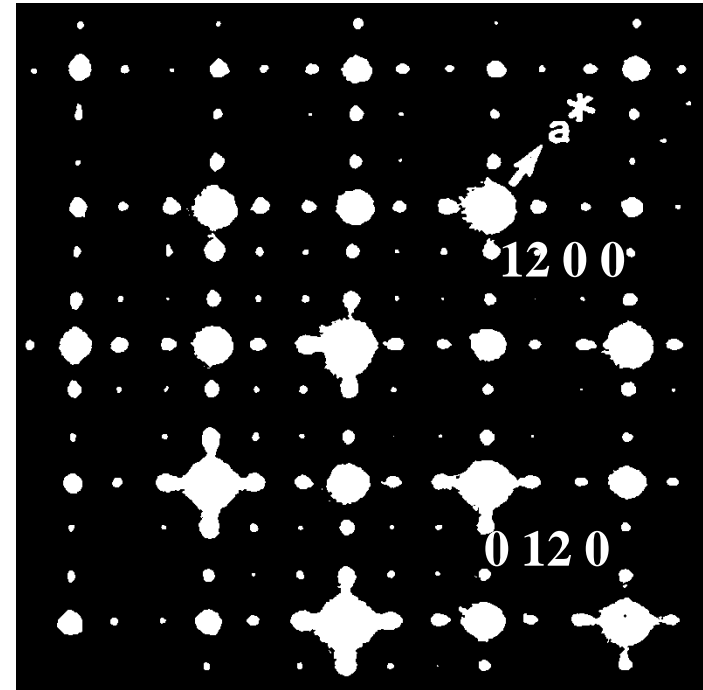
320 K



Cubic $Fd\bar{3}m$
 $a = 8,248 (1) \text{ \AA}$

x 9


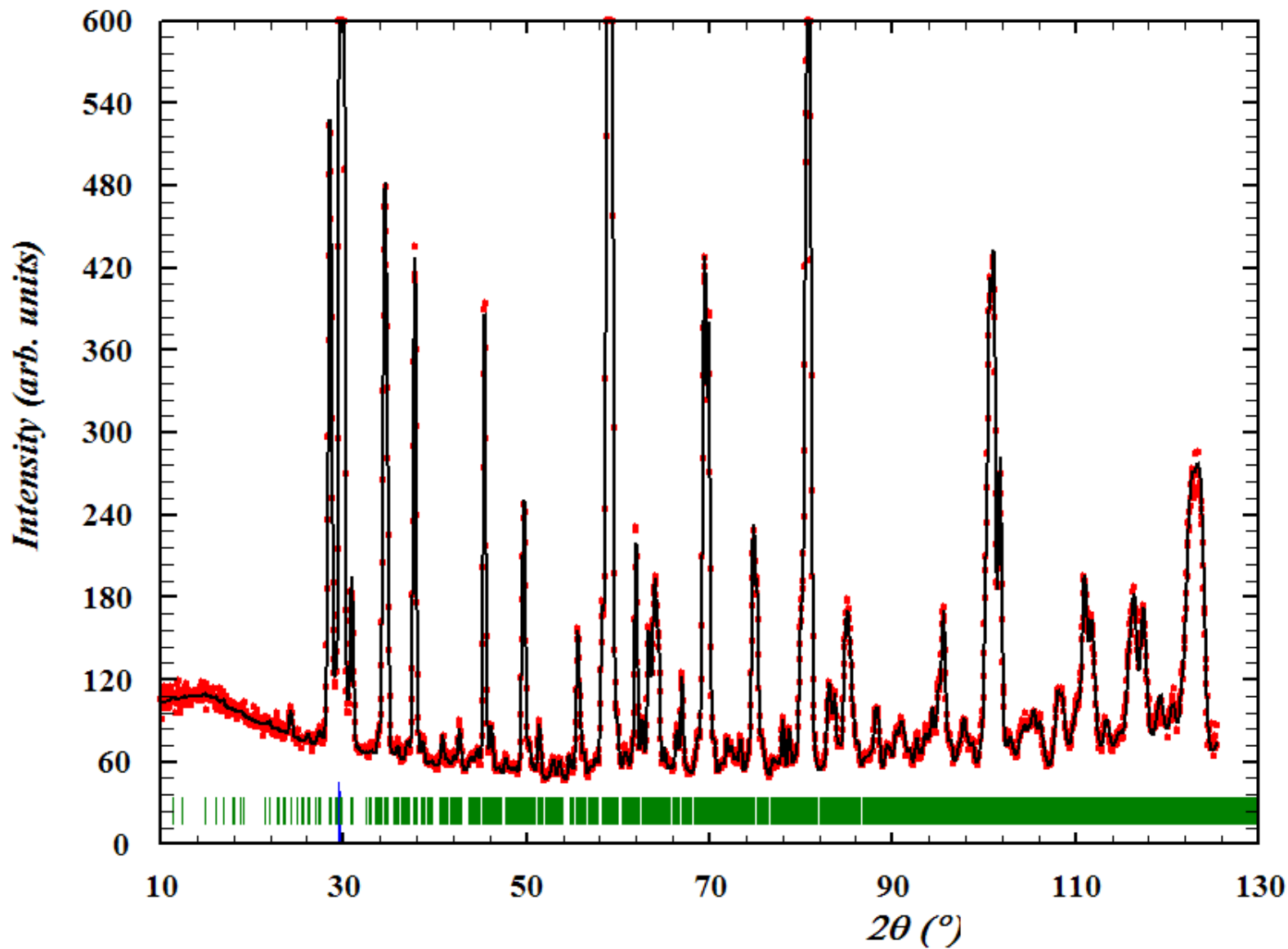
230 K



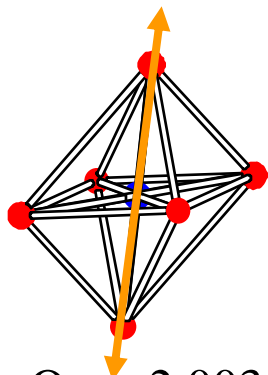
Orthorhombic $Fddd$
 $a = 24.7435(5) \text{ \AA}$
 $b = 24.8402(5) \text{ \AA}$
 $c = 8.1989(1) \text{ \AA}$

J. Rodríguez-Carvajal *et al*, *PRL*, **81**, 4660 (1998)

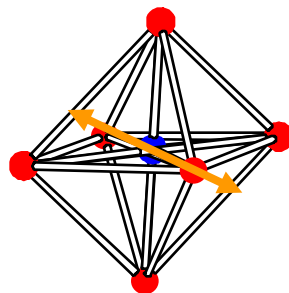
Refinement of the charge-ordered phase



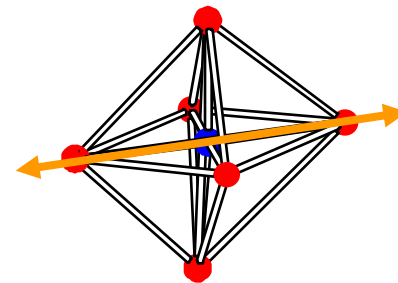
LiMn₂O₄: Partial Charge Ordering



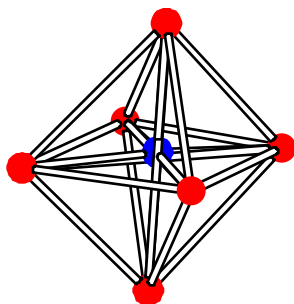
$\langle \text{Mn-O} \rangle = 2,003(2) \text{ \AA}$
Mn(1) = 3.20+
 $\Delta = 20.6$



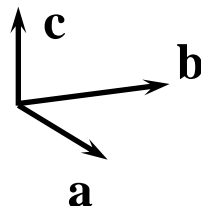
$\langle \text{Mn-O} \rangle = 1,996(4) \text{ \AA}$
Mn(2) = 3.27+
 $\Delta = 19.4$



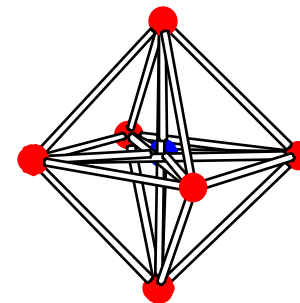
$\langle \text{Mn-O} \rangle = 2,020(5) \text{ \AA}$
Mn(3) = 3.12+
 $\Delta = 36.6$



$\langle \text{Mn-O} \rangle = 1,903(4) \text{ \AA}$
Mn(4) = 4.02+
 $\Delta = 4.6$

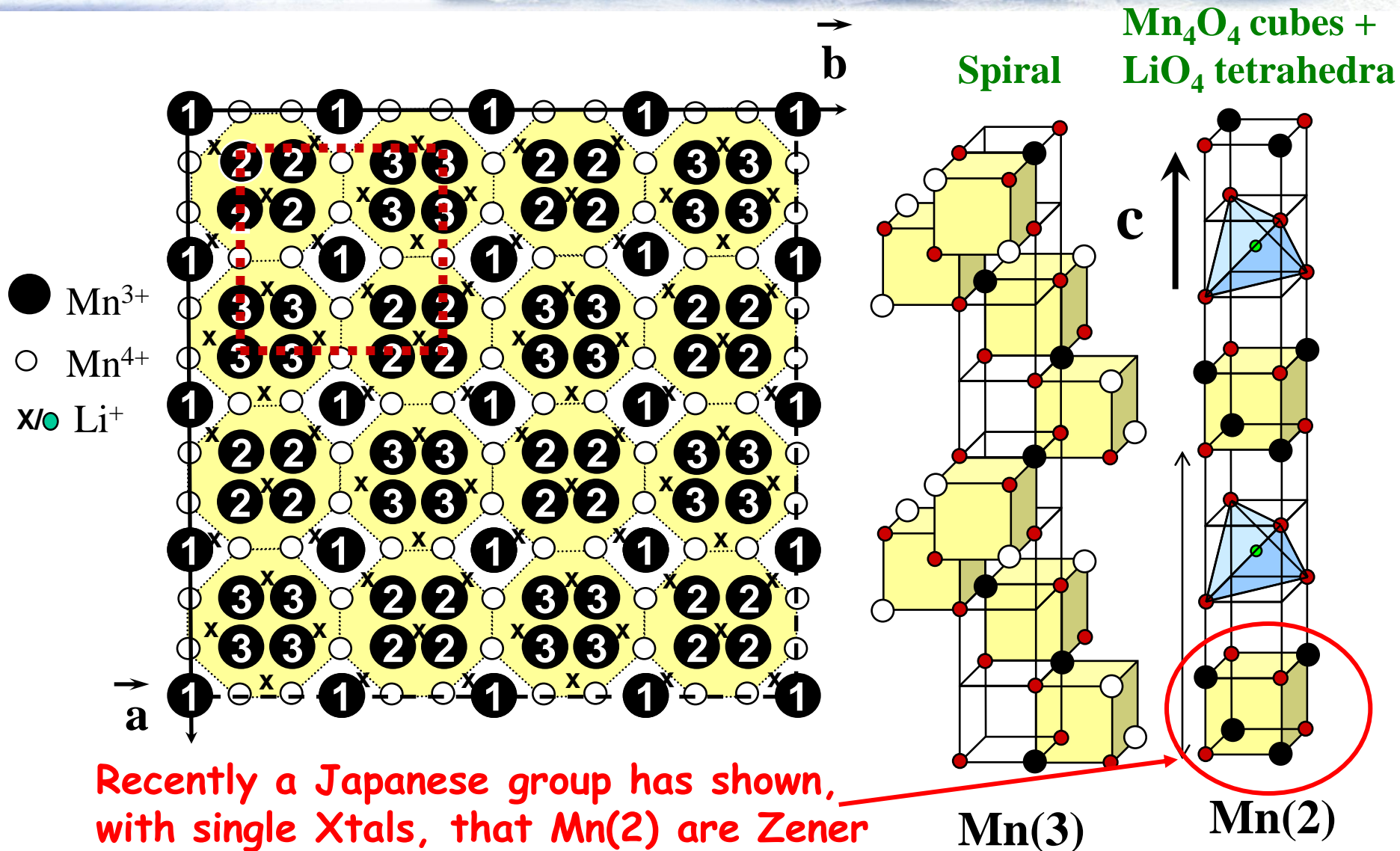


64 « Mn⁴⁺ »
80 « Mn³⁺ -like »
8 delocalised holes



$\langle \text{Mn-O} \rangle = 1,915(4) \text{ \AA}$
Mn(5) = 3.90+
 $\Delta = 6.1$

Distribution of Mn^{3+} and Mn^{4+} in the cell



Recently a Japanese group has shown, with single Xtals, that Mn(2) are Zener Polarons involving 4Mn