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RMCProfile: Big Box modelling of crystalline to amorphous materials

Matt Tucker (ORNL, USA)

Wojciech Slawinski (Warsaw)

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ESRF

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ANALYSIS OF DIFFRACTION DATA IN REAL SPACE

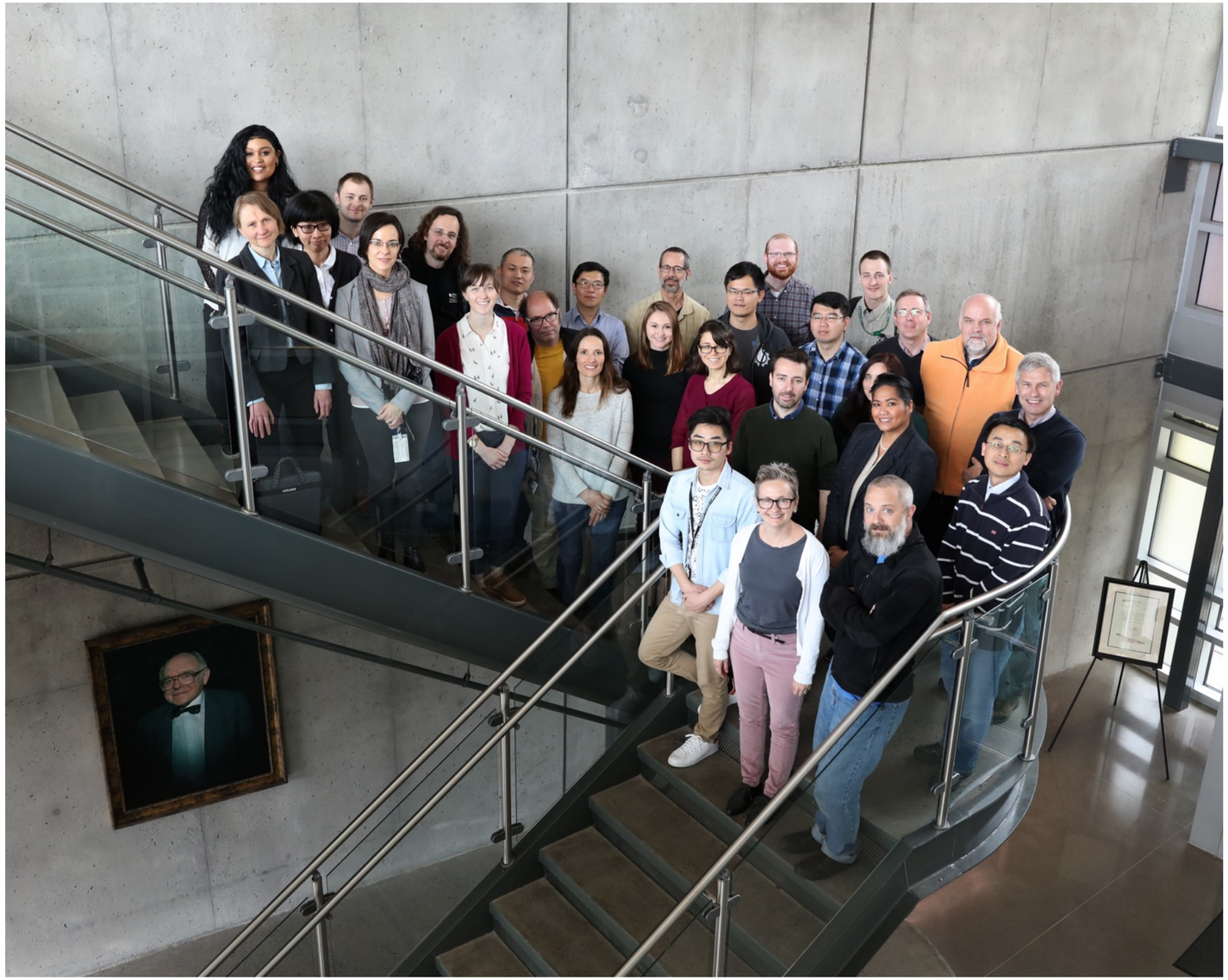
16-21 OCT 2022

GRENOBLE, FRANCE

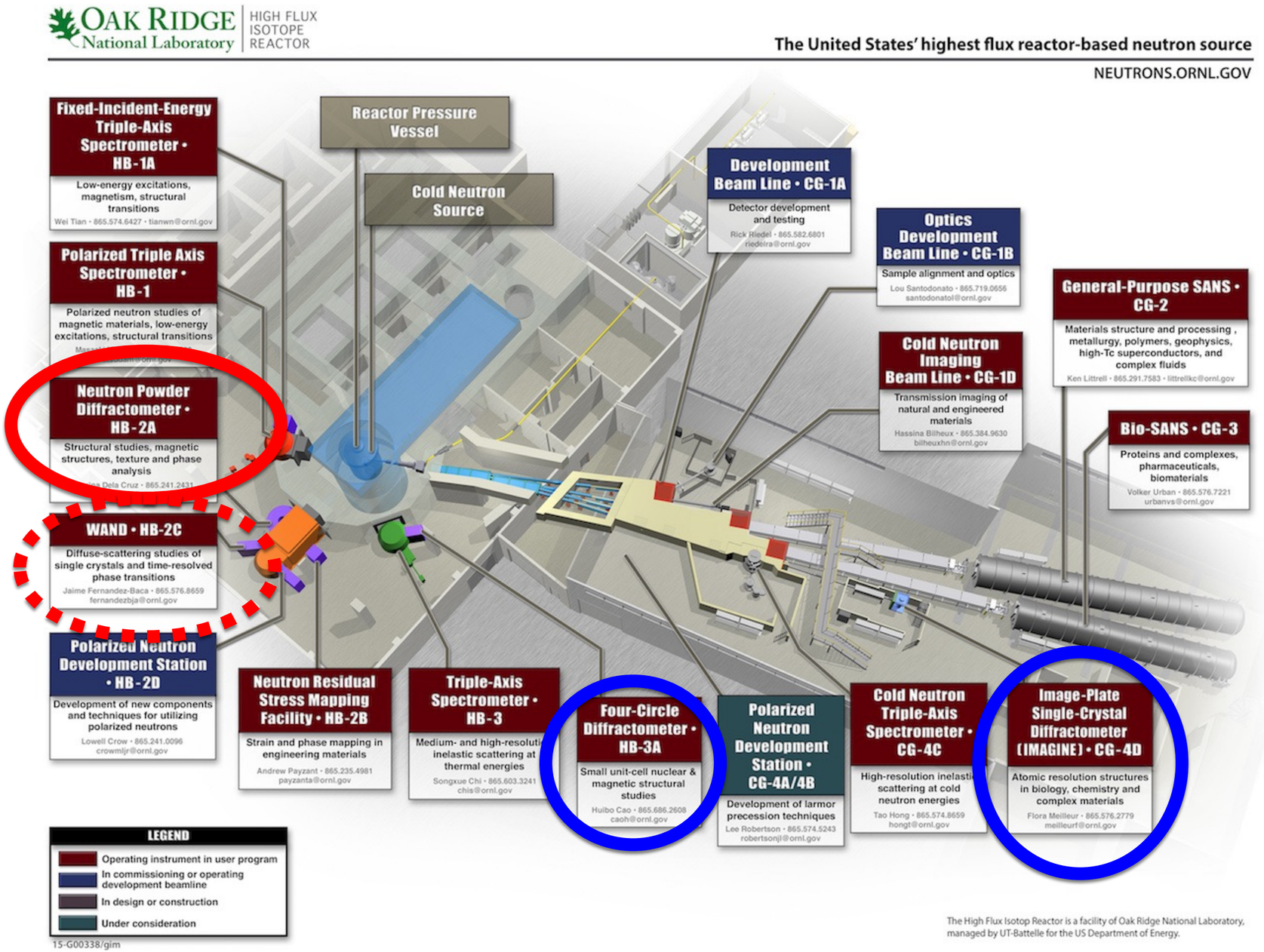
Diffraction Section @ ORNL

Powder
Diffraction Group

Single Crystal
Diffraction Group



High Flux Isotope Reactor



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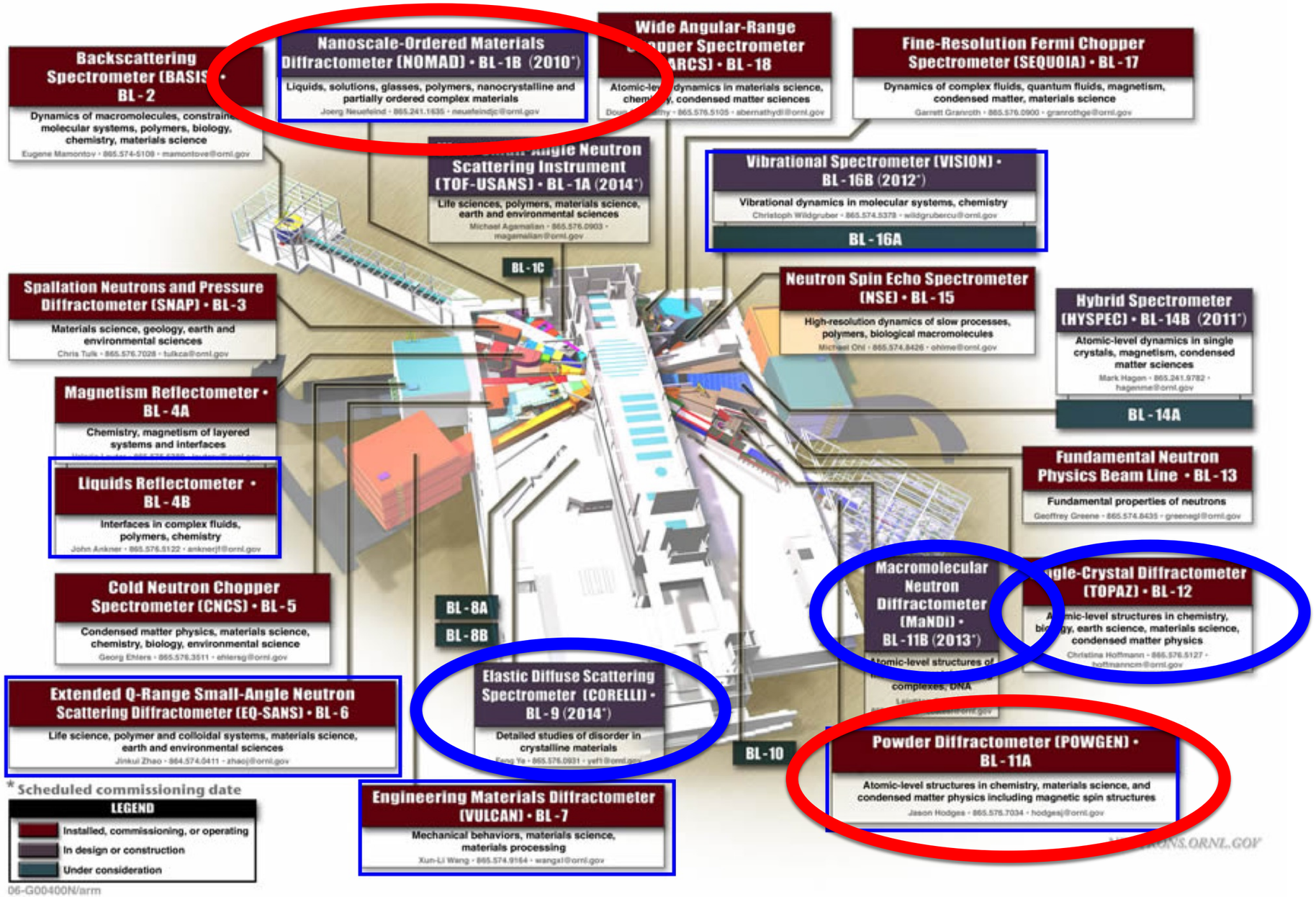
Spallation Neutron Source



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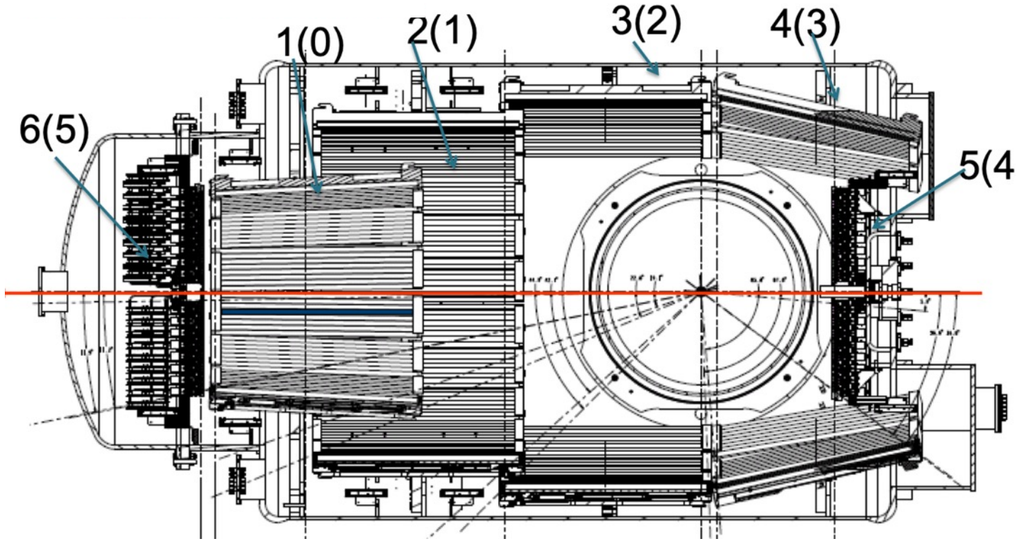
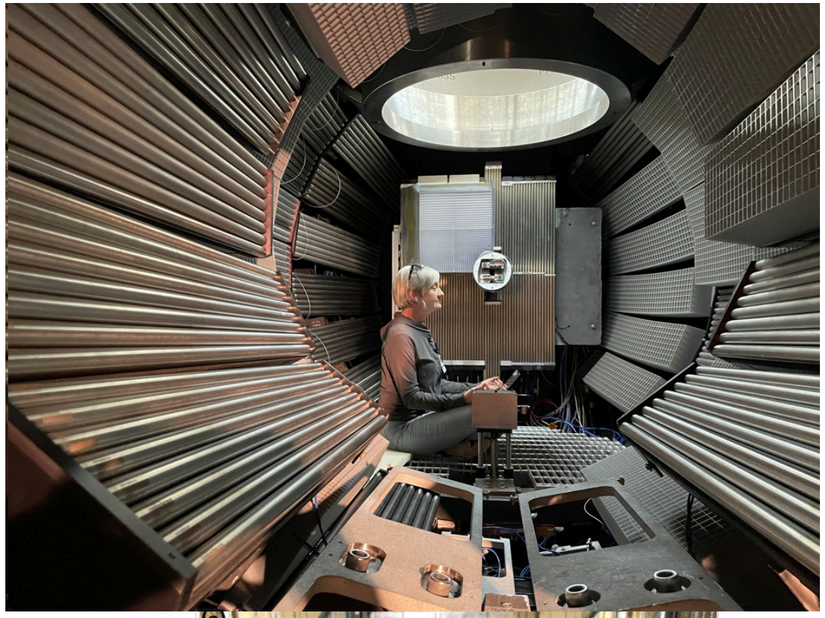
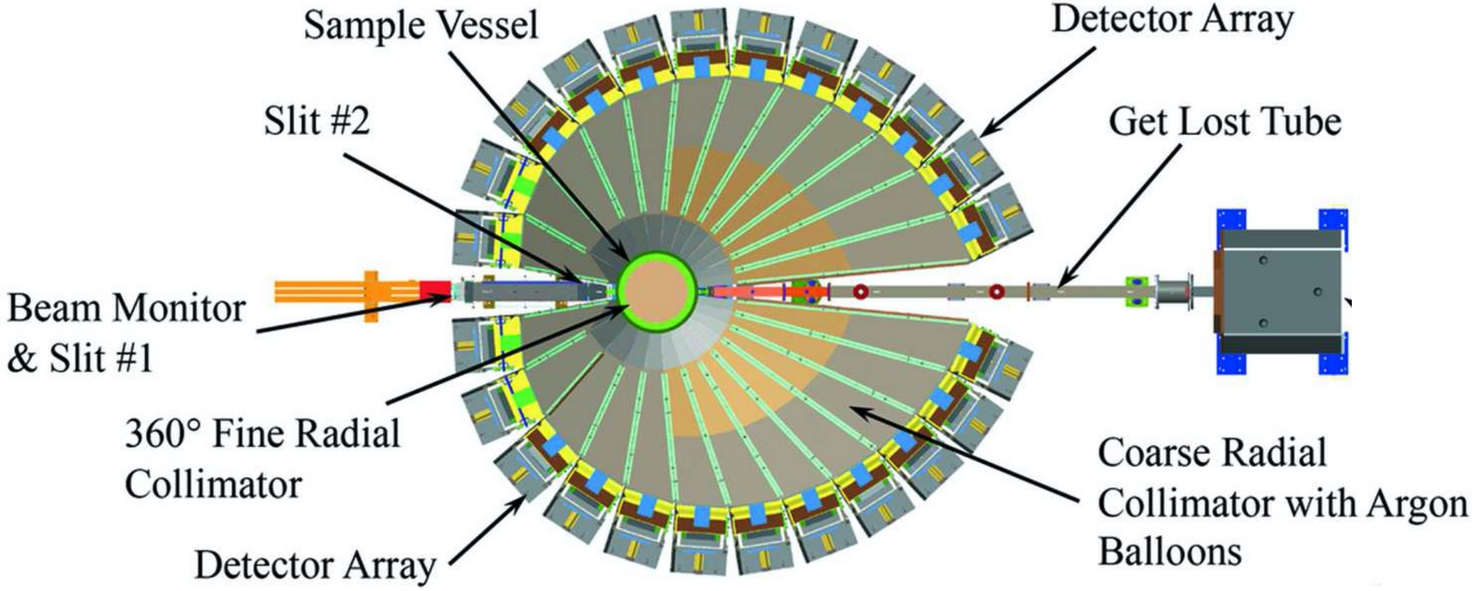
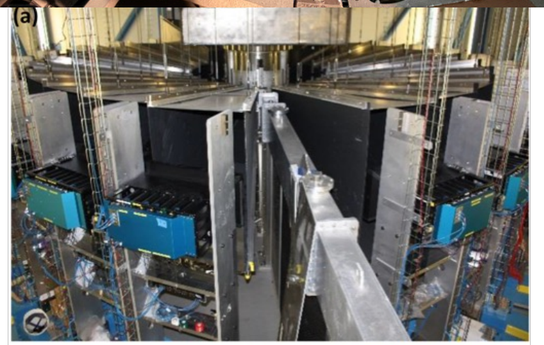
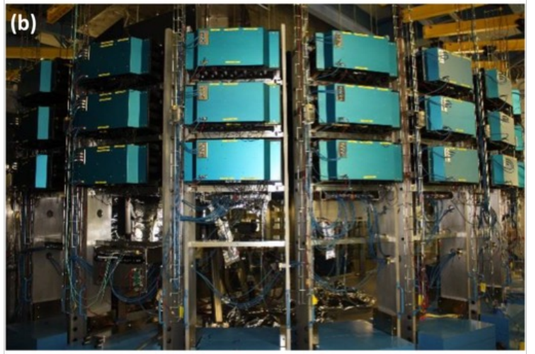
NOMAD and POWGEN











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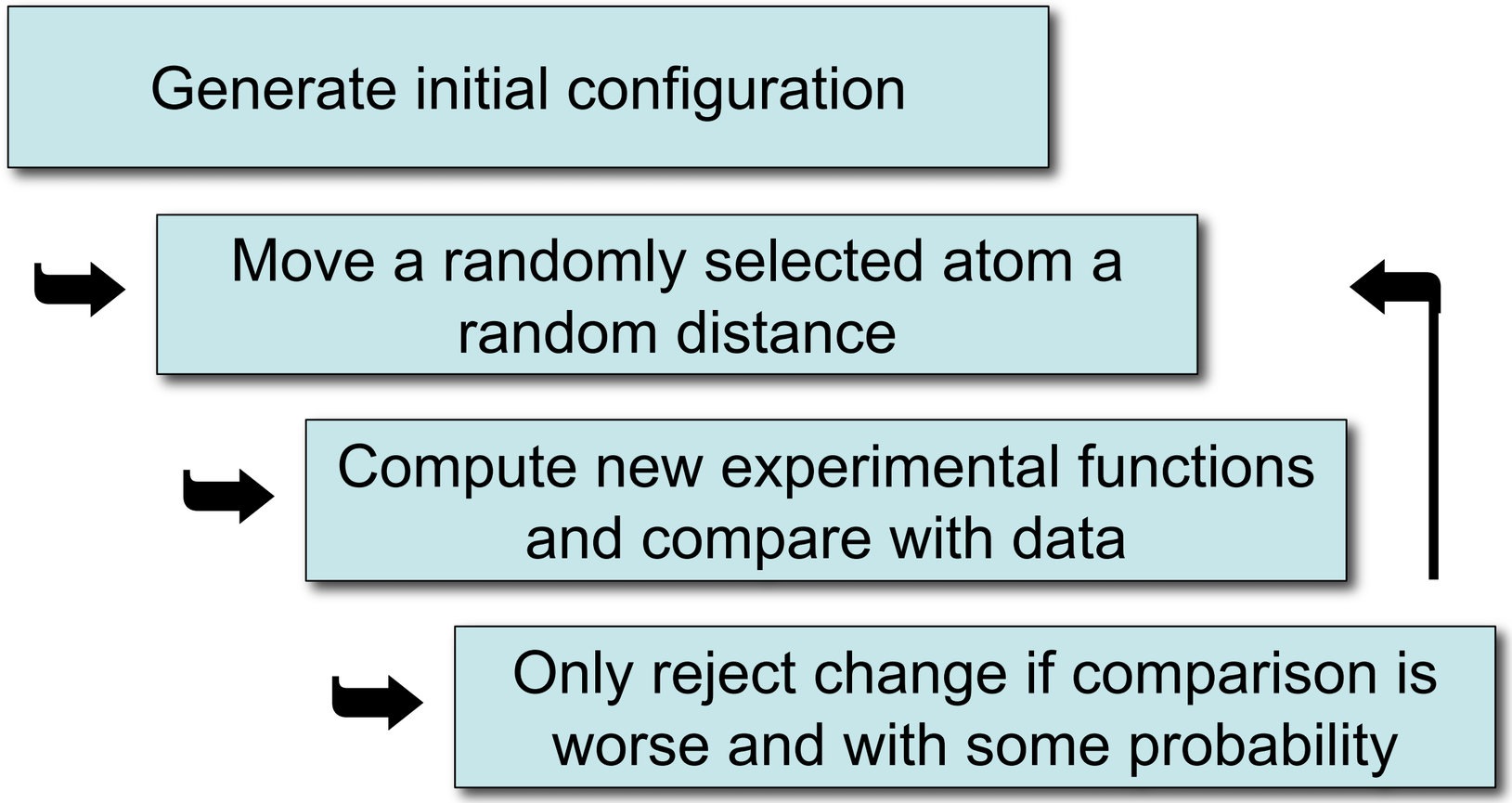
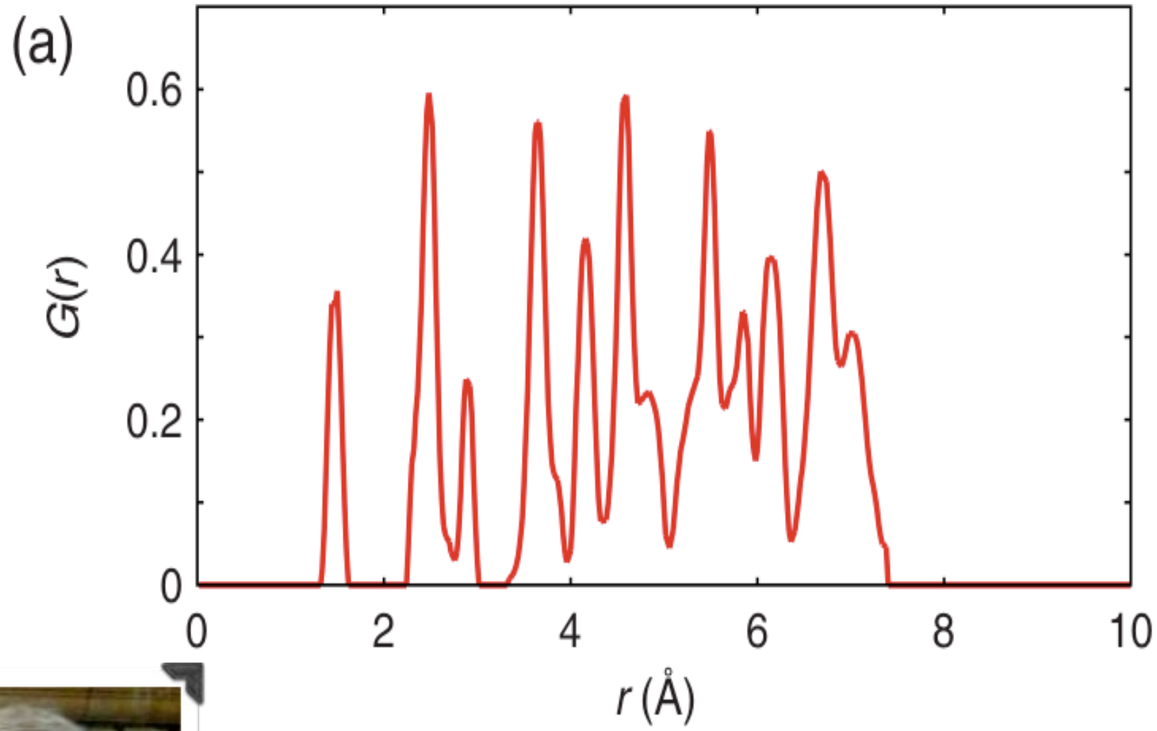
The Reverse Monte Carlo algorithm



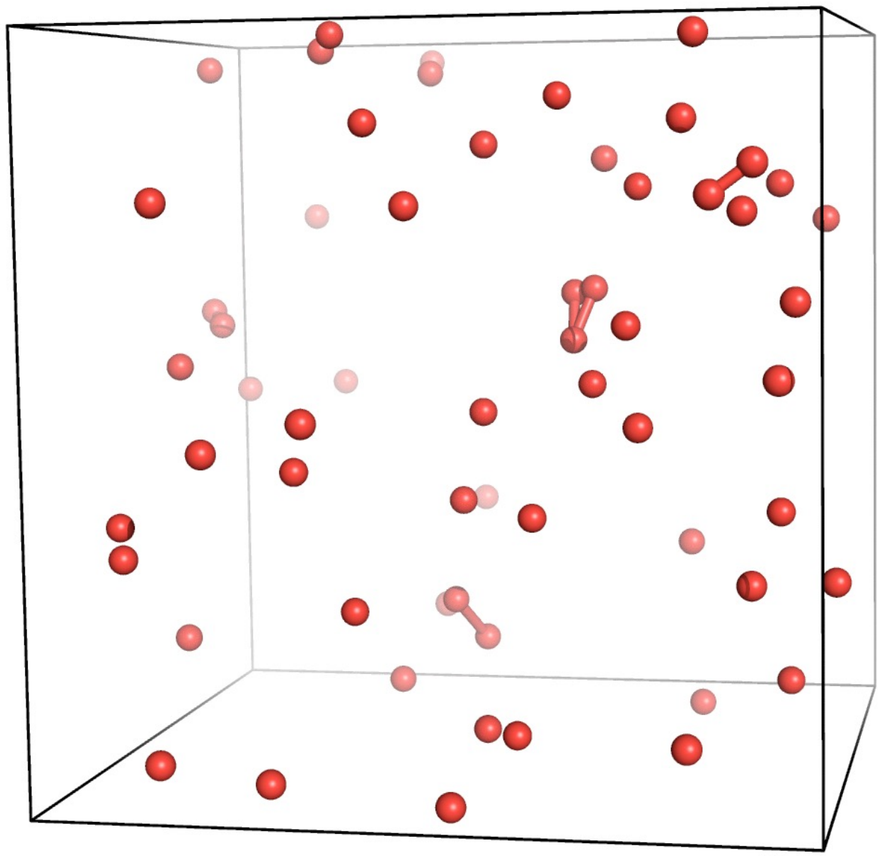
RMC in action: C60



Cliffe M J *et al*
PRL **104** (2010) 1255013

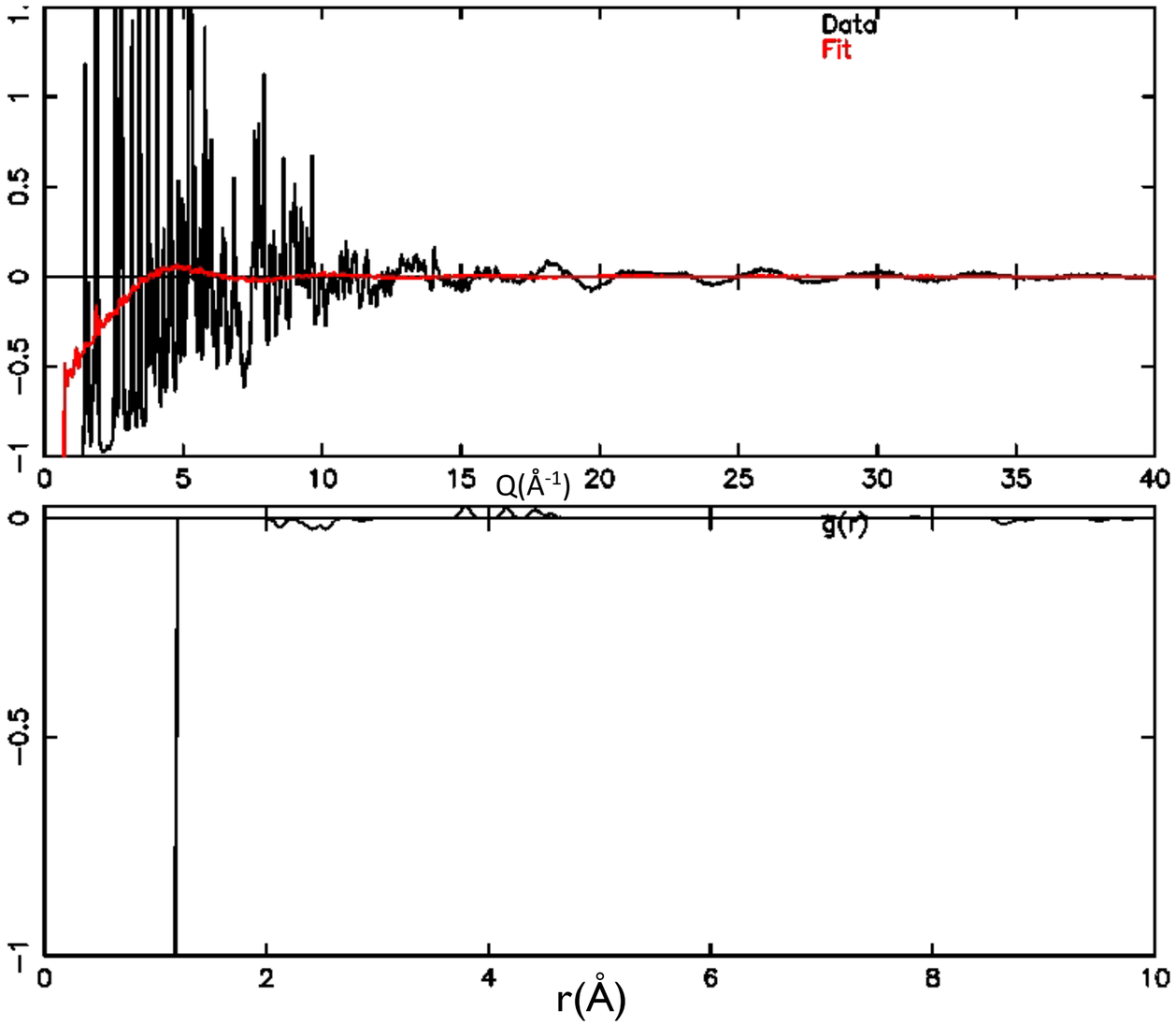
INVERT

RMC in action: C60

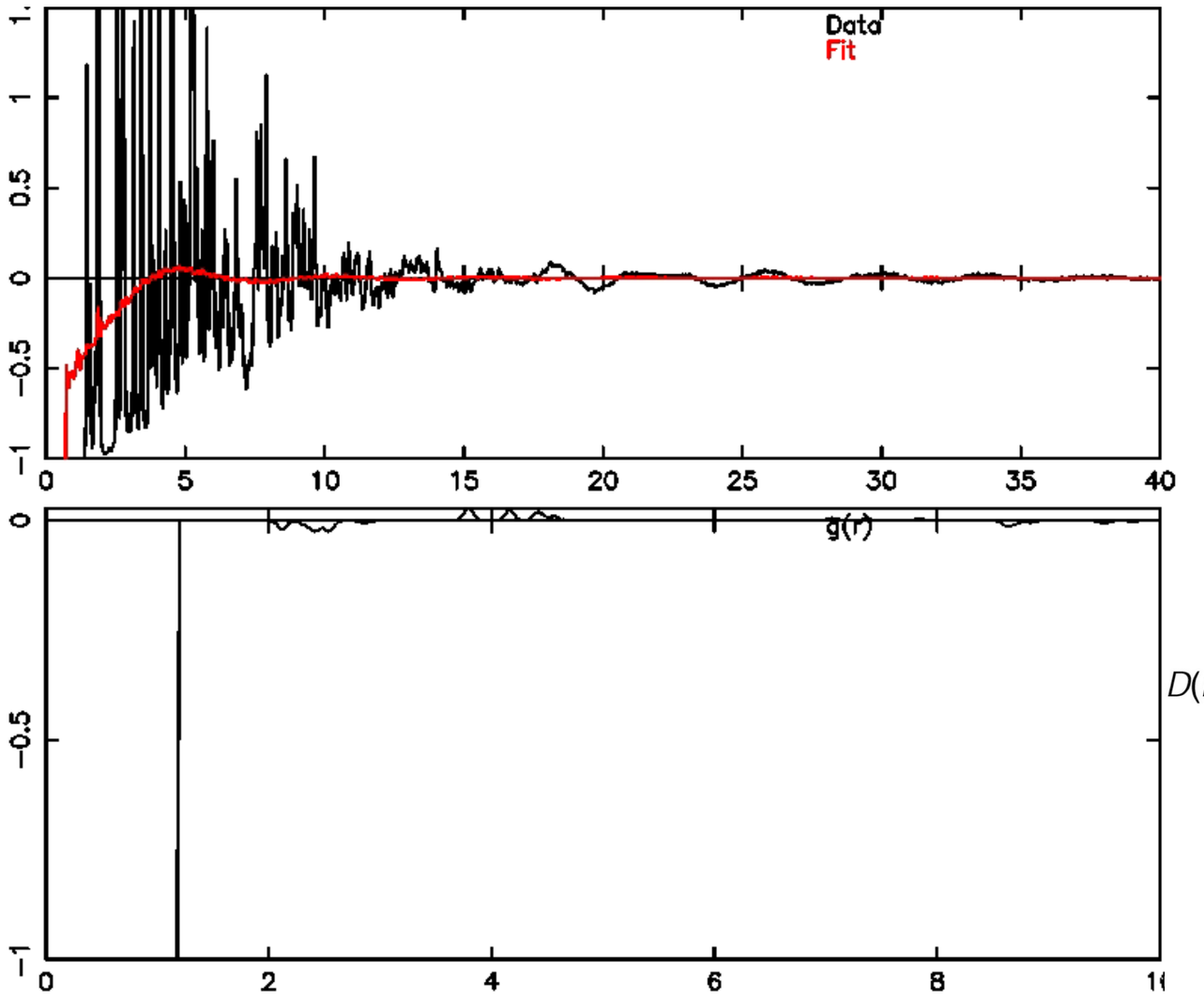


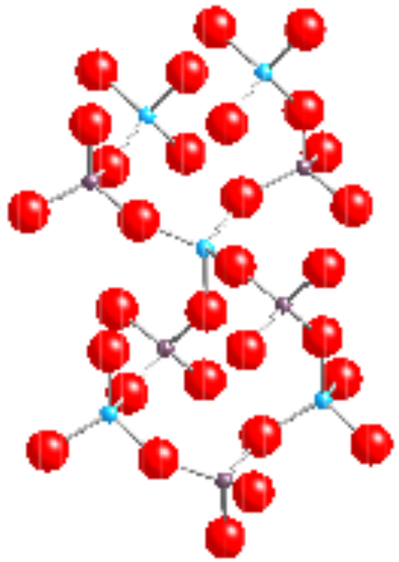
Cliffe M J *et al, PRL* **104** (2010) 1255013

RMC in action

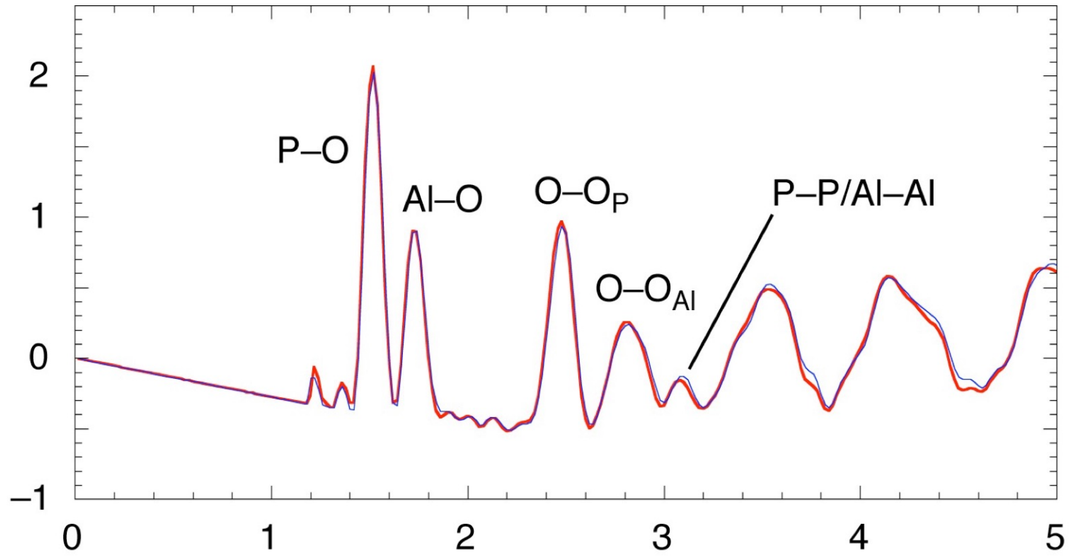


RMC in action





$AlPO_4$



RMC programs

rmcprofile.pages.ornl.gov

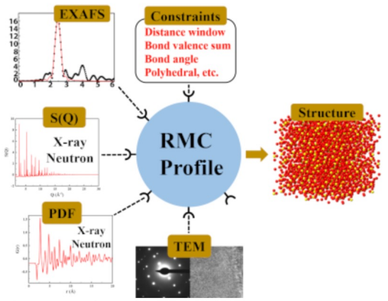
RMC PROFILE

SEARCH ABOUT POSTS PACKAGE COMMUNITY OTHERS

RMCProfile

Reverse Monte Carlo for crystalline and disordered materials

Welcome to the RMCProfile website hosted at Oak Ridge National Laboratory (ORNL) in US. Here you can download the RMCProfile software, view documentation and examples, join community for discussion and learn about updates of the package, etc.



This version of RMC was built from the original RMCA code of McGreevy & Pusztai to determine the local structure of crystalline materials while still being capable of analyzing disordered systems. The current version of RMCProfile results from a collaboration between scientists at ISIS facility (UK), Spallation Neutron Source (SNS at Oak Ridge National Laboratory, US), University of Cambridge (UK), University of Oxford (UK), Queen Mary University of London

https://rmcprofile.pages.ornl.gov/#

External Links

Monte Carlo

DISCUS
RMC++
HRMC
fullrmc
RMC for EXAFS
EPSR
RMC at ISIS facility
RMCProfile website - legacy
Wikipedia for RMC

RMC PROFILE

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ssion. Discussions are hosted as
ssion, one does need a GitHub

ology

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Facilities

US - ORNL - NOMAD
US - ORNL - POWGEN
US - APS - 11-ID-B
US - BNL - 28-ID-1
UK - ISIS - GEM
UK - ISIS - Polaris
UK - Diamond - I15-1
Japan - J-PARC - NOVA
Japan - SPRING8 - BL14B1
Japan - SPRING8 - BL22XU
Europe - ESS - HEIMDAL
China - SSRF - BL13W1

Yuanpeng Zhang (ORNL)

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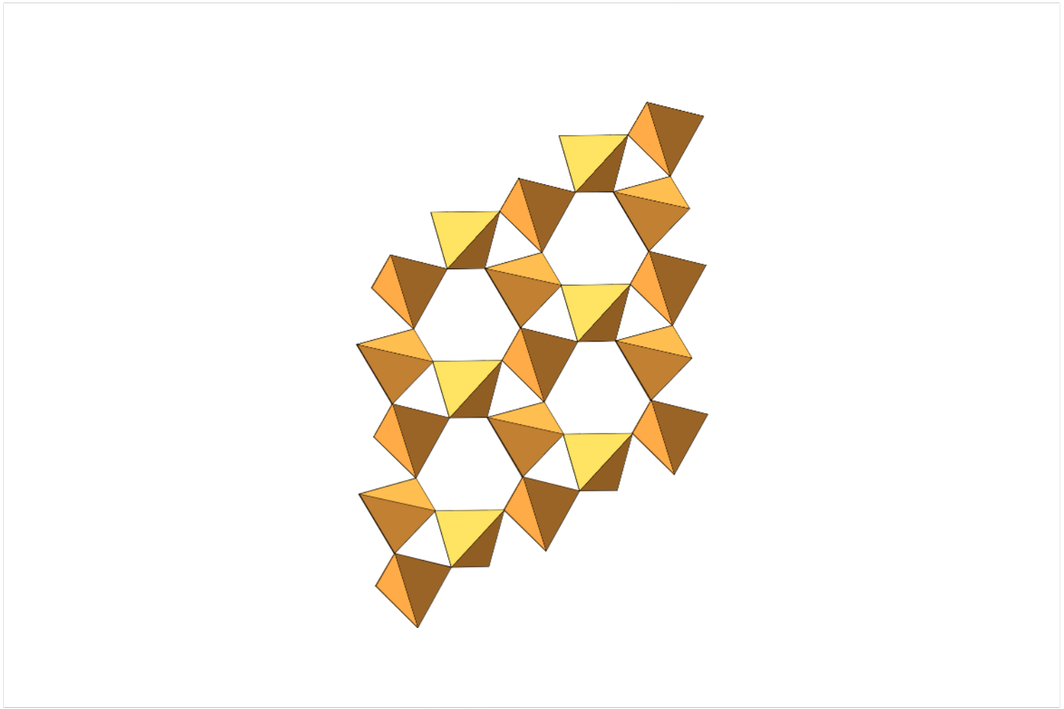
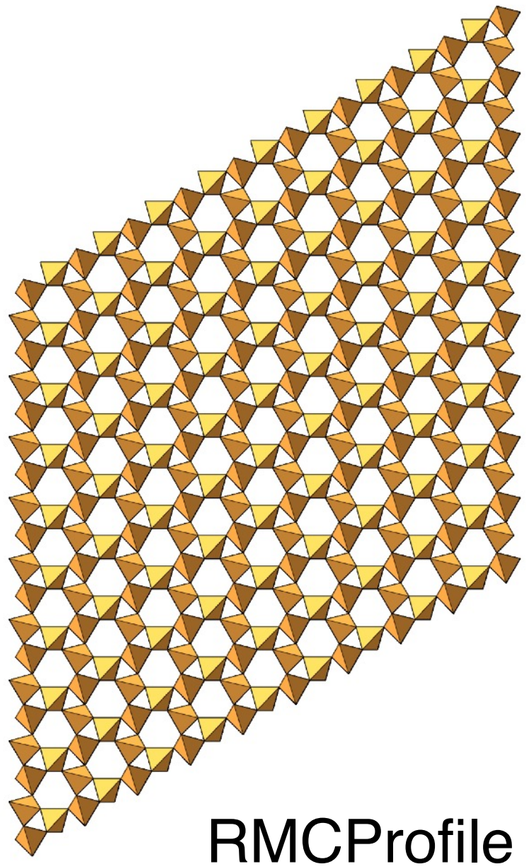
Big box vs small box models

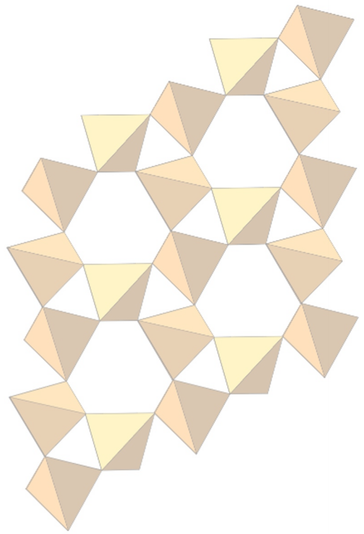


PDFgui
(r-space Reitveld)

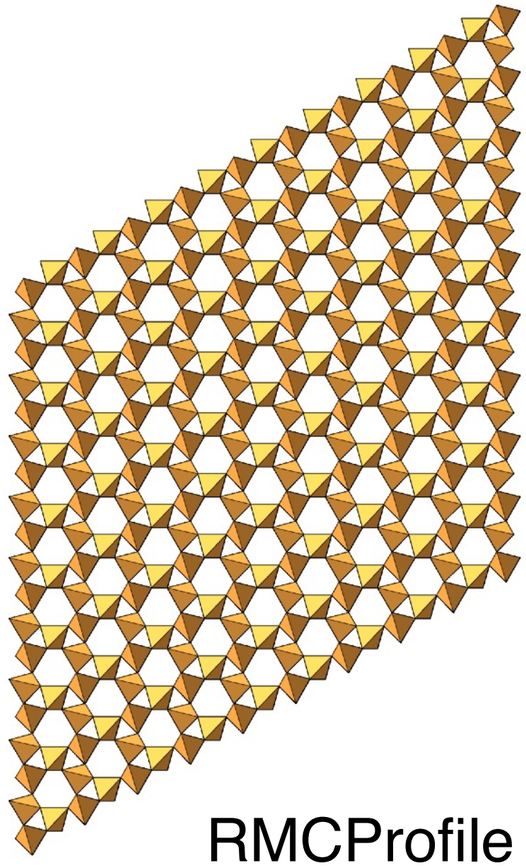


RMCProfile
(Reverse Monte Carlo)

Big box models



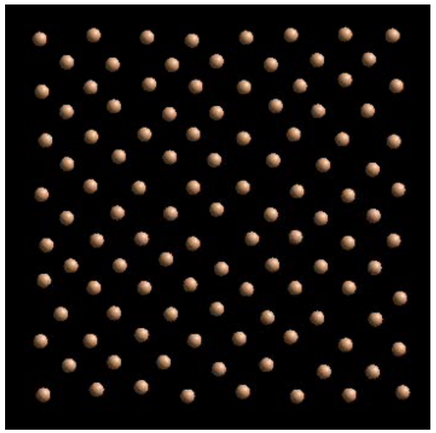
PDFgui
(r-space Reitveld)



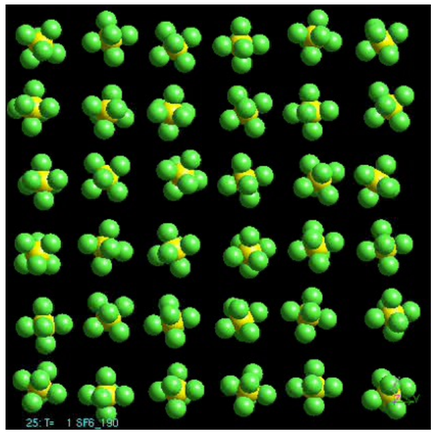
RMCProfile
(Reverse Monte Carlo)

Disordered materials

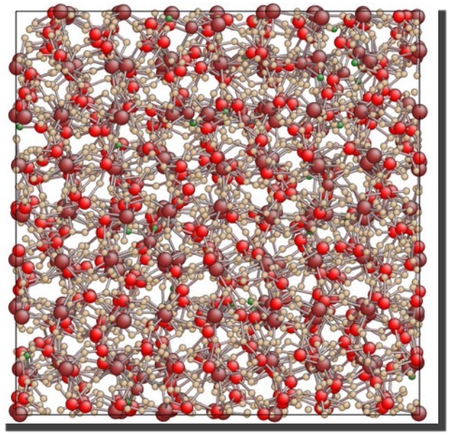
Simple crystals



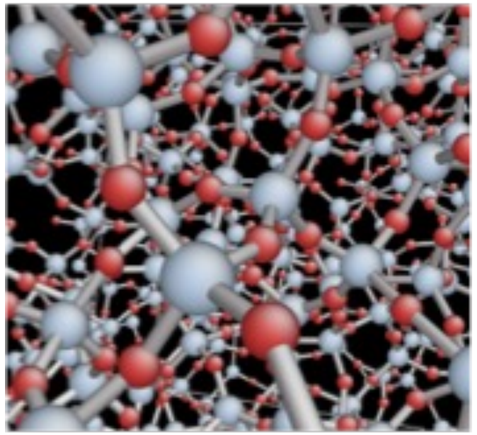
Disordered crystals



Amorphisation



Amorphous



RMCProfile

The RMC Method

Reverse Monte Carlo Simulation:
a new technique for the determination of disordered structures

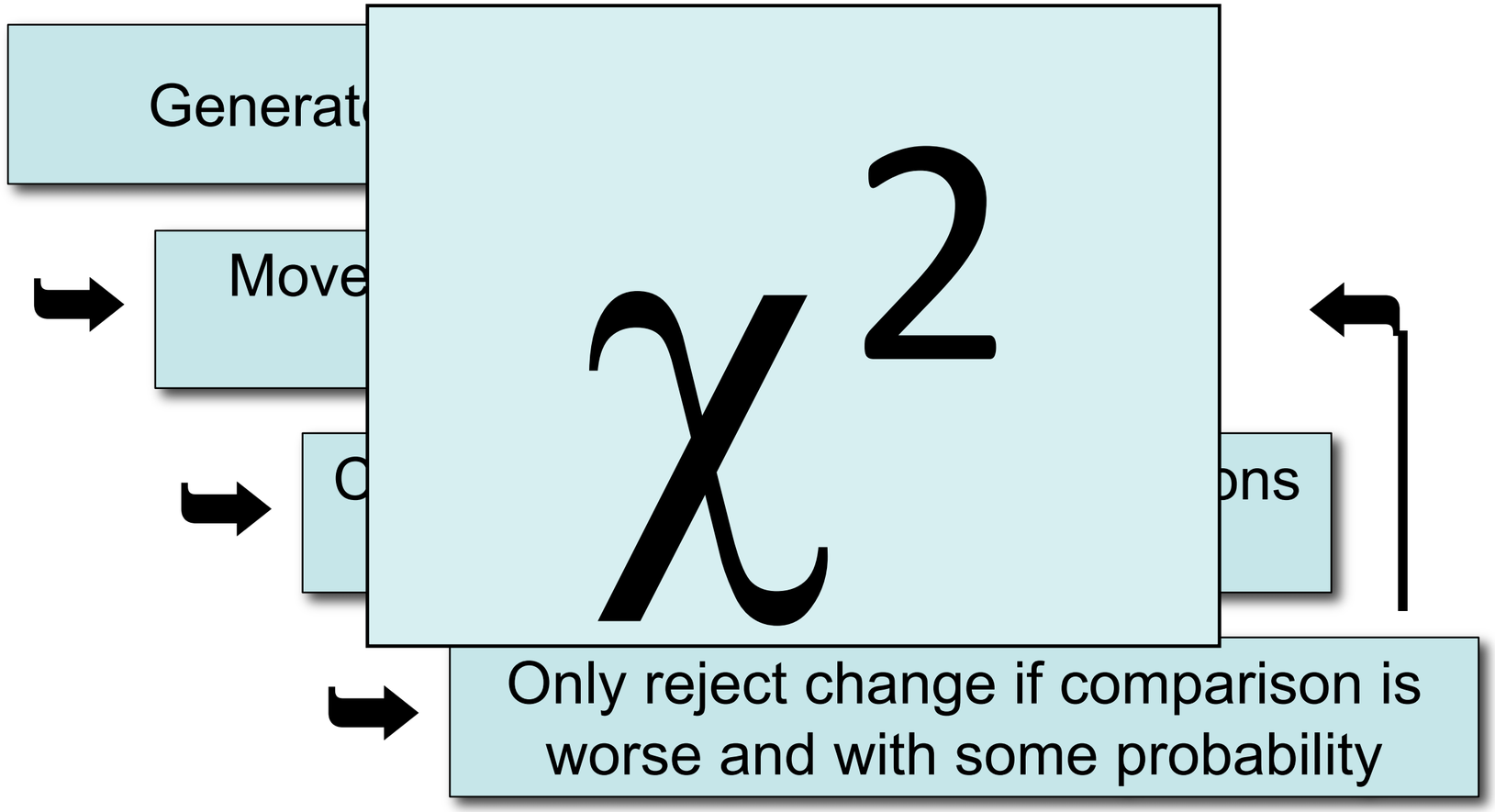
McGreevy R L and Pusztai L, *Molecular Simulation* **1**(1988) 359

We have developed a new technique, based on the standard Monte Carlo simulation method with Markov chain sampling, where a set of three dimensional particle configurations are generated that are consistent with the experimentally measured structure factor, A(Q), and radial distribution function, g(r), of a liquid or other disordered system. Consistency is determined by a standard χ^2 test using the experimental errors. No input potential is required. We present initial results for liquid argon. Since the technique can work directly from the structure factor it promises to be extremely powerful for modelling the structures of glasses or amorphous materials. It also has many other advantages in multicomponent systems and as a tool for experimental data analysis.

Key words: Monte Carlo, structure factor, radial distribution function, liquid, glass.

PACS numbers: 02.50, 61.25, 61.40.

The Reverse Monte Carlo algorithm



RMC χ^2

$$\chi^2_{RMC} = \sum_j \chi^2_j$$

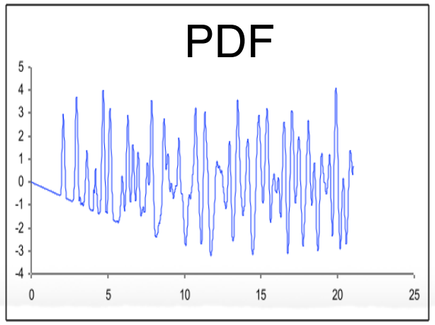
$$\chi^2_{F(Q)} = \sum_j \left[F_{calc}(Q_j) - F_{box}(Q_j)\right]^2 / \sigma^2_{F(Q)}(Q_j)$$ Total scattering

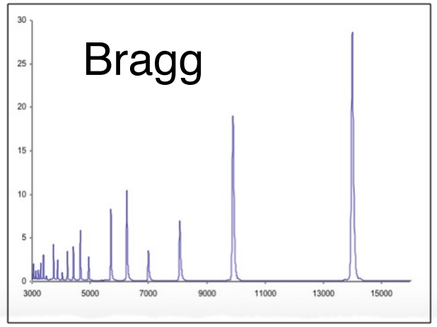
$$\chi^2_{G(r)} = \sum_j \left[G_{calc}(r_j) - G_{expt}(r_j)\right]^2 / \sigma^2_{G(r)}(r_j)$$ PDF

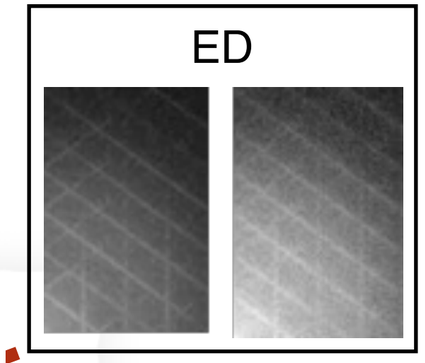
$$\chi^2_{profile} = \sum_j \left[I_{calc}(t_j) - s' I_{expt}(t_j)\right]^2 / \sigma^2_{I(i)}(t_j)$$ Bragg profile

$$\chi^2_f = \sum_k w_k \left[f_k^{calc} - f_k^{req}\right]^2$$ Constraints/restraints

RMC PROFILE







PRL 96, 047209 (2006) PHYSICAL REVIEW LETTERS week ending 3 FEBRUARY 2006

Magnetic Structure of MnO at 10 K from Total Neutron Scattering Data

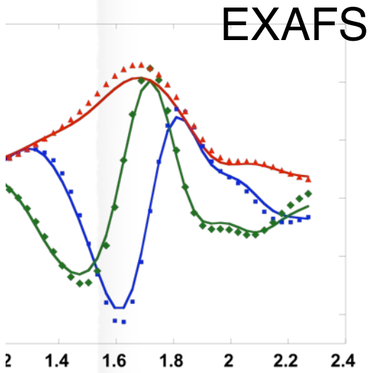
Andrew L. Goodwin,[1] Matthew G. Tucker,[1] Martin T. Dove,[1,*] and David A. Keen[2,3]

[1]*Department of Earth Sciences, Cambridge University, Downing Street, Cambridge CB2 3EQ, United Kingdom*
[2]*Department of Physics, Oxford University, Clarendon Laboratory, Parks Road, Oxford OX1 3PU, United Kingdom*
[3]*ISIS Facility, Rutherford Appleton Laboratory, Chilton, Didcot, Oxfordshire OX11 0QX, United Kingdom*

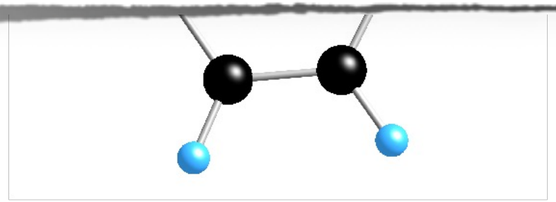
(Received 24 October 2005; published 2 February 2006)

Total neutron scattering data from a powdered sample of MnO collected at 10 K have been analyzed using the reverse Monte Carlo method to refine the nuclear and magnetic structure. The results give the first unambiguous assignment of the average magnetic structure. The magnetic moments are aligned ferromagnetically within (111) sheets with the magnetization vectors of alternate sheets along axes parallel and antiparallel to the $\langle 11\bar{2} \rangle$ directions, albeit with a small modulated out-of-plane component. Small displacements of Mn and O (modulated with the same periodicity) accompany the magnetic ordering and both atomic and magnetic structures may be described in the monoclinic space group $C2$.

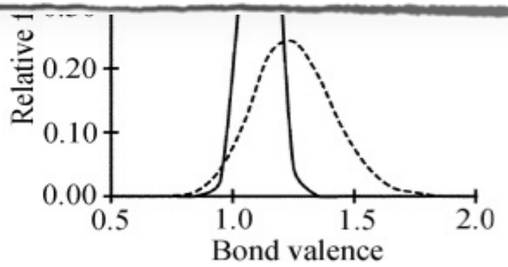
DOI: 10.1103/PhysRevLett.96.047209 PACS numbers: 75.25.+z, 02.70.Uu, 61.12.Ex

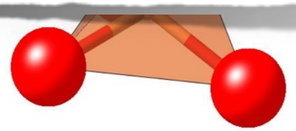


Polyhedral restraints



Molecular potentials





M G Tucker et al, *J. Phys.: Condens. Matter* **19,** 335218 (2007).

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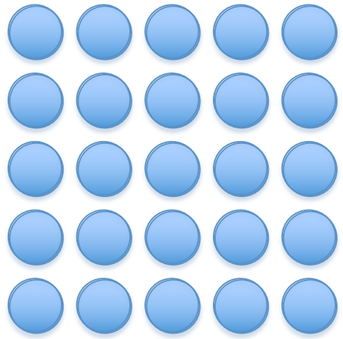
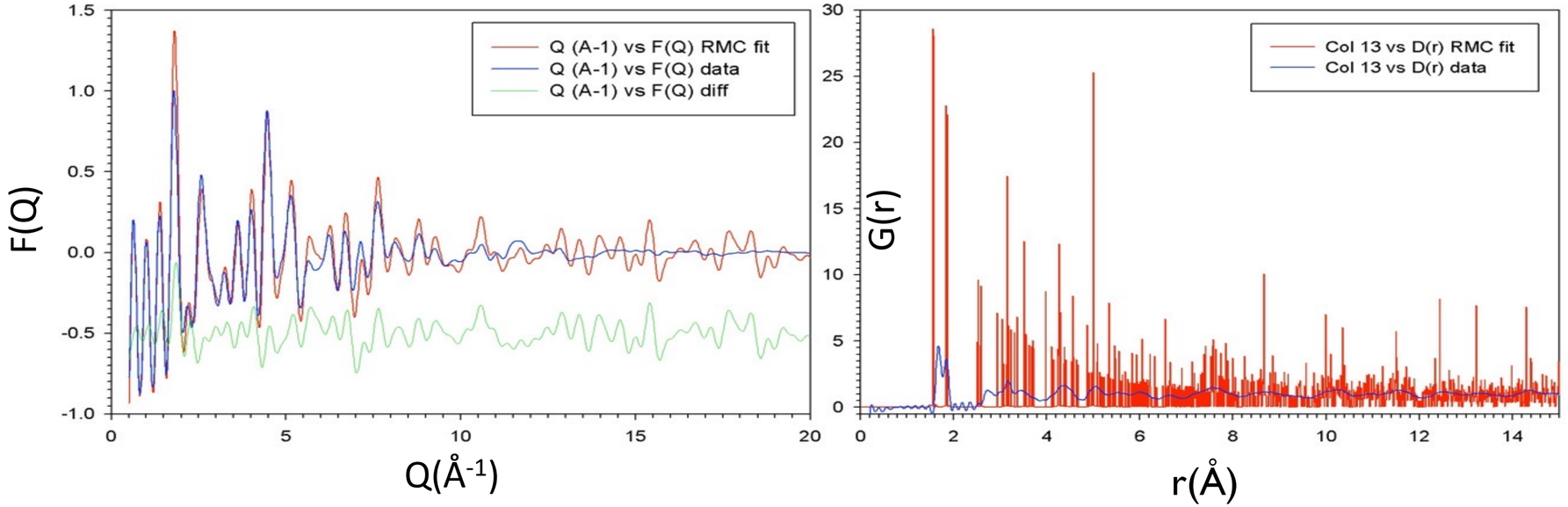
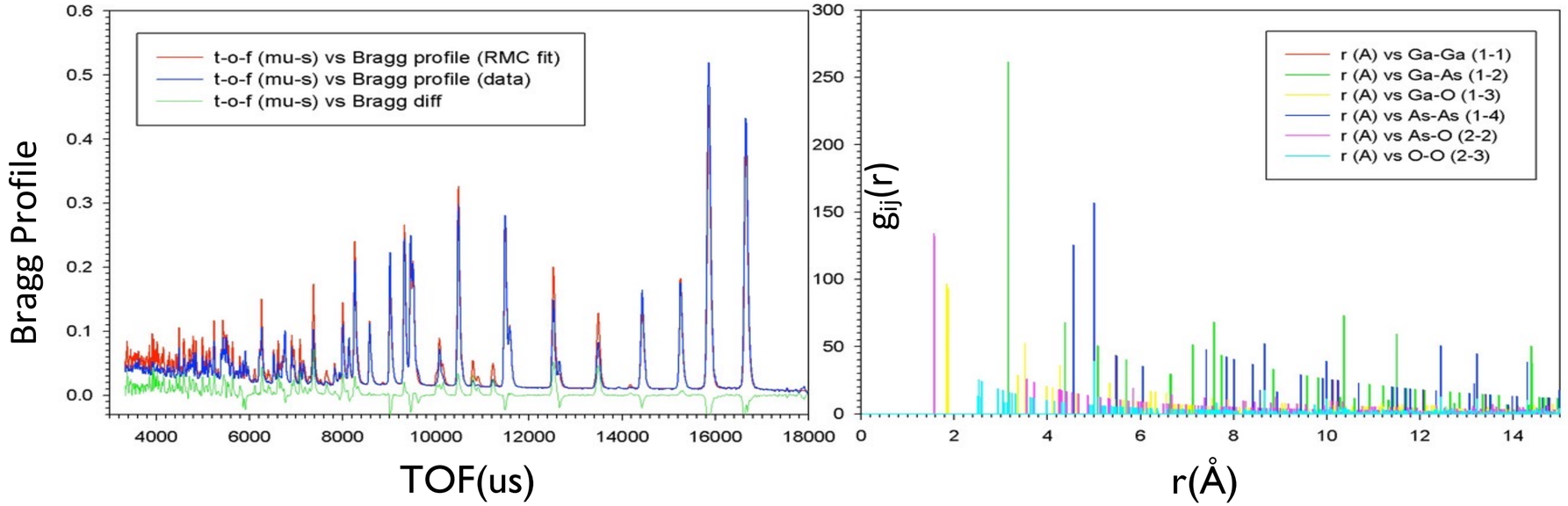
RMCProfile refinement



GaAsO4 RT.files





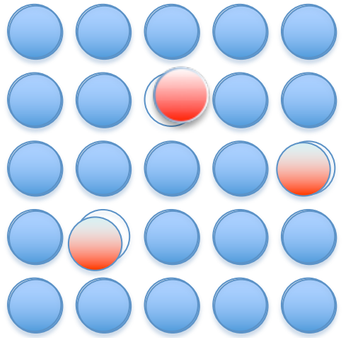
4608 atoms

0 moves

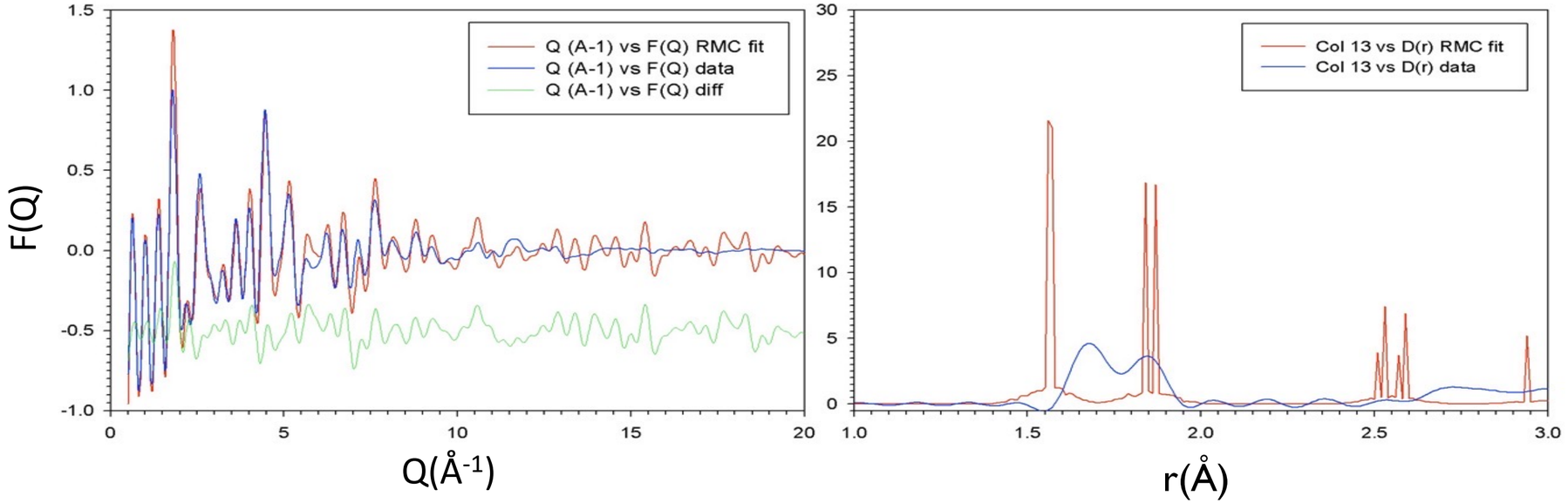
$\chi^2 = 3023$

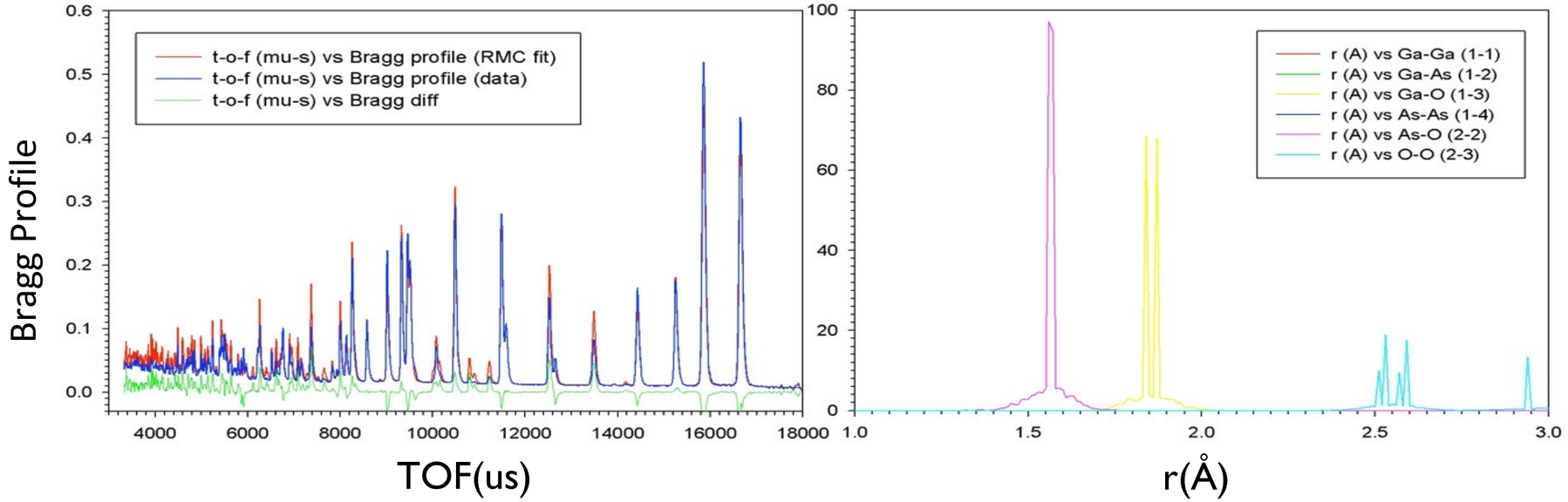
Slides from
Dave Keen

RMCProfile refinement



GaAsO4 RT.files





4608 atoms

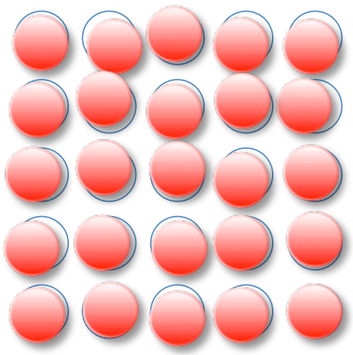
834 moves

$\chi^2 = 1629$

Slides from
Dave Keen

OAK RIDGE National Laboratory

RMCProfile refinement

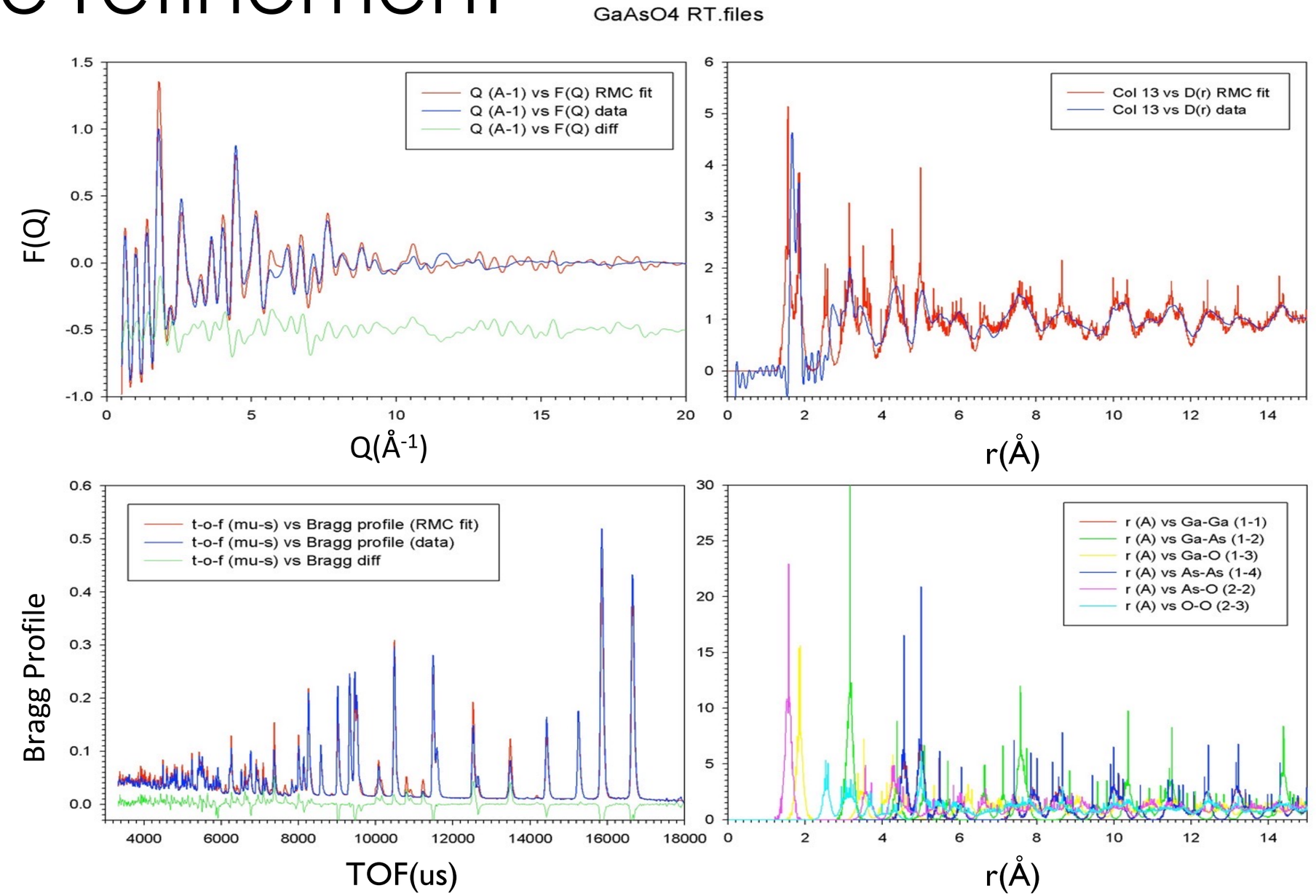


4608 atoms

4699 moves

$\chi^2 = 92.3$

Slides from
Dave Keen



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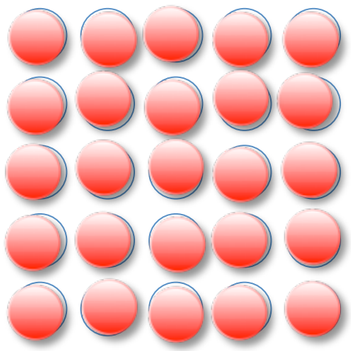
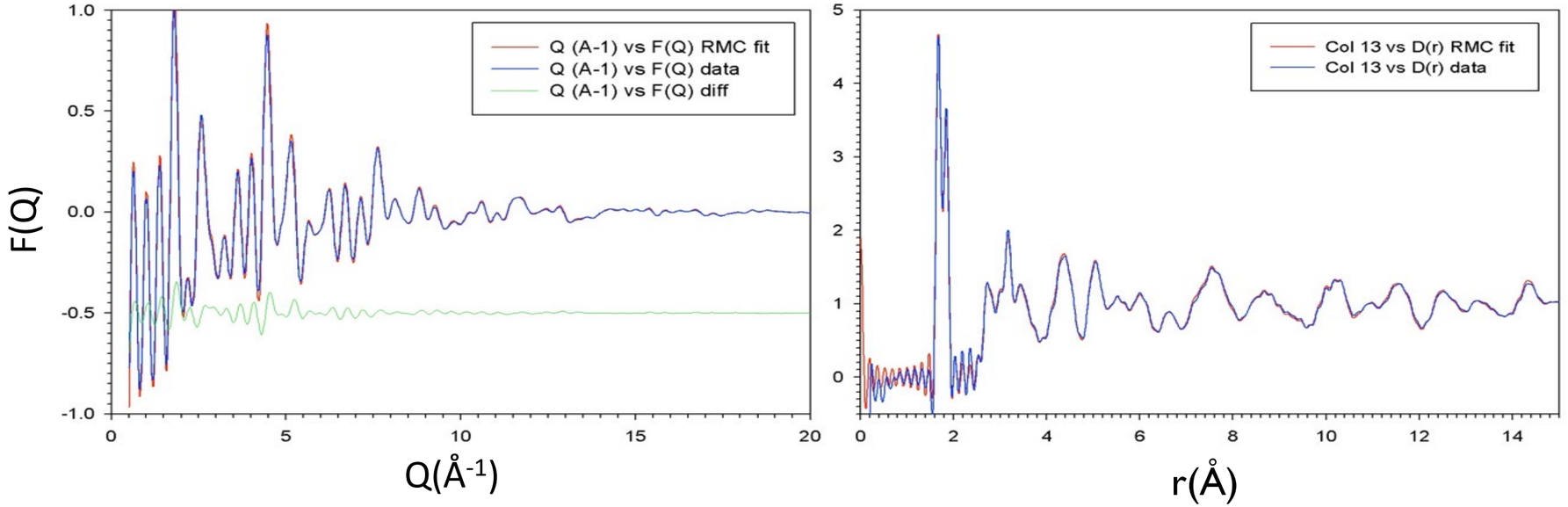
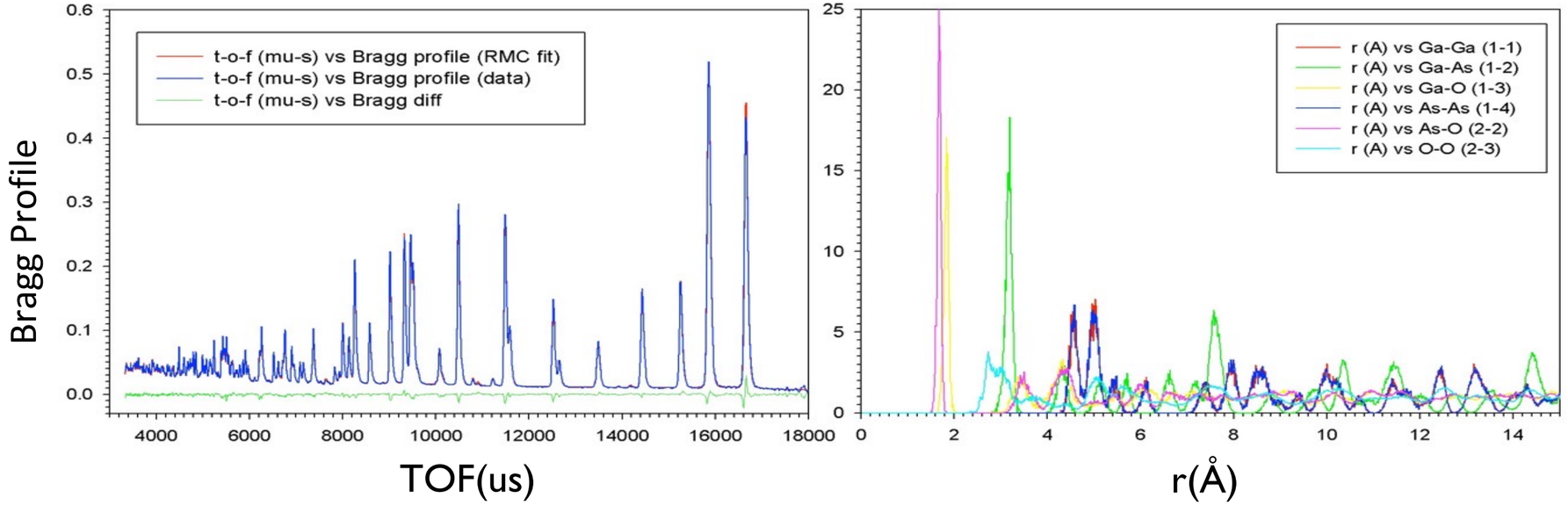
RMCProfile refinement



GaAsO4 RT.files





4608 atoms

RMC converged

$\chi^2 = 3.43$

Slides from
Dave Keen

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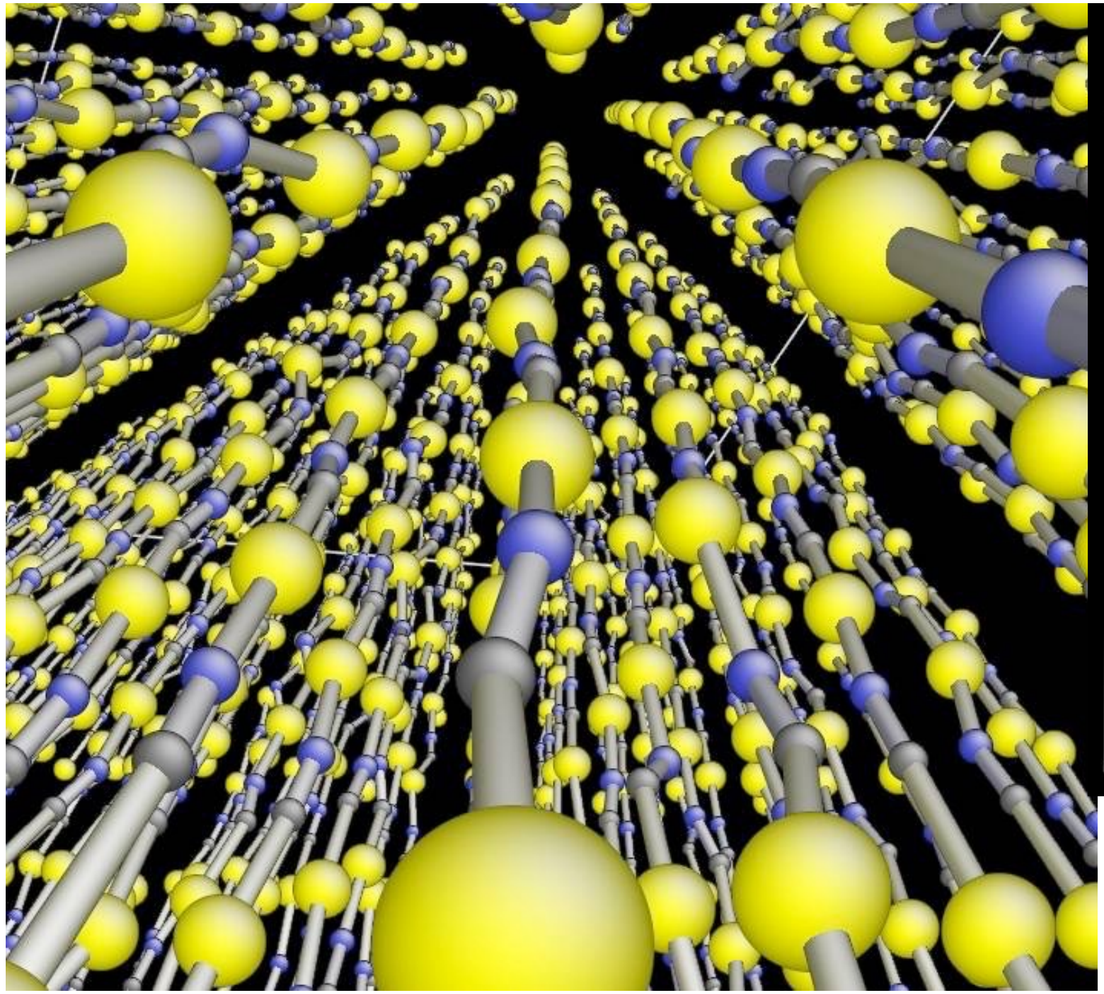
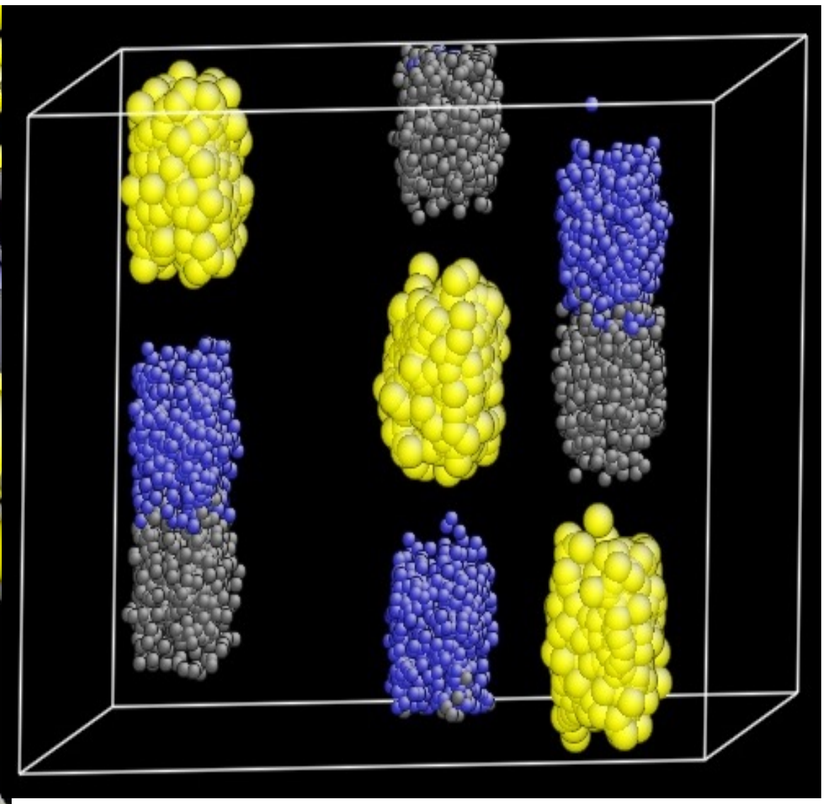
Big box models





Collapsed to unit cell

Introduction to the RMCProfile7 program

Atomic configuration

- usually a supercell of the crystallographic unit cell
- 10000 – 50000 atoms

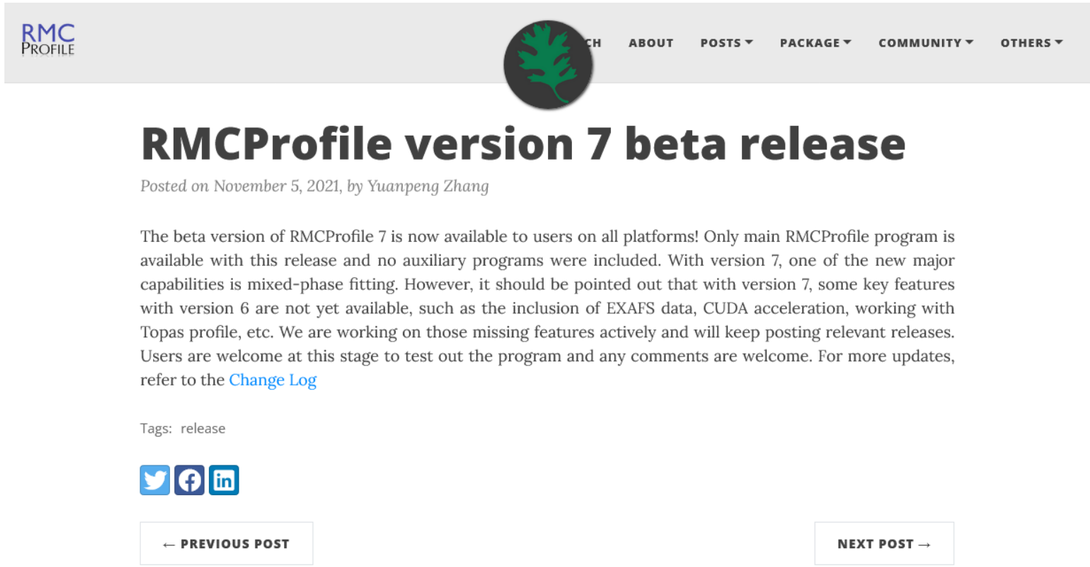
RMCProfile can fit

- real space data , i.e. G(r), D(r), T(r)
- reciprocal space data, i.e. F(Q)
- Bragg data (using GSAS output)
- magnetic scattering in reciprocal space
- EXAFS (k- and/or r-space)
- (Electron) single crystal diffuse

Variety of additional constraints

- minimum distance “hard sphere Monte Carlo”
- distance window
- molecular potentials (distance and angle)
- polyhedral constraint
- coordination number

https://rmcprofile.pages.ornl.gov



OAK RIDGE National Laboratory

Introduction to the RMCProfile7 program

WHAT'S NEW!

Multiphase refinement

- calculation of all dataset types can be done for multiple phases*

RMCProfile can fit

- real space dataset (X-Ray and neutron) can be calculated as a back Fourier Transform of reciprocal space dataset
- full GSAS-II compatibility (easy Bragg data extraction)

Variety of additional constraints

- molecular potentials (distances, angles, torsion angle, inversion angle, planarity + variants)
- potentials and swaps are now compatible

Novel move type

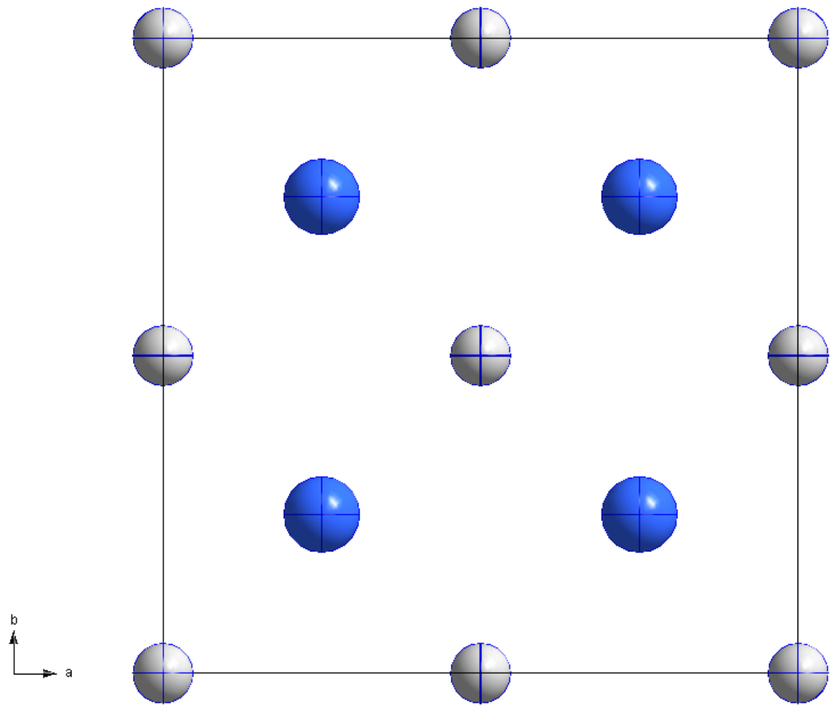
- molecule (rigid body) type move
- swap between atoms and atoms, atoms to molecules and molecules to molecules

*W. Sławiński, J. Appl. Cryst. (2018). **51**, 919–923

RMCProfile7 – multiple phases

WHAT'S NEW!

CaF_2 and CeO_2 with 50%/50% mass fraction

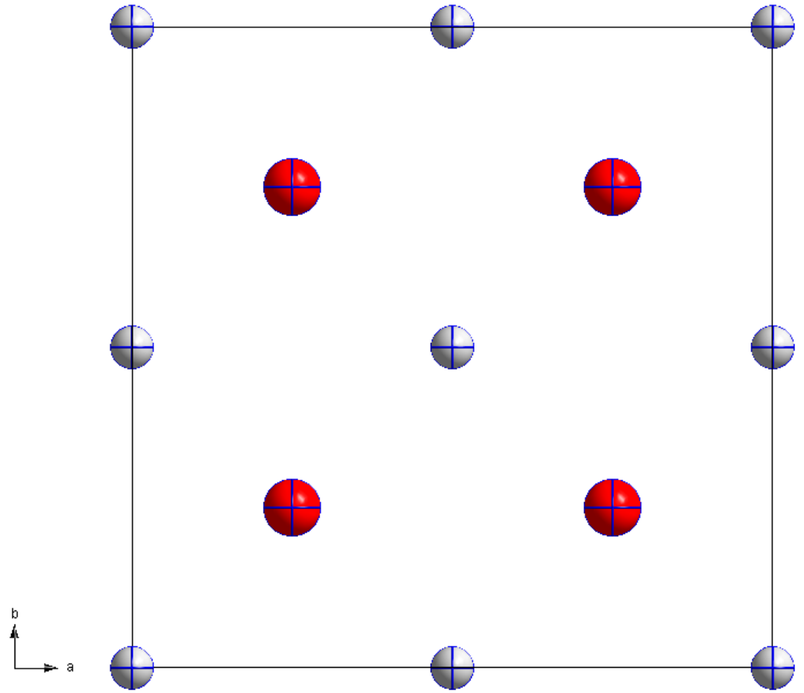


General	
Origin	50_50_CAF2_CEO2

Bibliographic data

Phase data	
Space-group	F m -3 m (225) - cubic
Cell	a=5.4625(0) Å V=162.99(0) Å^3 Z=4

Atomic parameters						
Atom	Wyck.	Site	x/a	y/b	z/c	U [Å^2]
Ca1	4a	m-3m	0	0	0	0.0058(1)
F1	8c	-43m	1/4	1/4	1/4	0.0092(0)



General	
Origin	50_50_CAF2_CEO2

Bibliographic data

Phase data	
Space-group	F m -3 m (225) - cubic
Cell	a=5.4076(0) Å V=158.13(0) Å^3 Z=4

Atomic parameters						
Atom	Wyck.	Site	x/a	y/b	z/c	U [Å^2]
Ce1_2	4a	m-3m	0	0	0	0.0028(1)
O1_2	8c	-43m	1/4	1/4	1/4	0.0050(1)

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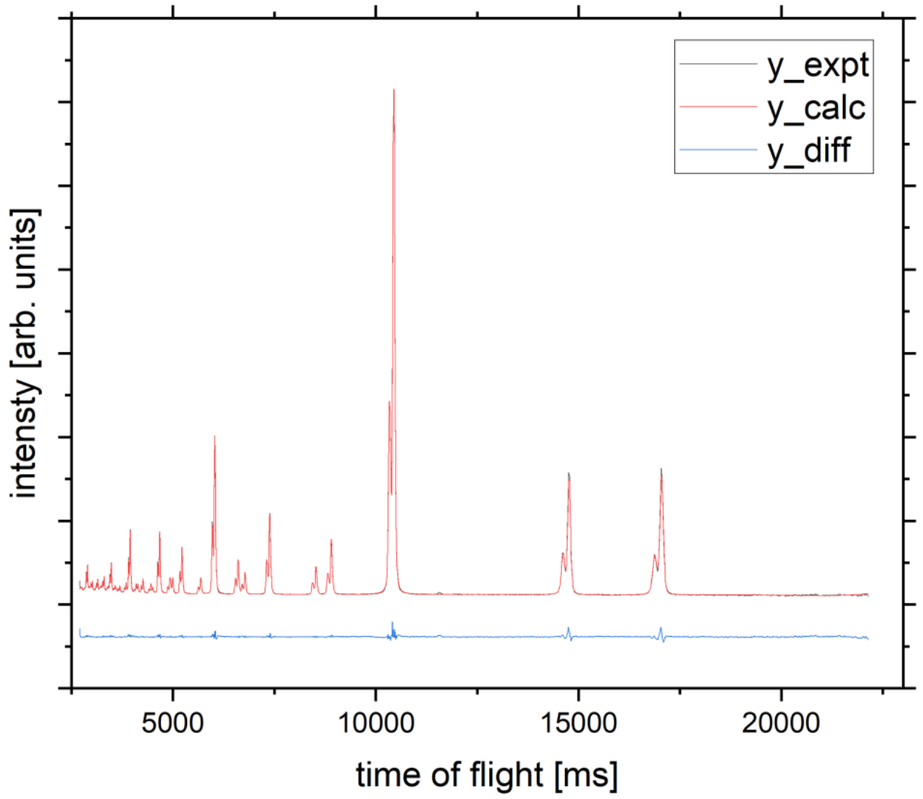
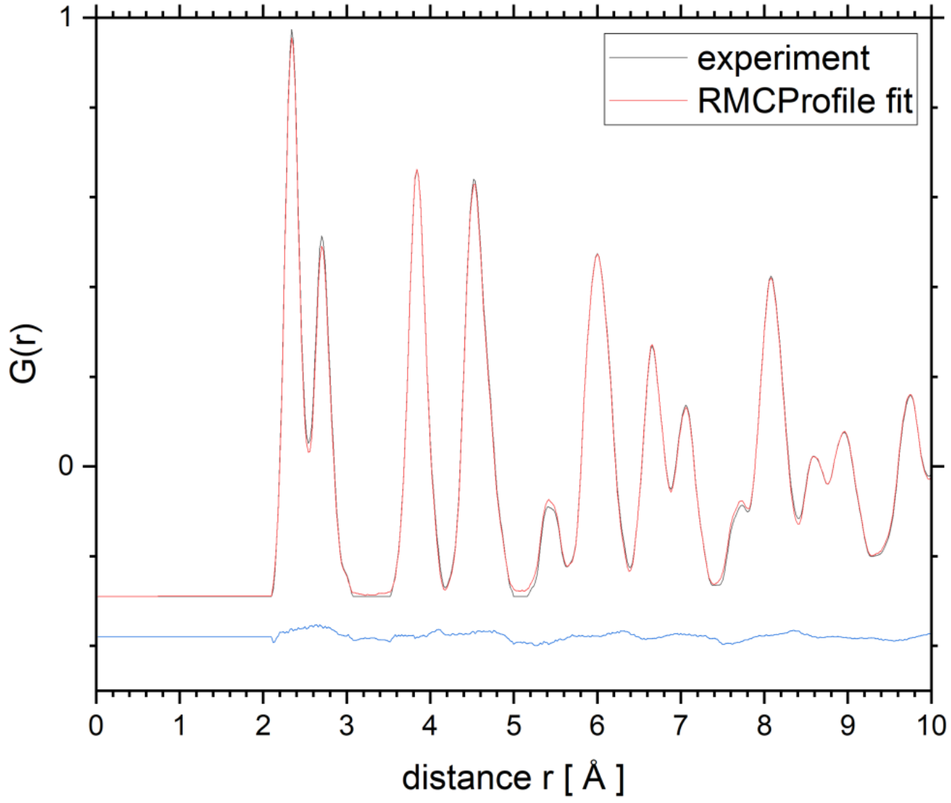
ADD2022

RMCProfile7 – multiple phases

WHAT'S NEW!

CaF_2 and CeO_2 with 50%/50% mass fraction (69.5%/30.5% mole fraction)



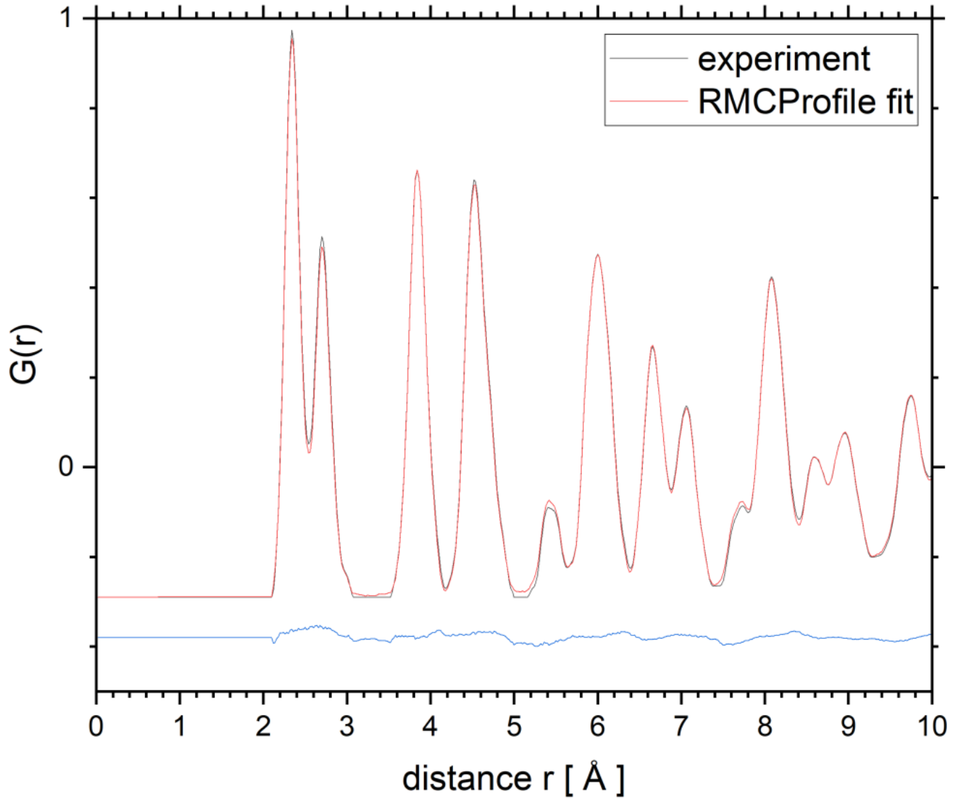


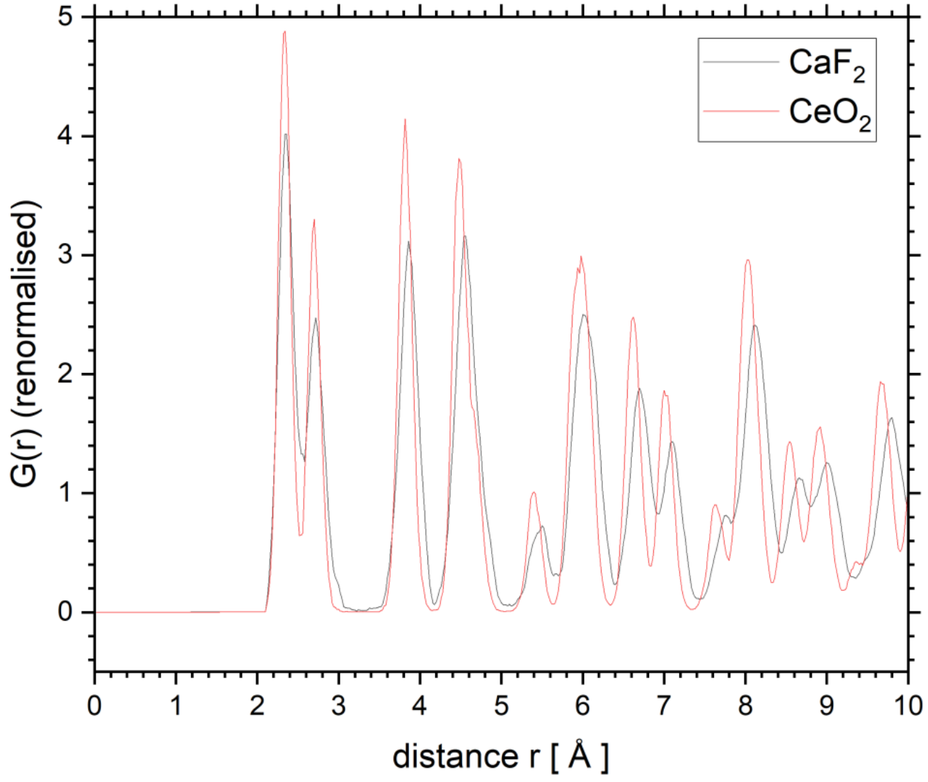
data collected POLARIS @ ISIS Neutron and Muon Source, UK

RMCProfile7 – multiple phases

WHAT'S NEW!

CaF_2 and CeO_2 with 50%/50% mass fraction (69.5%/30.5% mole fraction)

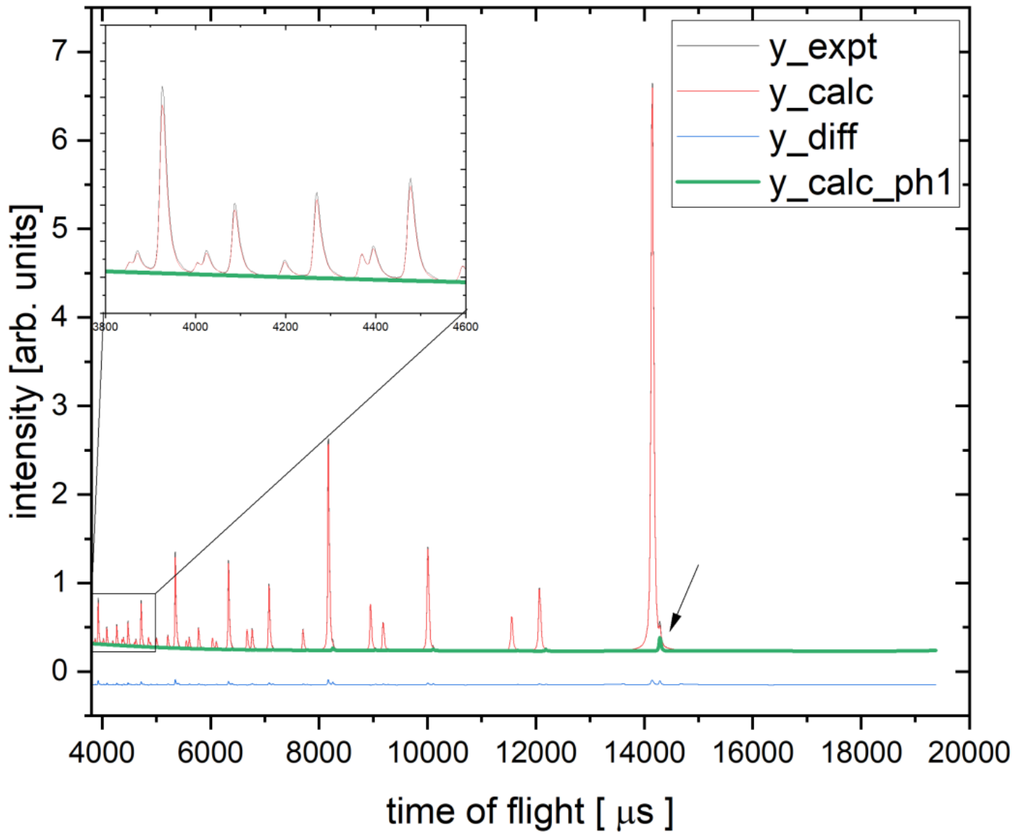


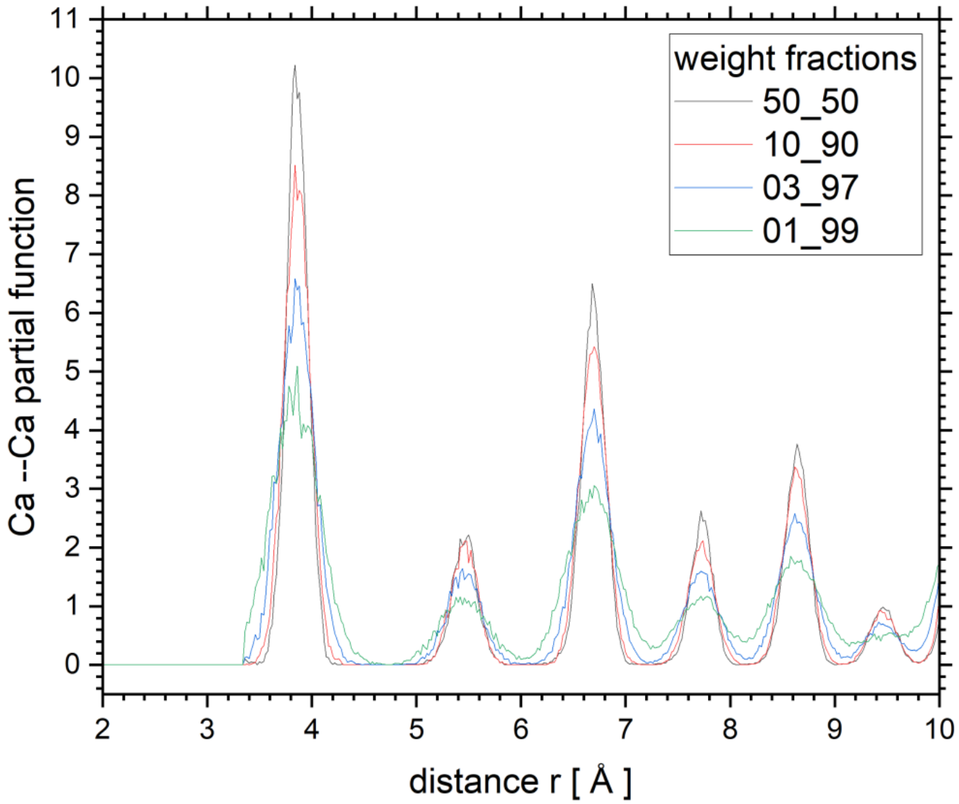


RMCProfile7 – multiple phases

BE AWARE OF

CaF_2 and CeO_2 with different mass fractions





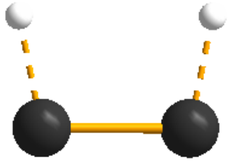
RMCProfile7 – potentials

WHAT'S NEW!

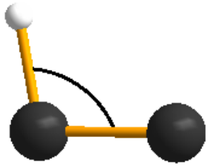
RMCProfile7 can search for collections of atoms which will be restrained by potentials

BOND

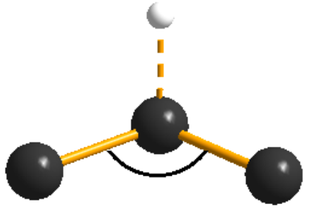
BOND-CENTRAL



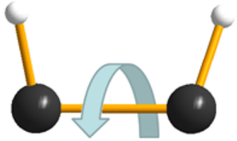
ANGLE



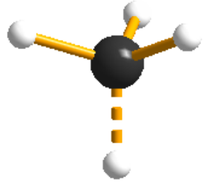
ANGLE-CENTRAL



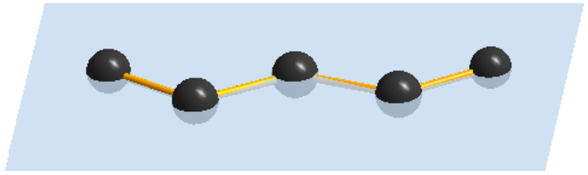
DIHEDRAL ANGLE



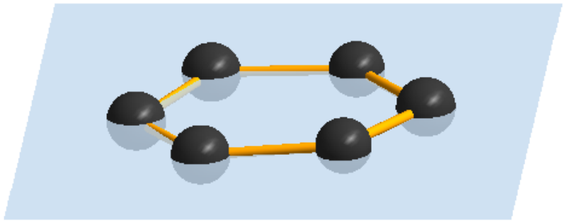
INVERSION ANGLE



PLANAR



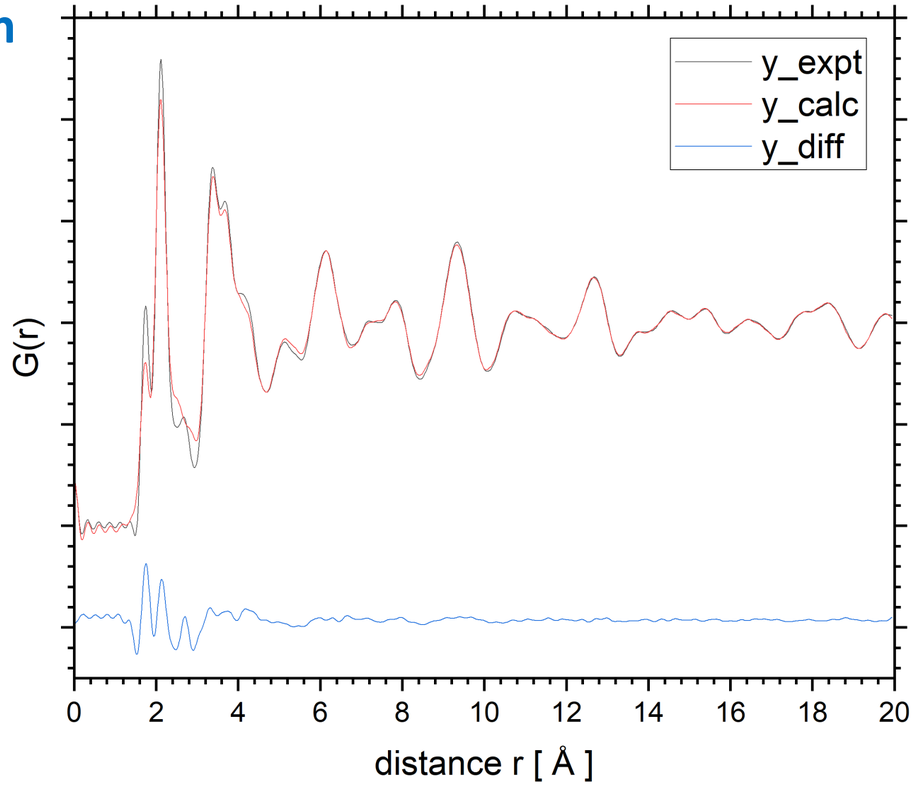
PLANAR RING

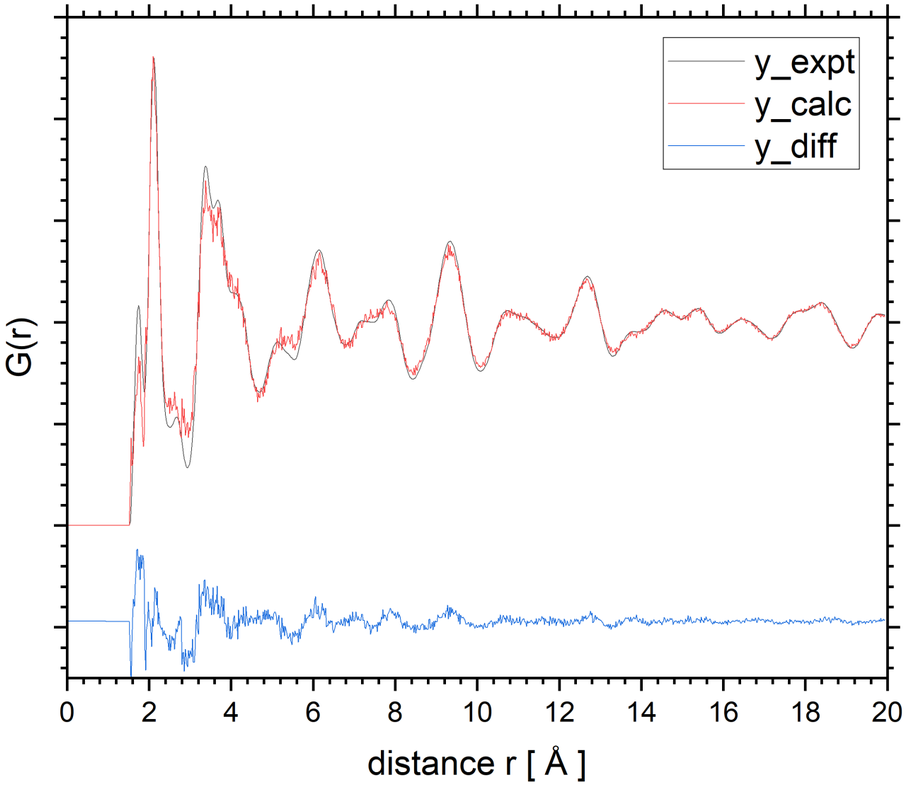


RMCProfile7 – real space data calculation

WHAT'S NEW!

RMCProfile7 can calculate REAL SPACE data as a back Fourier Transform of F(Q) data





Might be usefull for

- **X-Ray PDF data due to the nature of scattering factor f(Q) dependence**
- **Neutron data measured with limited Q_{max}**

RMCProfile7 – molecule type move

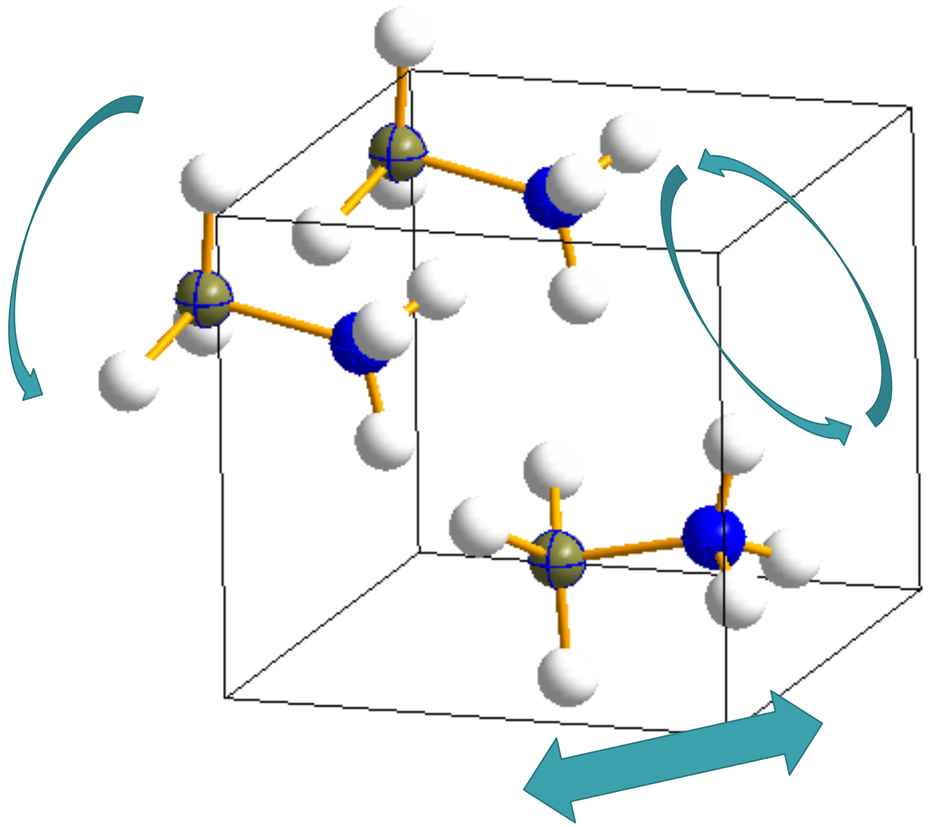
WHAT'S NEW!

RMCProfile7 can:

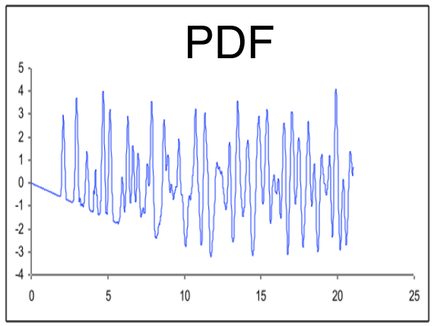
- **translate,**
- **rotate,**
- **swap**

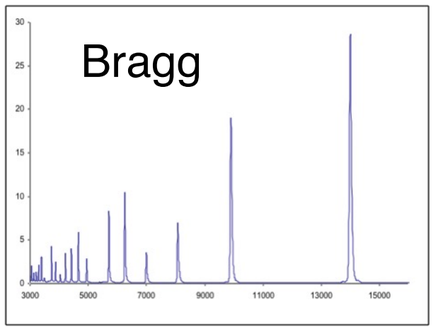
molecules (rigid bodies)

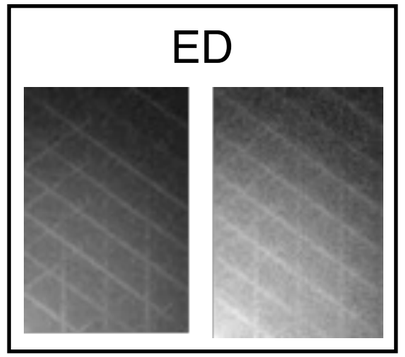
but also swap atoms with moleculed

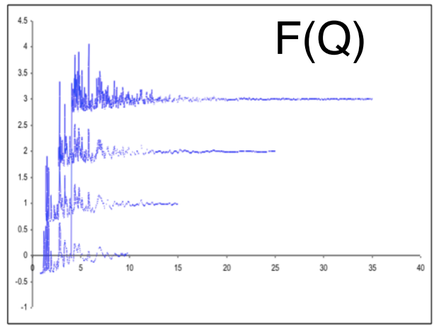


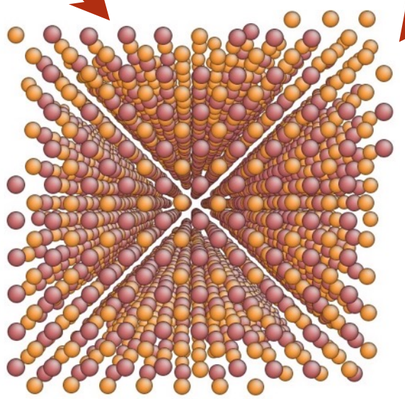
RMC Profile

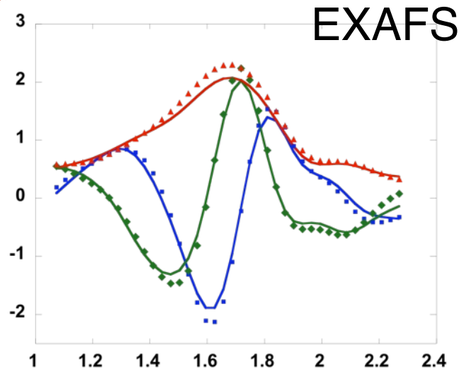


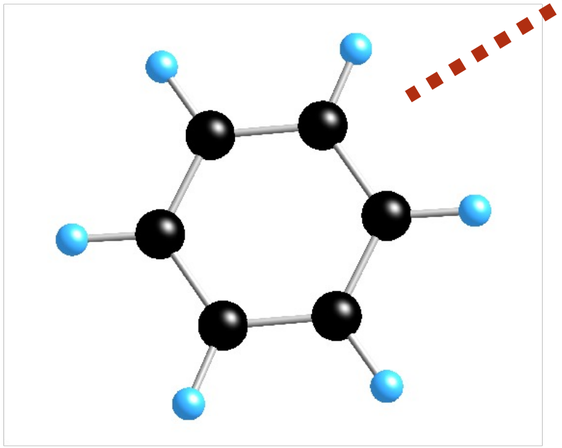




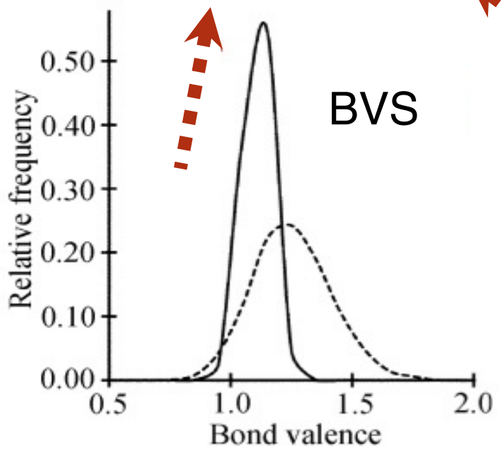


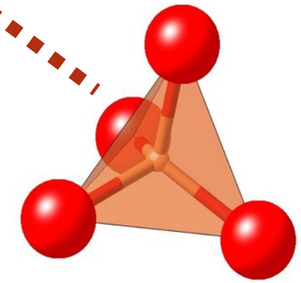






Molecular potentials





Polyhedral restraints

M G Tucker et al, *J. Phys.: Condens. Matter* **19,** 335218 (2007).

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Yuanpeng Zhang (ORNL)

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Thank you!

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