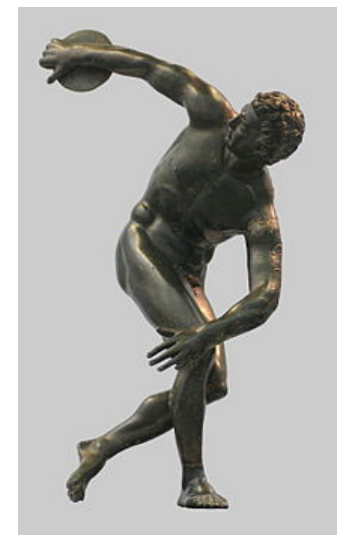




DISCUS,
Simulation and refinement of disordered
crystal structures

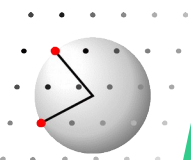
DIffuse **SC**attering ~~And~~ **S**tructure simulation
U_{nd}



DISCUS Workshop 2018

Reinhard B. Neder
Kristallographie und Strukturphysik
Friedrich-Alexander-University Erlangen-Nürnberg

reinhard.neder@fau.de

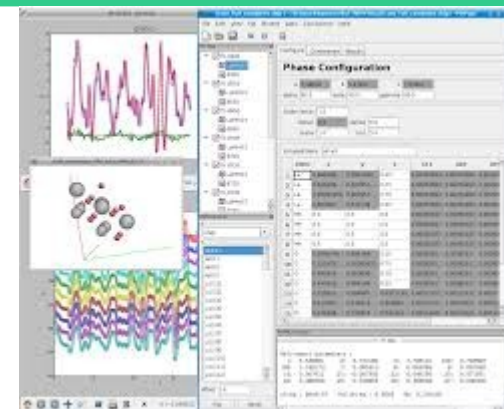
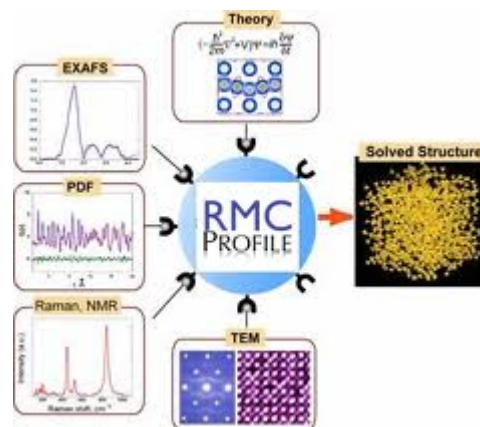


Small Box Modelling



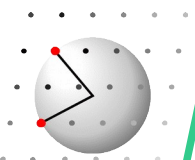
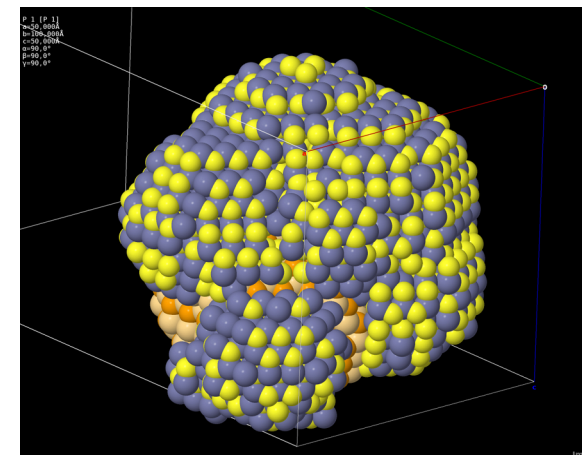
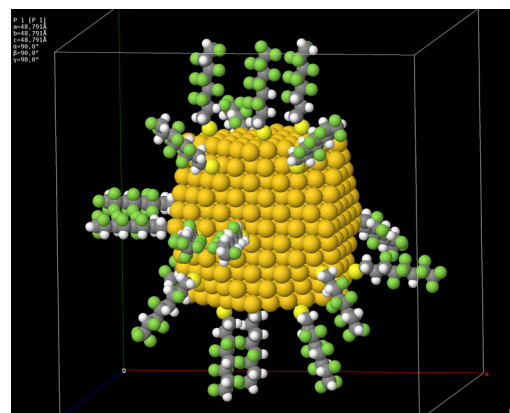
Large Box Modelling

Reverse Monte Carlo
RMCprofile
RMC_POT++
DISCUS



Bottom Up Modelling

DISCUS



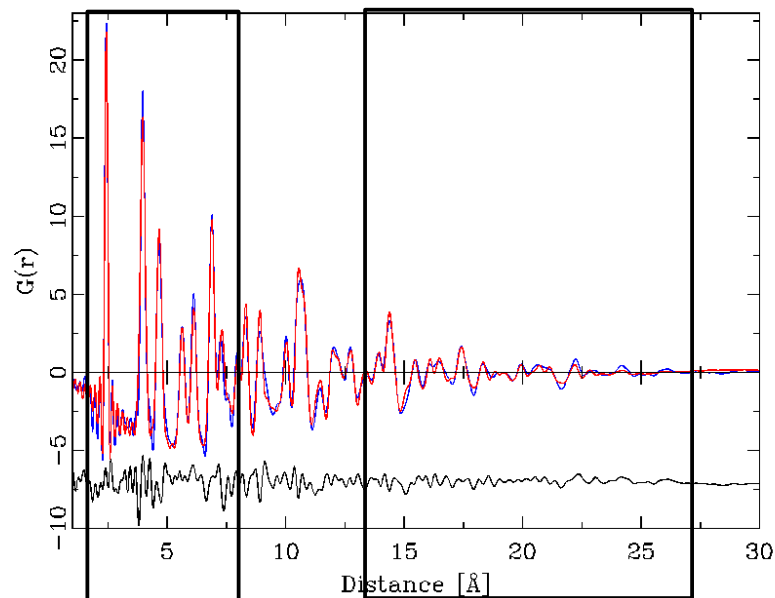
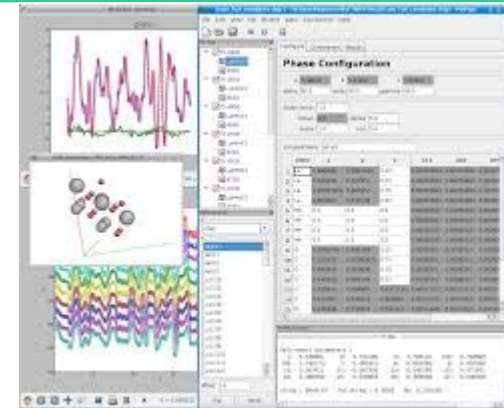
Small Box Modelling



Rietveld type refinement

Calculate PDF from single unit cell or **small** block of cells

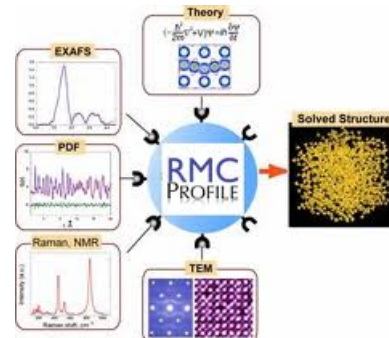
Determine changes in local versus moderate structure by
Separate refinement in restricted r-range



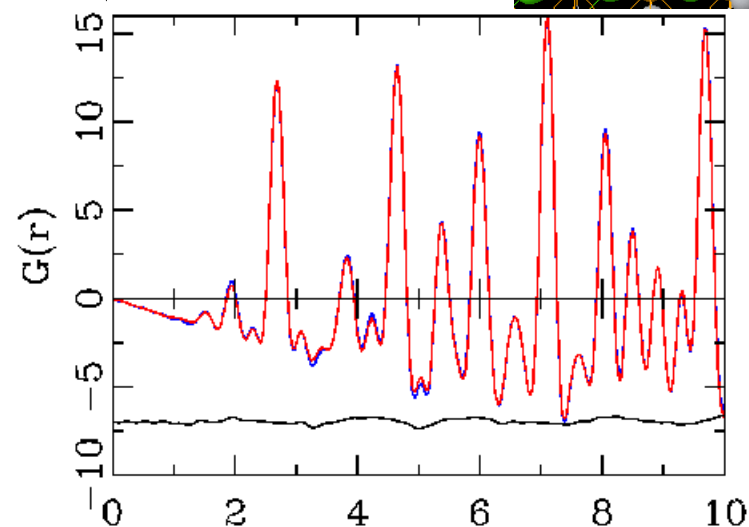
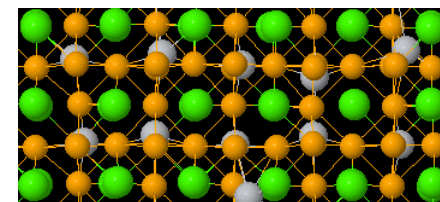
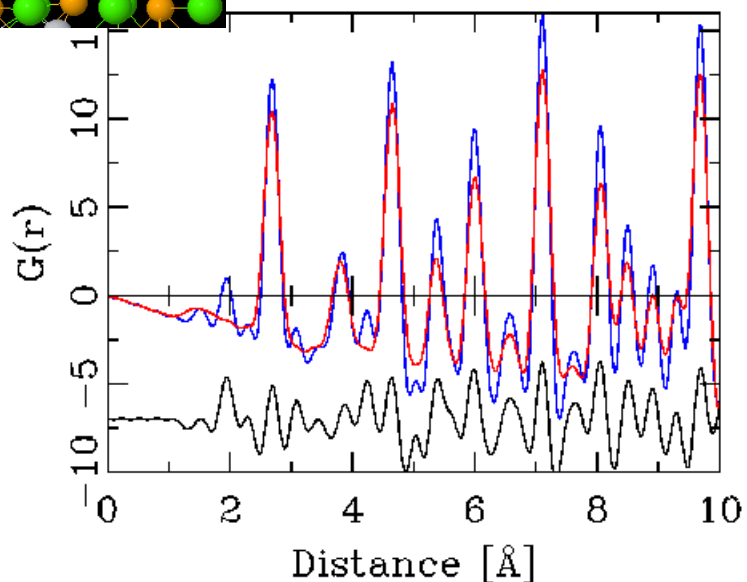
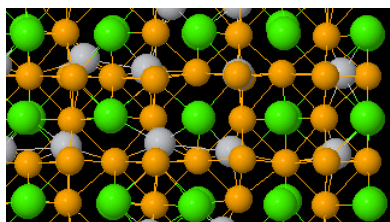
Large Box Modelling

Reverse Monte Carlo
RMCprofile
RMC_POT++

Model initial (random) **large** structure
Calculate PDF (+ powder + ...)



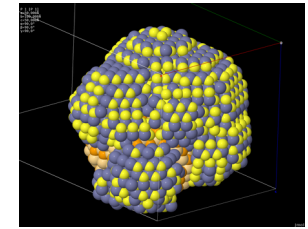
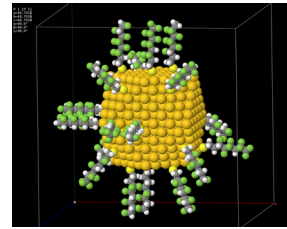
Modify structure (shift, exchange...)
while agreement improves
Usually requires constraints!



Analyze structure **afterwards**
Local coordination, shifts, etc.

Bottom Up Modelling

DISCUS



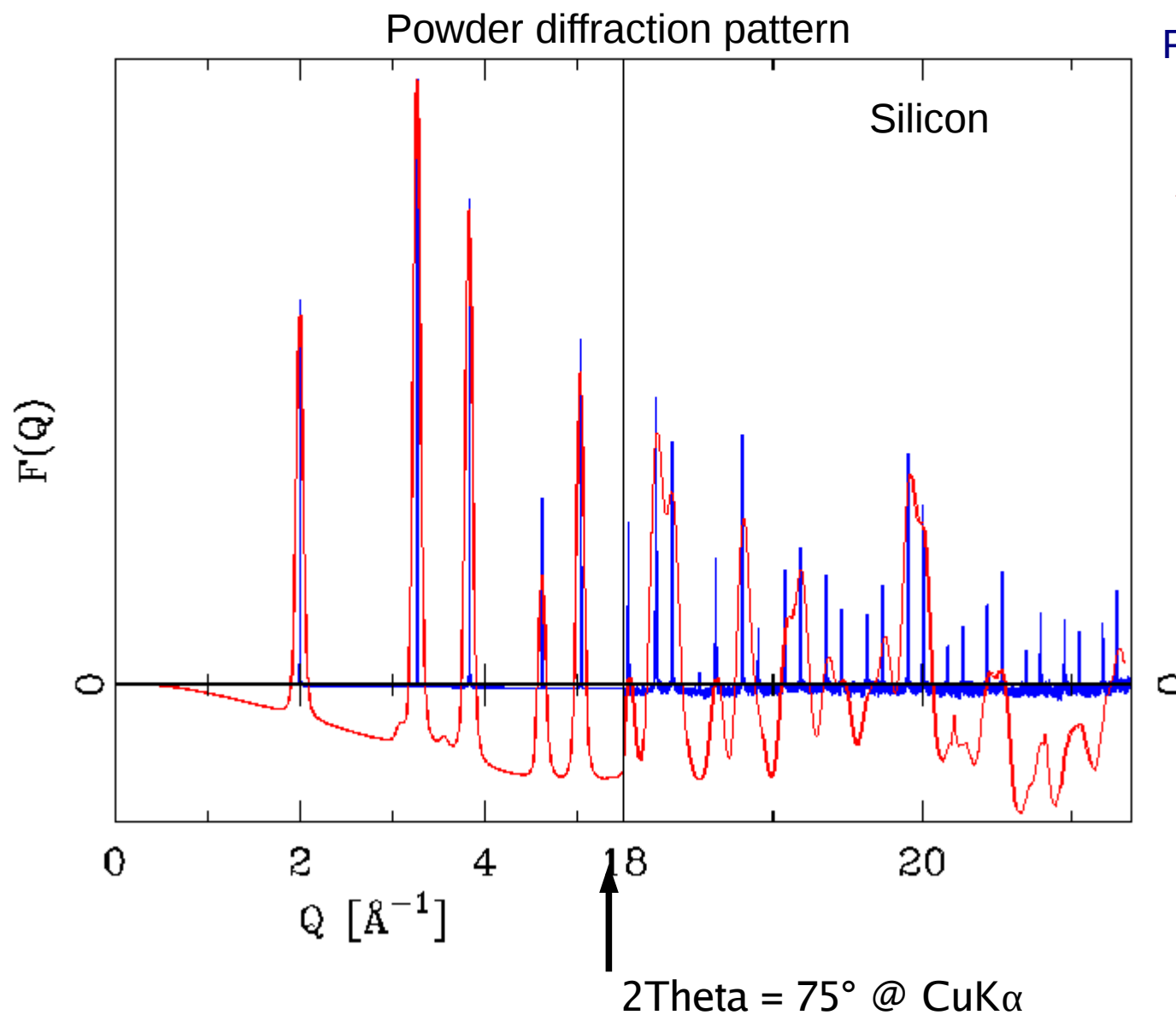
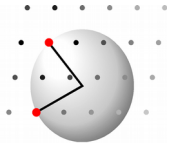
Define disorder rules to simulate complete structure

Non-periodic boundary conditions

Extended defect toolboxes

DISCUS program: tproffen.github.io/DiffuseCode/; Neder & Proffen (Oxford)

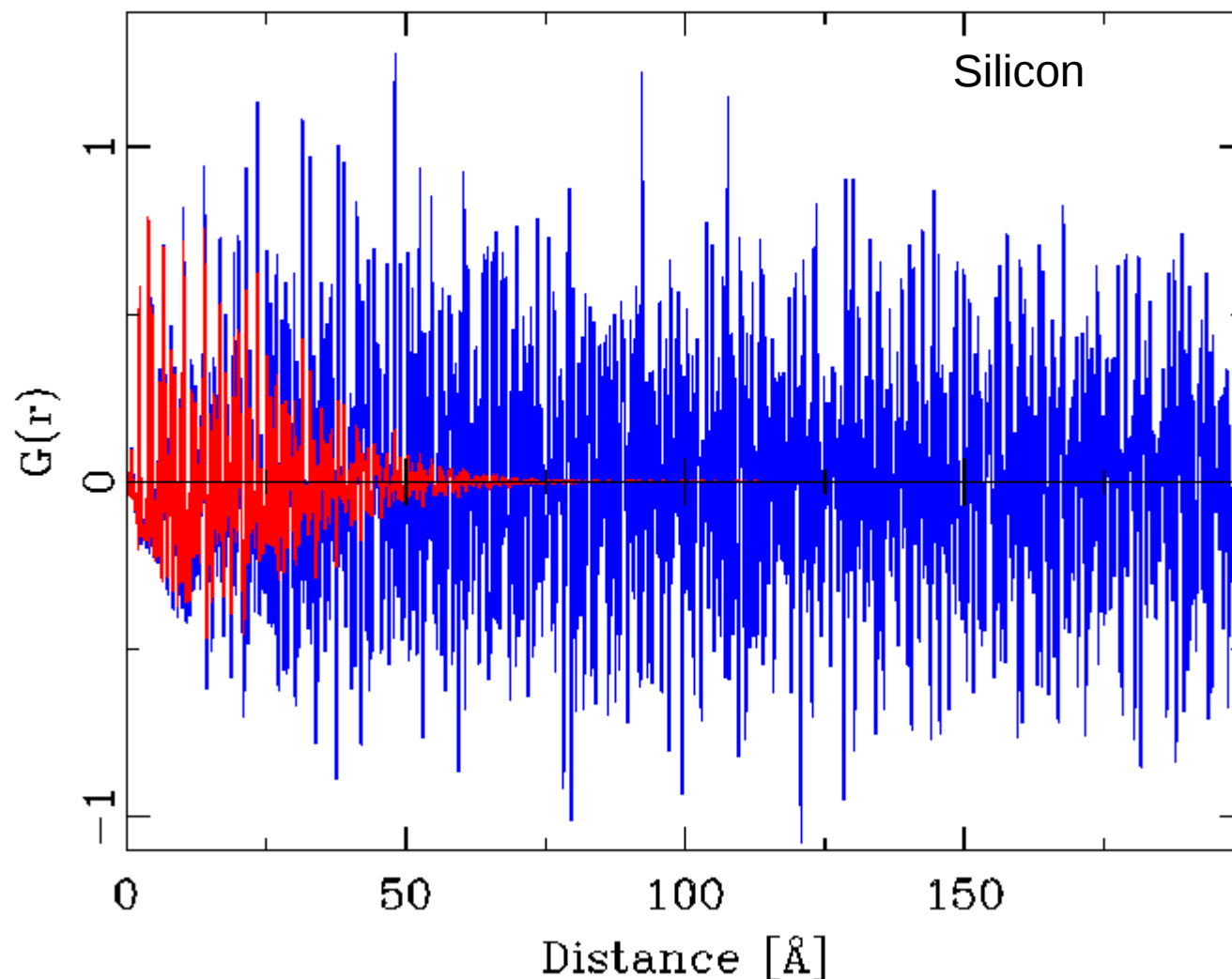
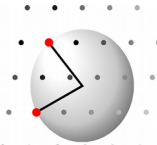




PSI high resolution data
~ 30 min

APS low resolution data
~ 1 min

Reflections well resolved
at PSI up to
and beyond 21.4 \AA^{-1}



PSI high resolution data
~ 30 min

APS low resolution data
~ 1 min

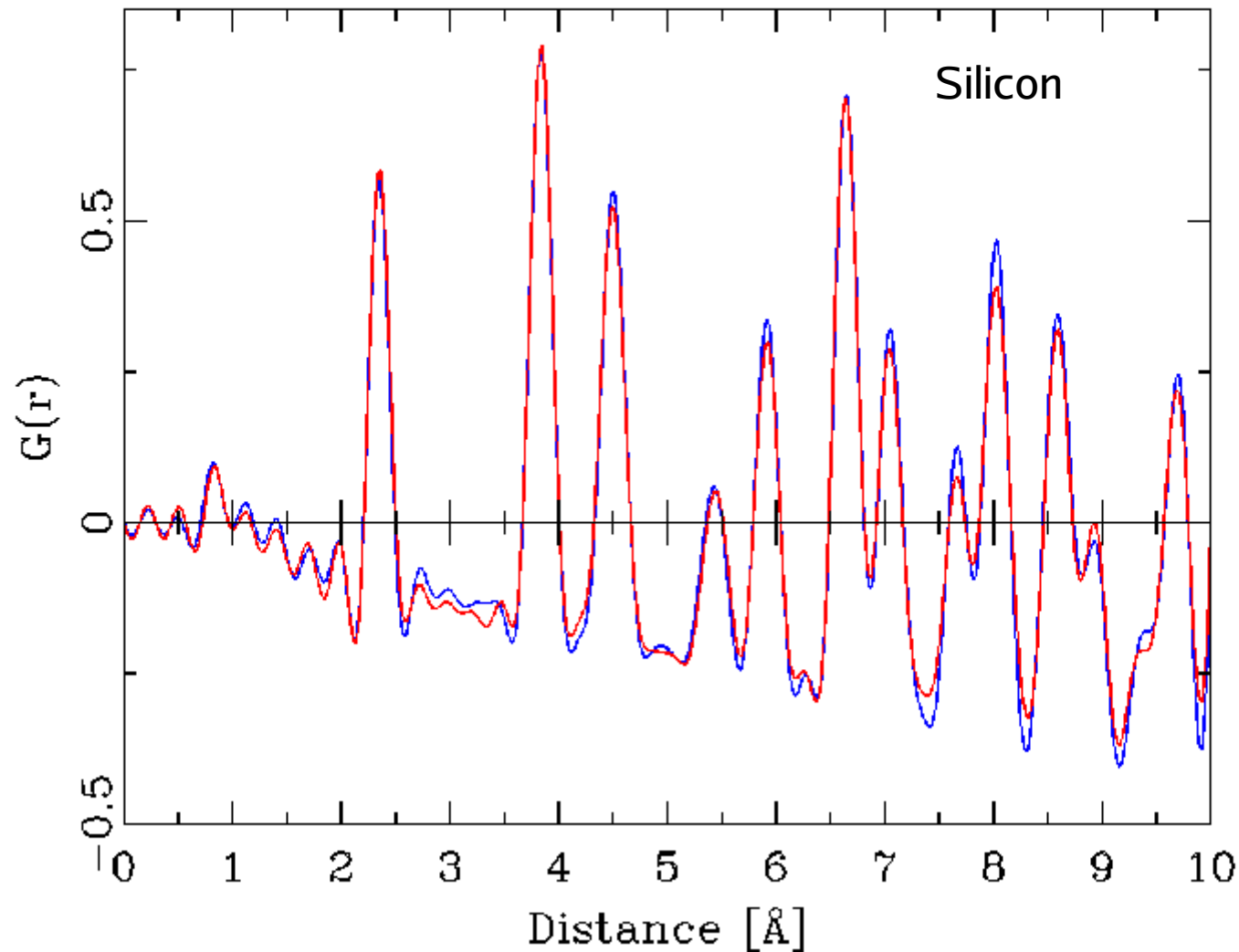
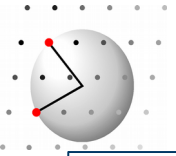
Reflections well resolved
at PSI up to
and beyond 21.4 \AA^{-1}

$G(r)$ @ PSI extends well
beyond 200 \AA !

$G(r)$ @ APS almost ZERO
beyond 70 \AA !

PDF peak height influences determination of nanoparticle diameter

High resolution NOT needed for small ($<5\text{nm}$) objects; only good if sample has narrow Bragg



PSI high resolution data
~ 30 min

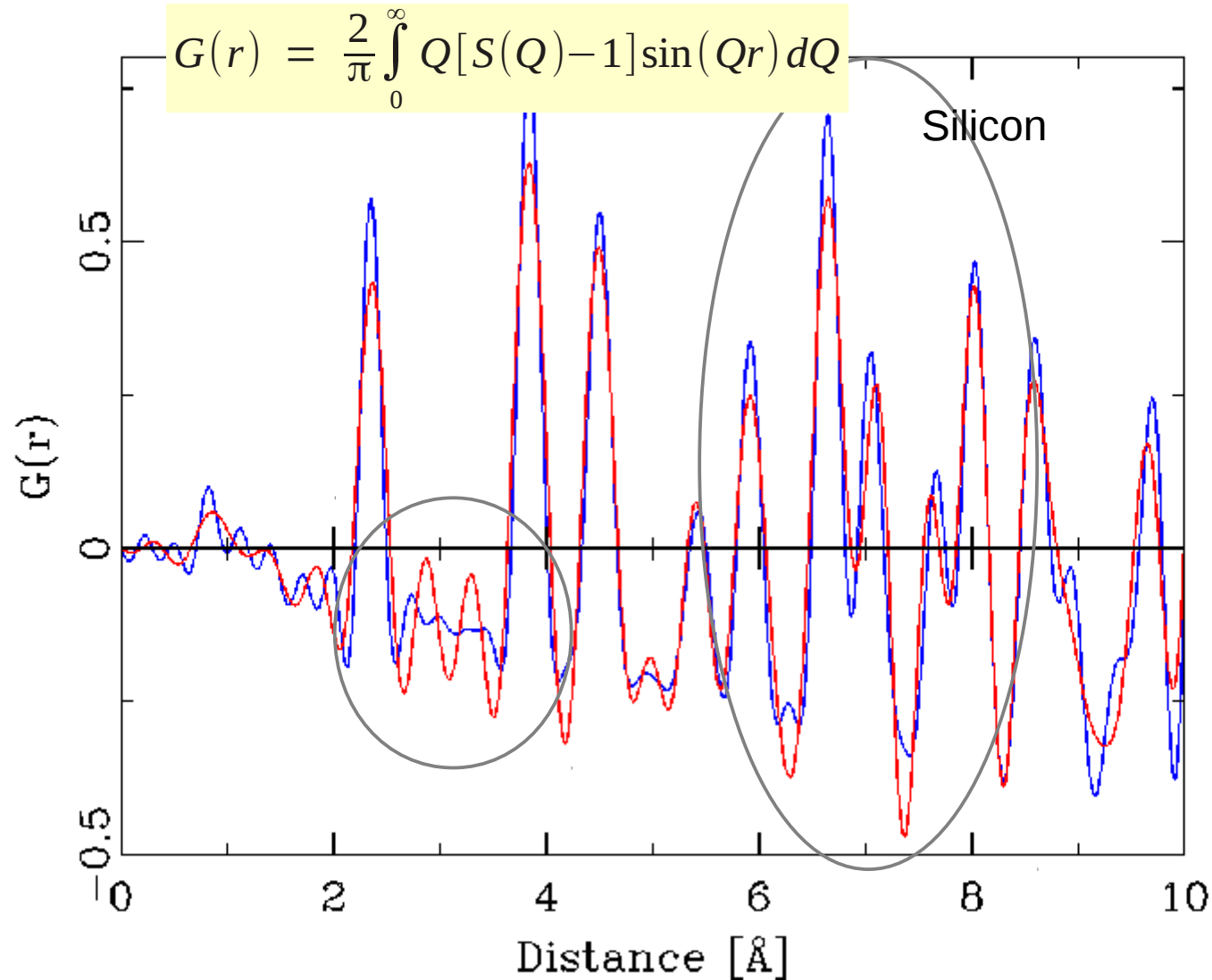
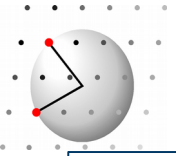
APS low resolution data
~ 1 min

Reflections well resolved
at PSI up to
and beyond 21.4 \AA^{-1}

$G(r)$ @ PSI extends well
beyond 200 \AA !

$G(r)$ @ APS almost ZERO
beyond 70 \AA !

Width of PDF maxima
is identical
(at least for low r -region)



PSI high resolution data
Silicon

$Q_{\max} = 21. \text{ \AA}^{-1}$

Synchrotron

$Q_{\max} = 15. \text{ \AA}^{-1}$

Mo-K α

Apparent shift of
smaller PDF maxima !?!

decreases with increasing
 Q_{\max}

Be careful with direct
interpretation of peak positions!

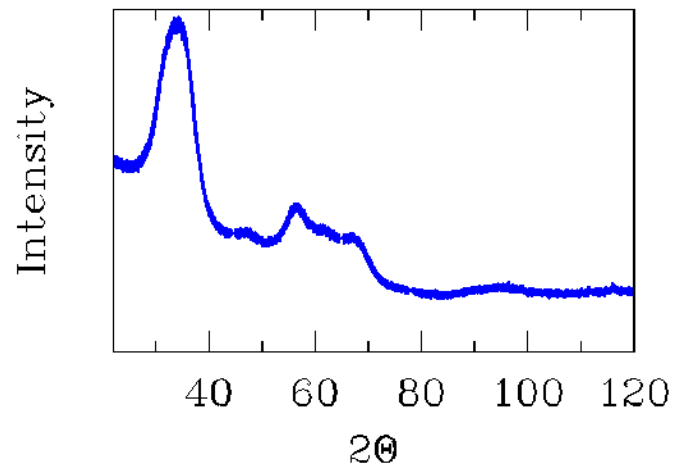
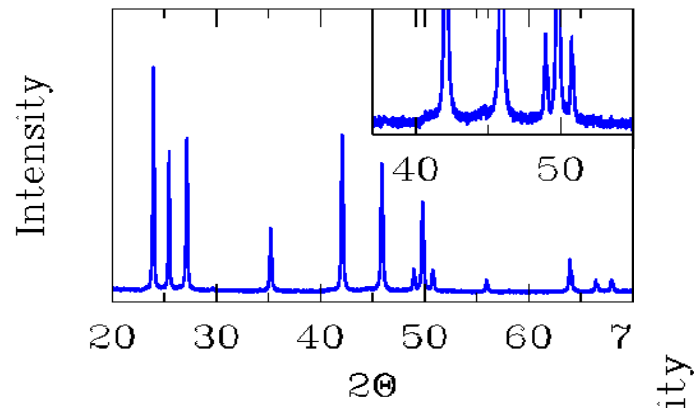
Local disorder in Si !?!

Anything that changes with
 Q_{\max} is not real

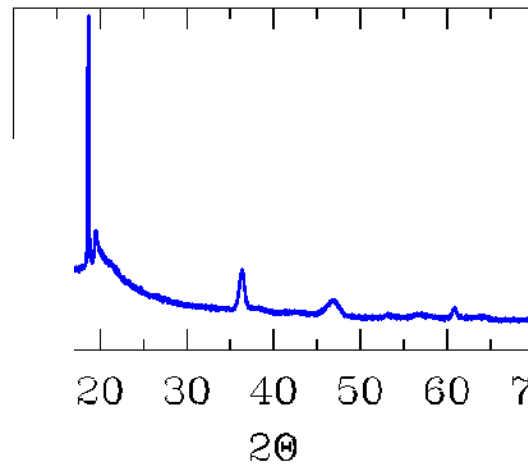
Always calculate PDF for several different values of Q_{\max}

Powder Diffuse Scattering

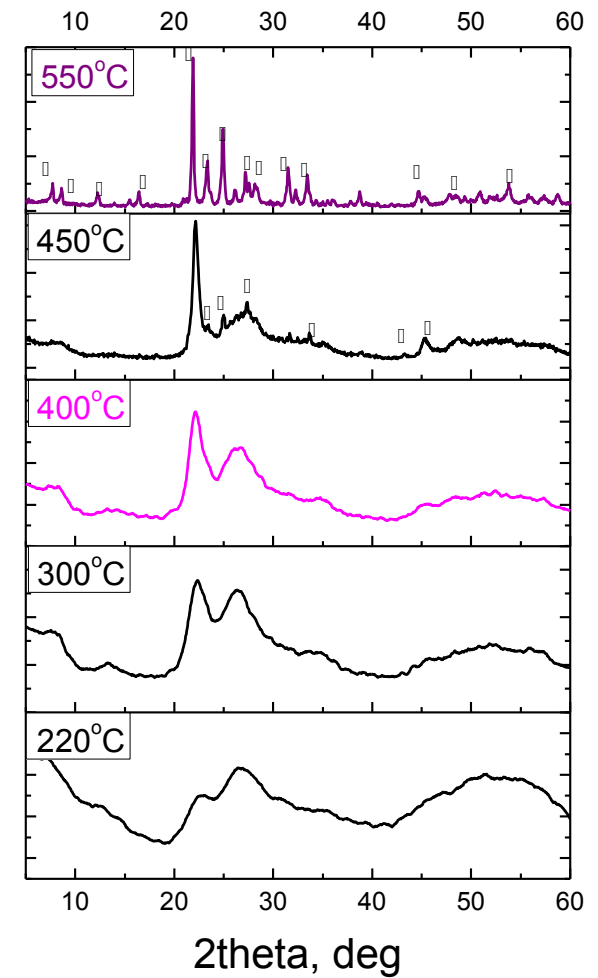
CdSe crystalline material



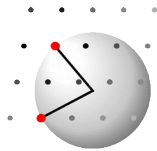
Nano crystalline ZnO



H_2TiO_3



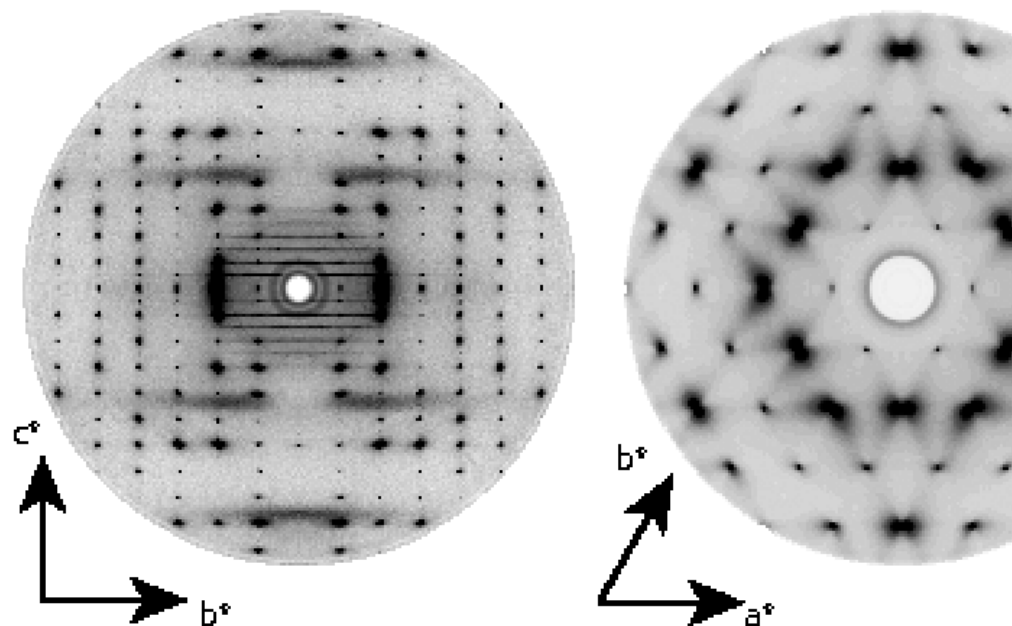
Mo-V-Nb oxides





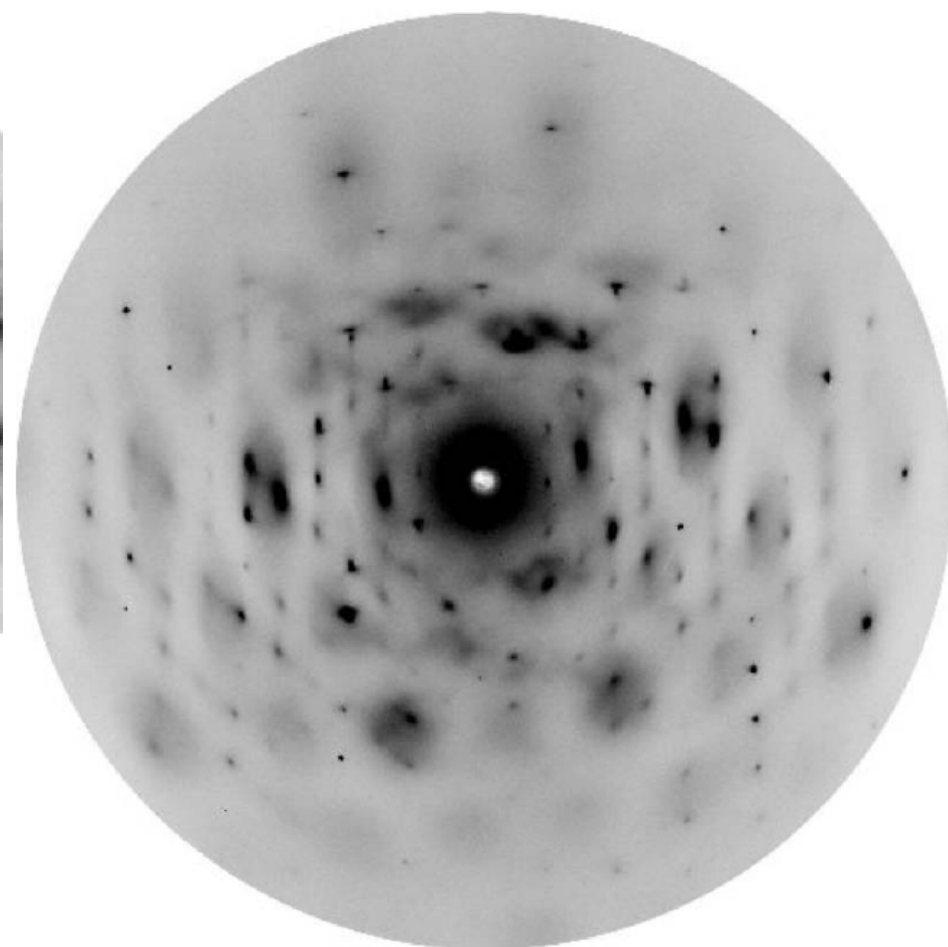
Singe Crystal Diffuse Scattering

Th Weber PhD München 1994

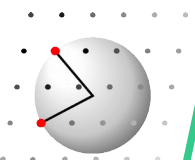


Alkane chains in Urea

Diffuse scattering by $0.05 \mu\text{m}^3$ single crystal
Neder et al Clays & Clay Minerals 1999

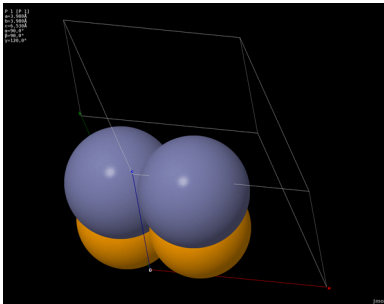


Pentachloronitrobenzene



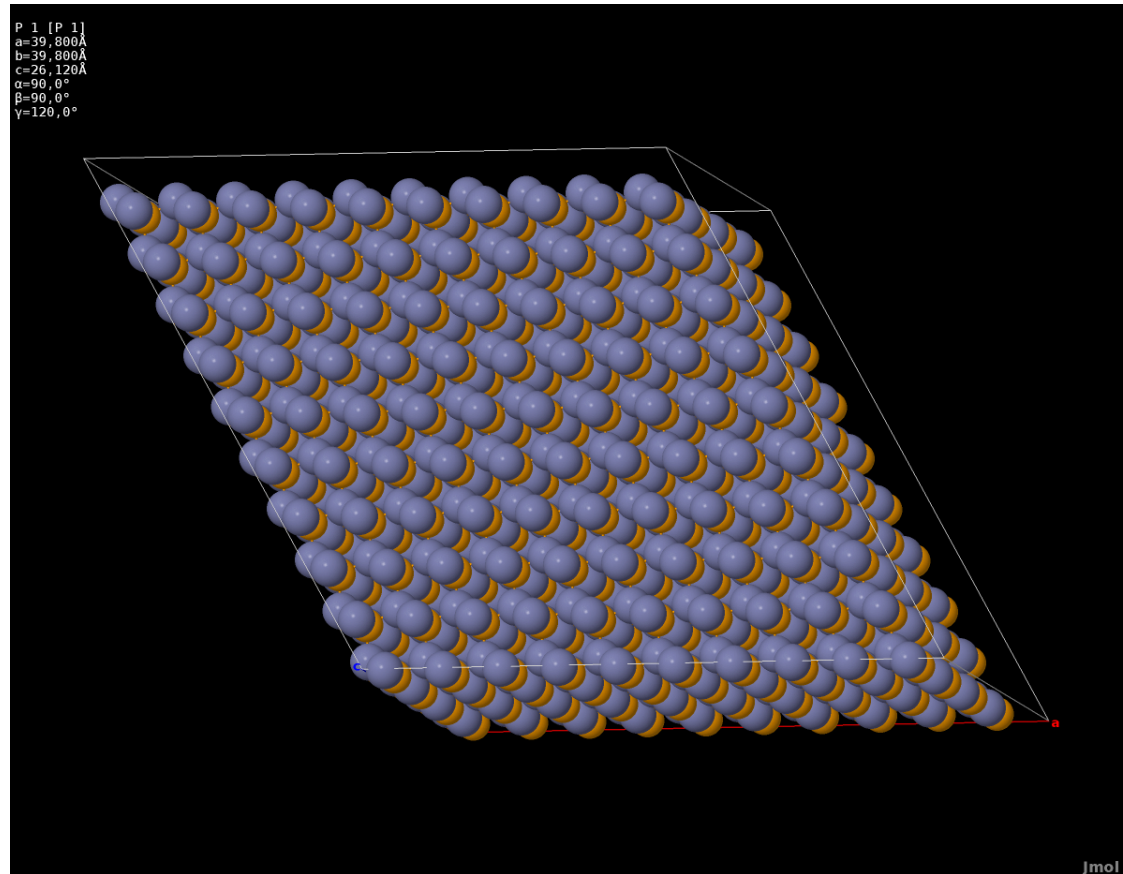
Goal

Simulate all things



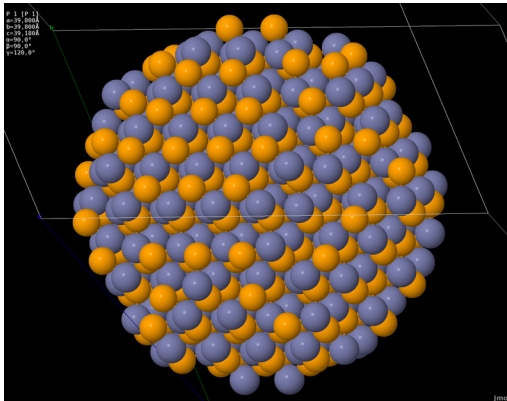
small

and large
which means periodic

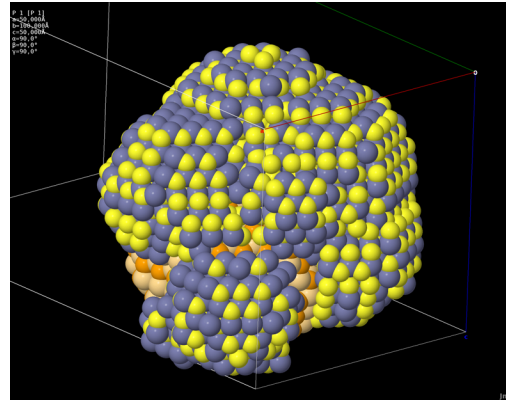


Goal

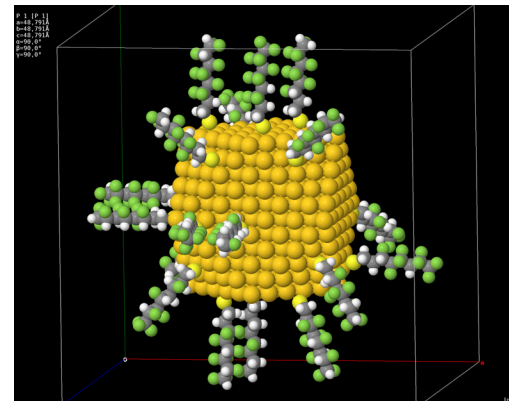
Simulate all things



simple

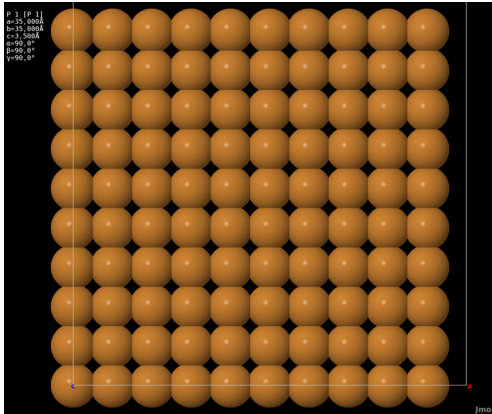


and complex

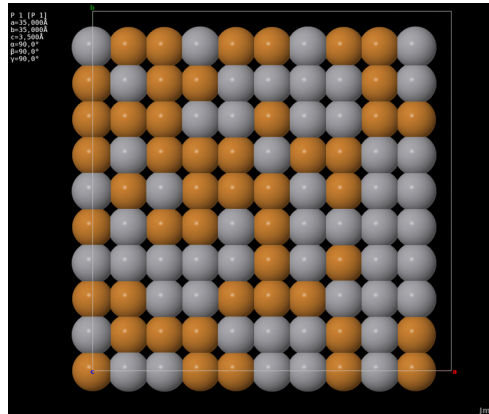


Goal

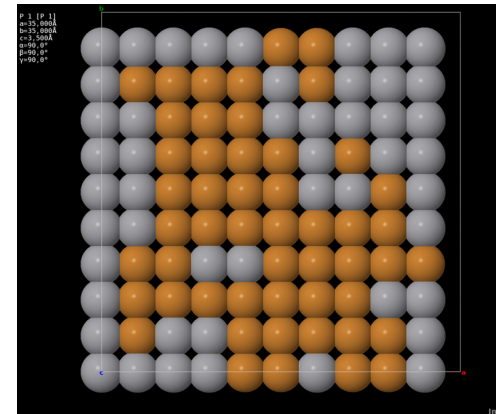
Simulate all things



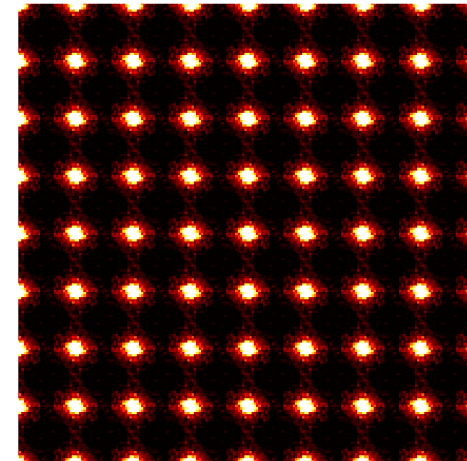
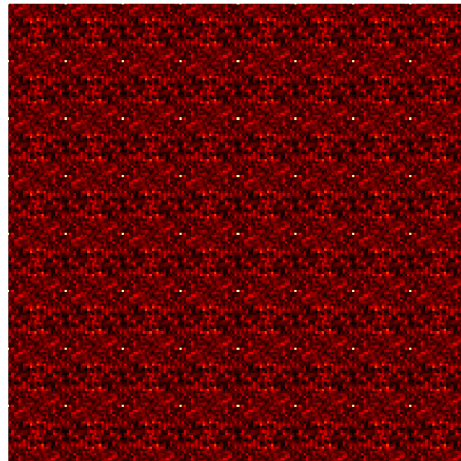
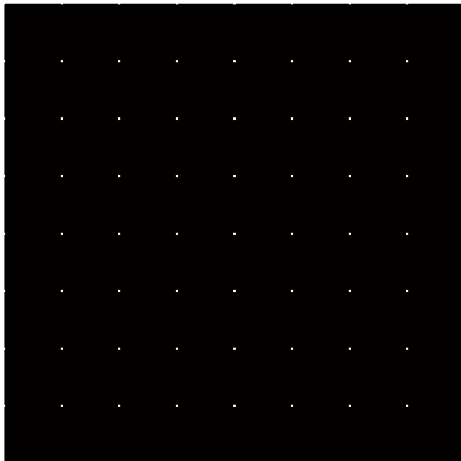
perfect



or random



and short range ordered





Differences to *Rietveld* or *single crystal Bragg* work

Rietveld / Single Crystal Bragg

Simulation of disorder

Just atoms in **one** asymmetric unit

list of ~ million atoms

sequence of atoms in computer
memory irrelevant

very often one needs relations
between „neighbors“

need to know sequence

Basic information for a simulation

Chemical composition

ZnSe

Space group, lattice constants

$F\bar{4}3m$ # 216; 5.41 Å

symmetry operations,
or;
generators, asymmetric unit

4 * 24 symmetry operations

choose generators to reproduce
sequence as in International Tables

(1) 1
(2) 2 0,0,z
(3) 2 0,y,0
(5) 3⁺ x,x,x
(13) m x,x,z
t(0, ½, ½); t(½, 0, ½)
t(1,0,0); t(0,1,0); t(0,0,1)

atoms, positions

Zn 4 a 0,0,0

S 4 c ¼, ¼, ¼

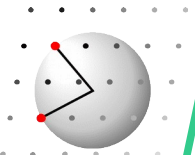
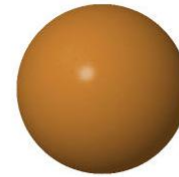
atomic displacement parameters

Typical Simulation

Read asymmetric unit

```
title    primitive cubic
spcgr    Pm-3m
cell     5.00, 5.00, 5.00, 90.0, 90.0, 90.0
atoms
Cu       0.000000, 0.000000, 0.000000, 0.8
```

Alternative: import CIF file
import SHELX file



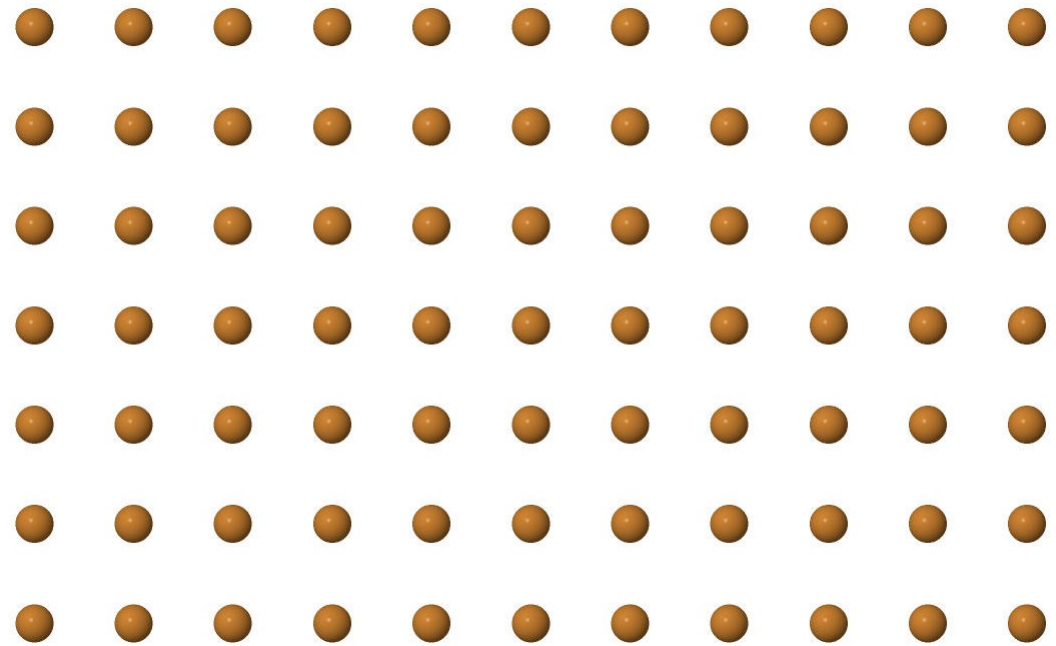
Typical Simulation

Read asymmetric unit

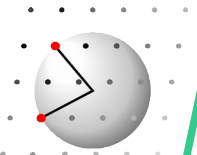
Expand to full unit cell

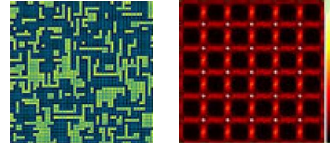
Expand to a block sized crystal

Keep shape simple!
Simulated crystal is small
compared to real sample!



Jmol





Typical Simulation

Read asymmetric unit

Expand to full unit cell

Expand to a block sized crystal

Introduce defects

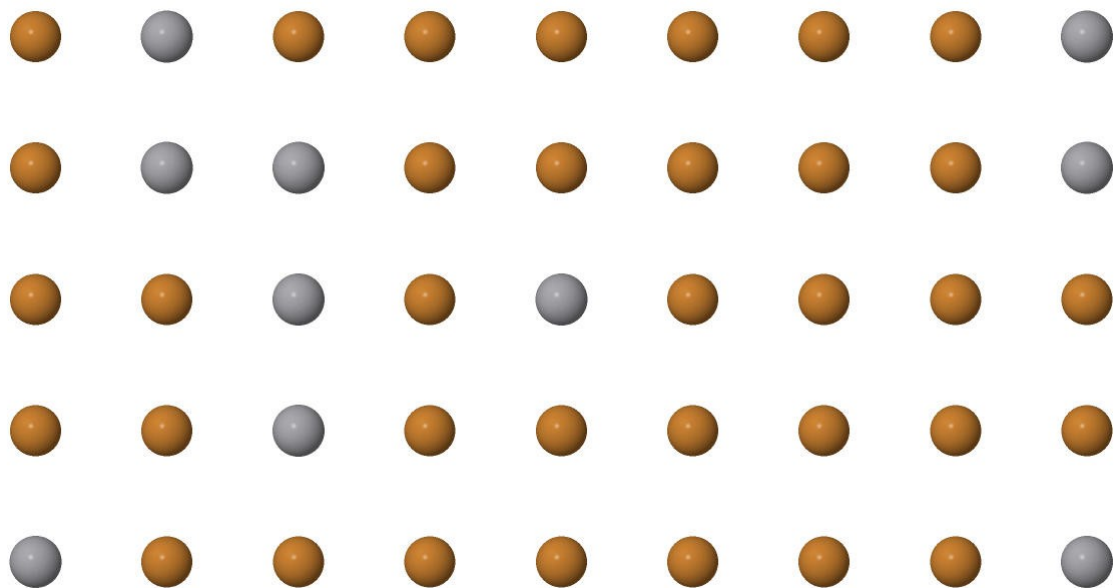
Randomly placed

atoms

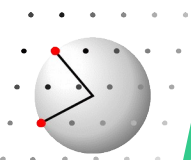
domains

shift atoms

...



Jmol



Typical Simulation

Read asymmetric unit

Expand to full unit cell

Expand to a block sized crystal

Introduce defects

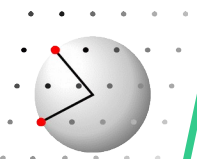
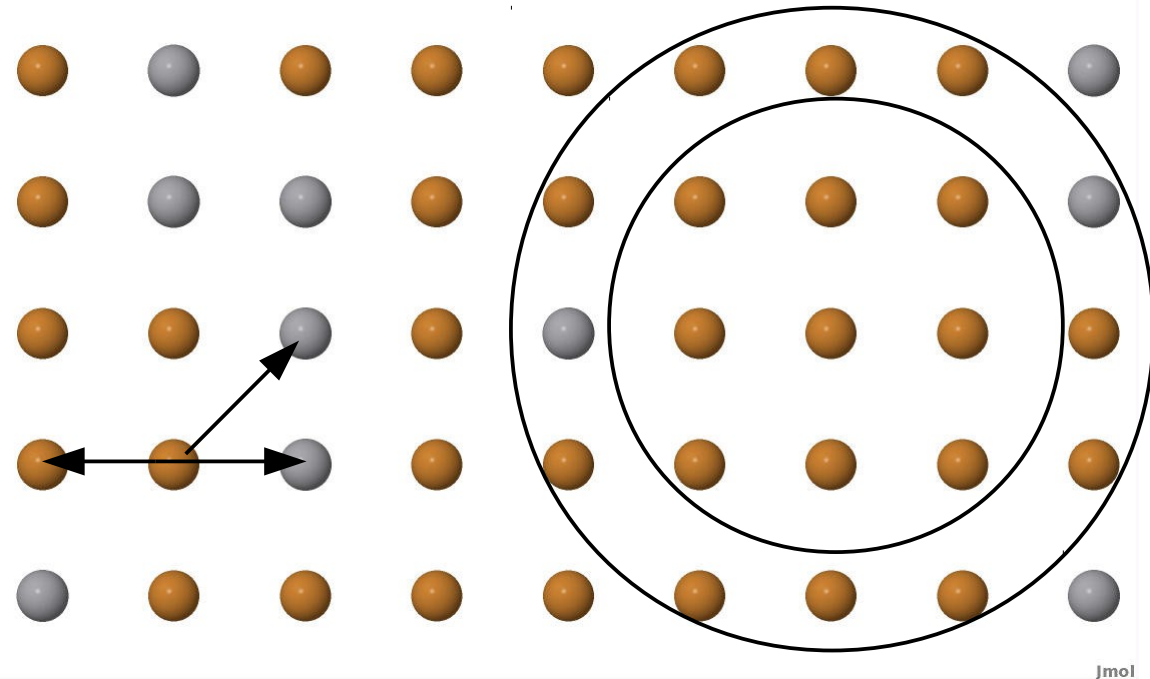
Randomly placed
atoms
domains
shift atoms
...

Introduce correlations

Neighbors in $[100]$ or ...

Neighbors in a shell

Atoms type A and B or ...





Typical Simulation

Read asymmetric unit

Expand to full unit cell

Expand to a block sized crystal

Introduce defects

Introduce correlations

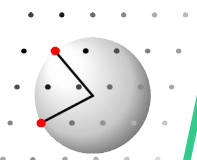
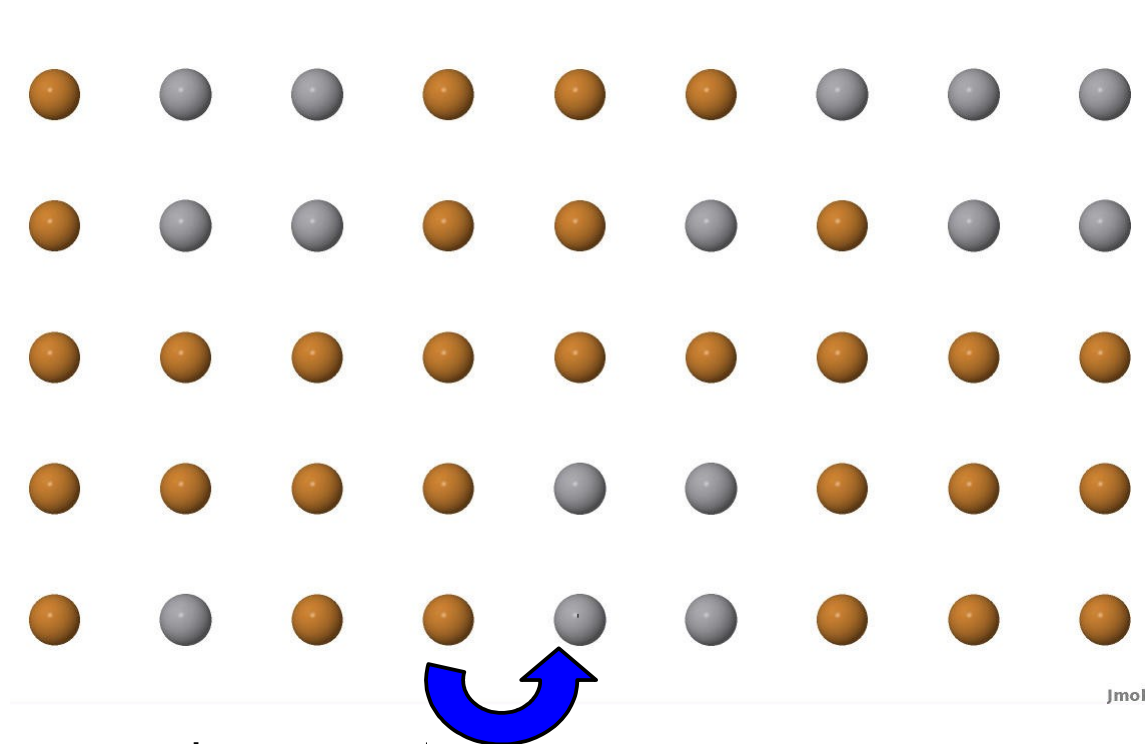
Modify crystal while
minimizing energy
associated with correlations

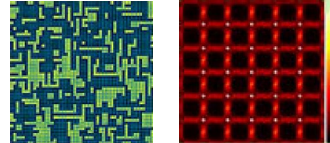
Ising models for chemical short range order

Distance potentials

Angular potentials

...





Typical Simulation

Read asymmetric unit



Expand to full unit cell

Expand to a block sized crystal



Introduce defects



Introduce correlations



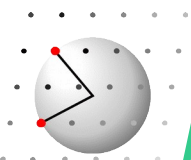
Modify crystal



Calculate: single crystal diffraction pattern
powder diffraction pattern
powder PDF
3D PDF

Refine: structure and disorder against experimental pattern

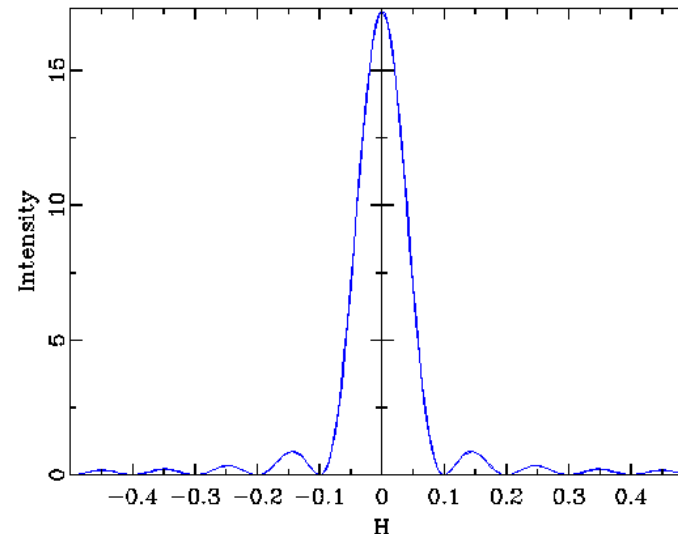
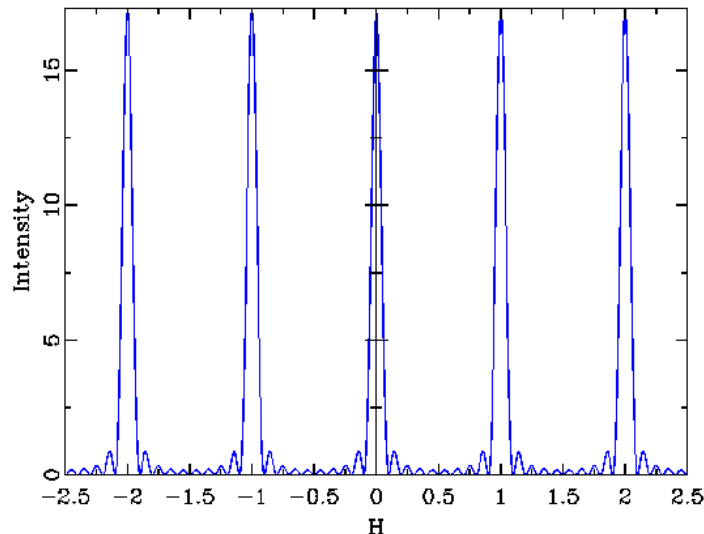
Jmol



Diffraction by simulated crystal structures

chain of 10 Si atoms

Calculated neutron diffraction pattern (intensity)



periodic reflections spaced at $1/(\text{Si-Si})$ detail of a reflection

zero points at $1/10$ reciprocal
lattice constants

Simulated crystals are small !!

Calculate diffraction pattern while avoiding finite size effects

Simulate Block of N unit cells; calculate in reciprocal space at $1/N \cdot a^*$

Hamlet: 2π or not 2π ?

Typical Simulation

Read asymmetric unit

Expand to full unit cell

Expand to a block sized crystal

Introduce defects

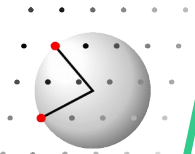
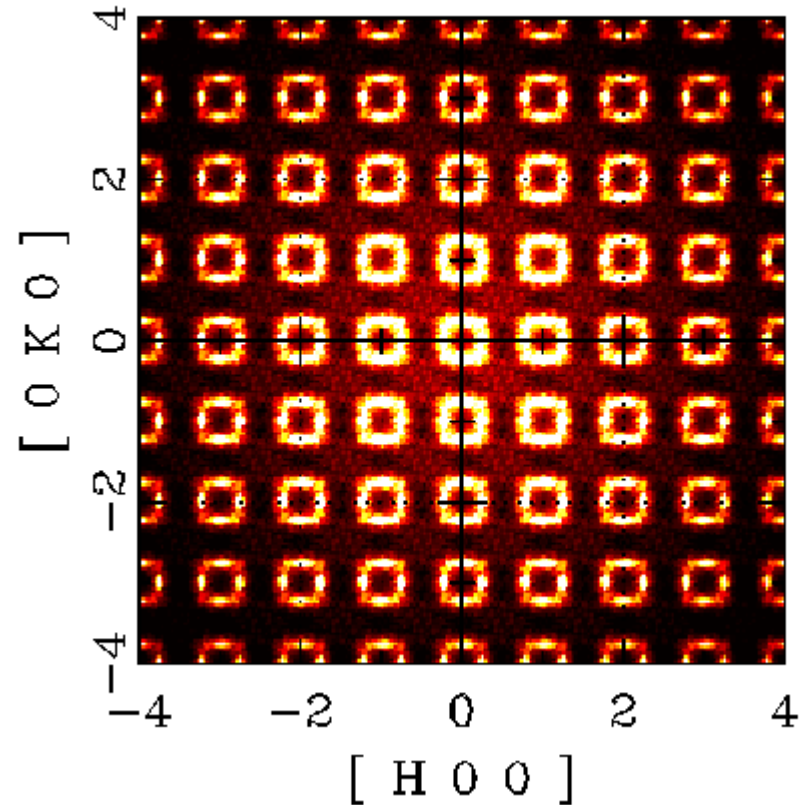
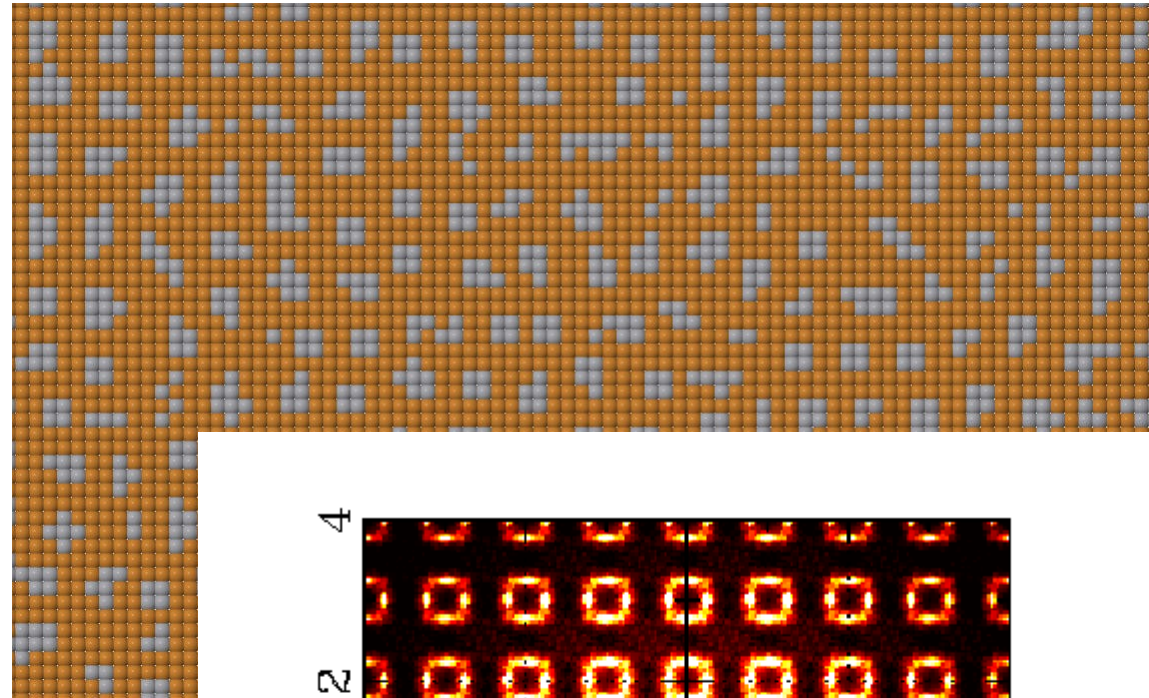
Introduce correlations

Modify crystal

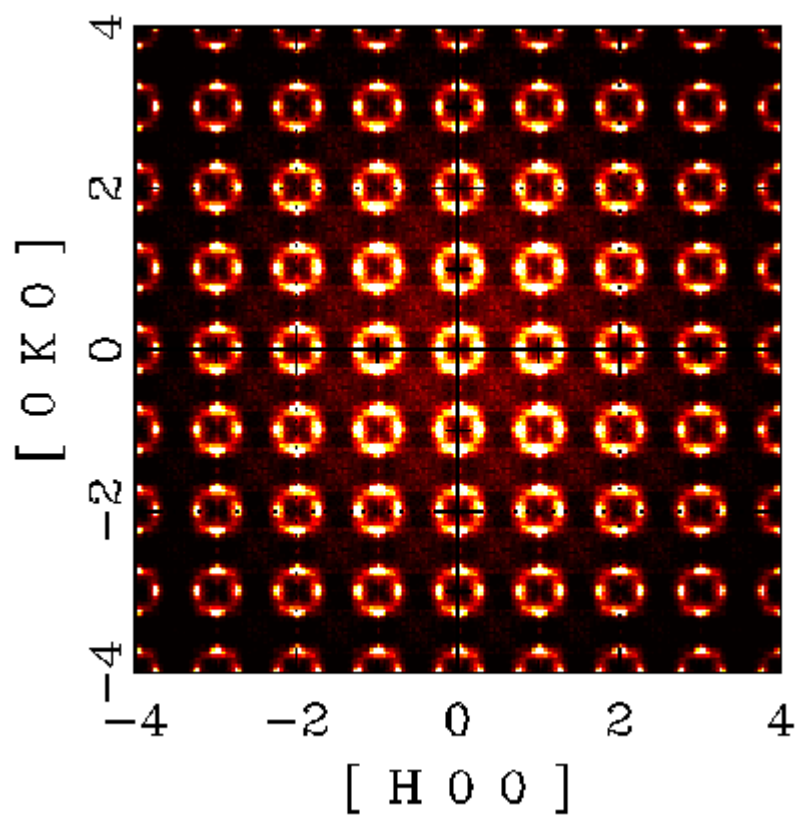
Calculate diffraction pattern / PDF

Compare to experimental data

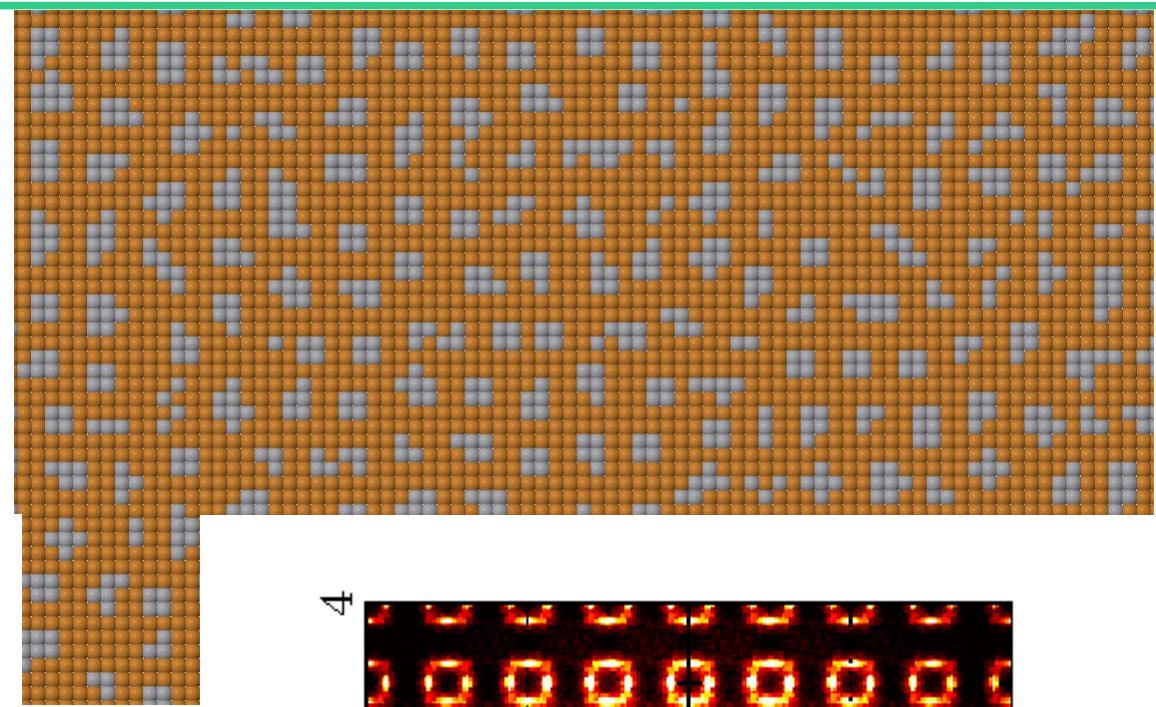
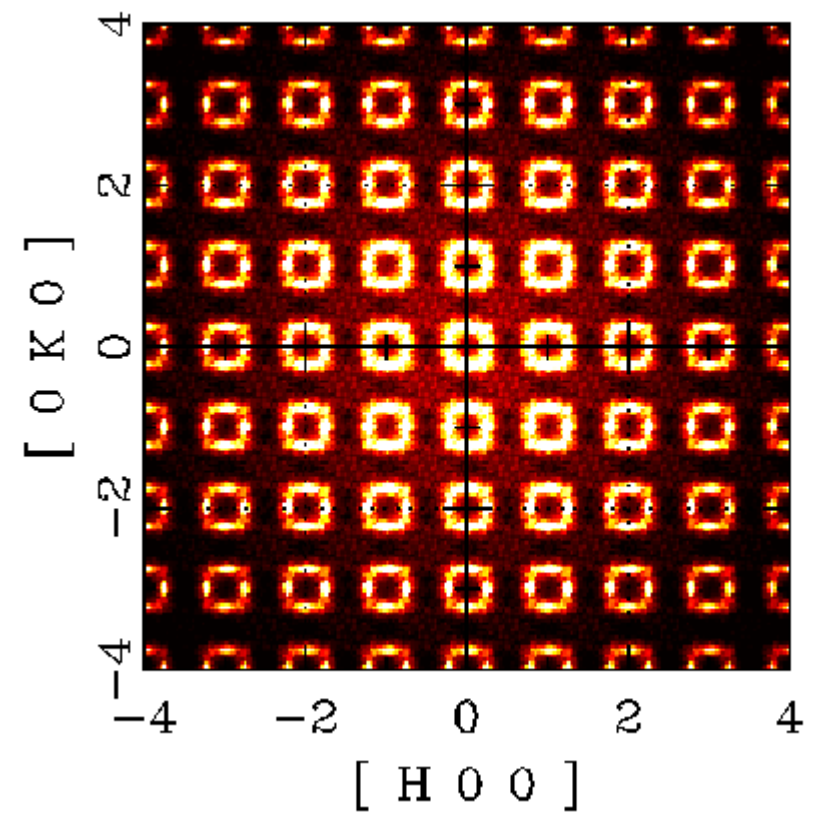
Refine disorder parameters

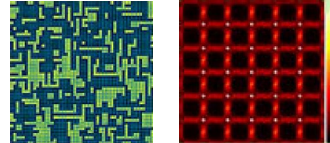


A word of CAUTION



logarithmic scale
Erlangen University

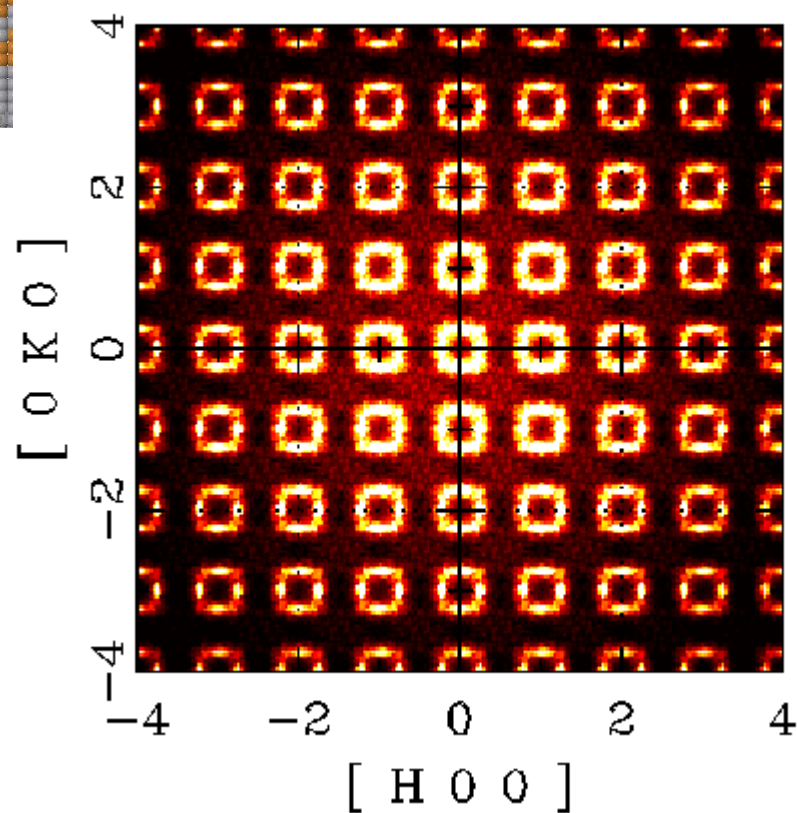
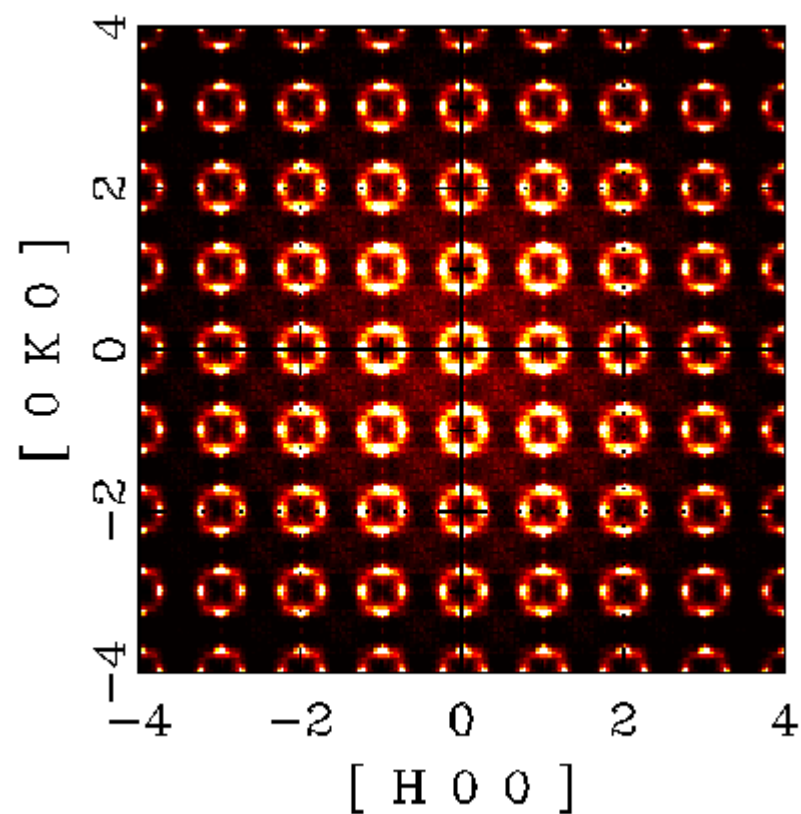
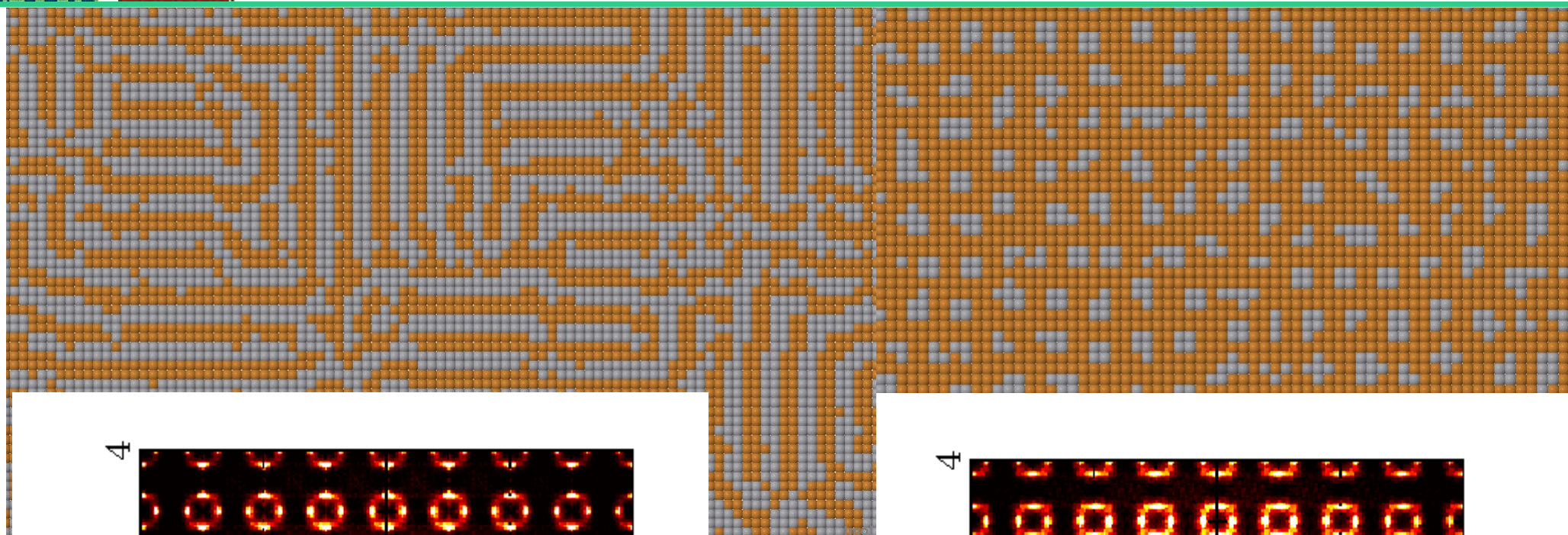




A:B = 1:1

Identical Correlations

A:B = 3:1

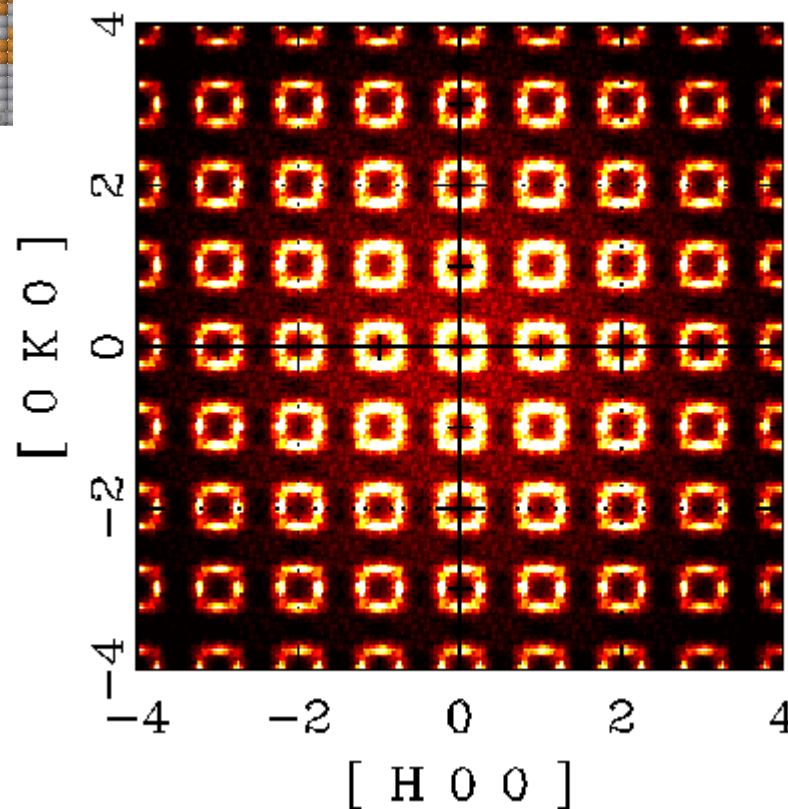
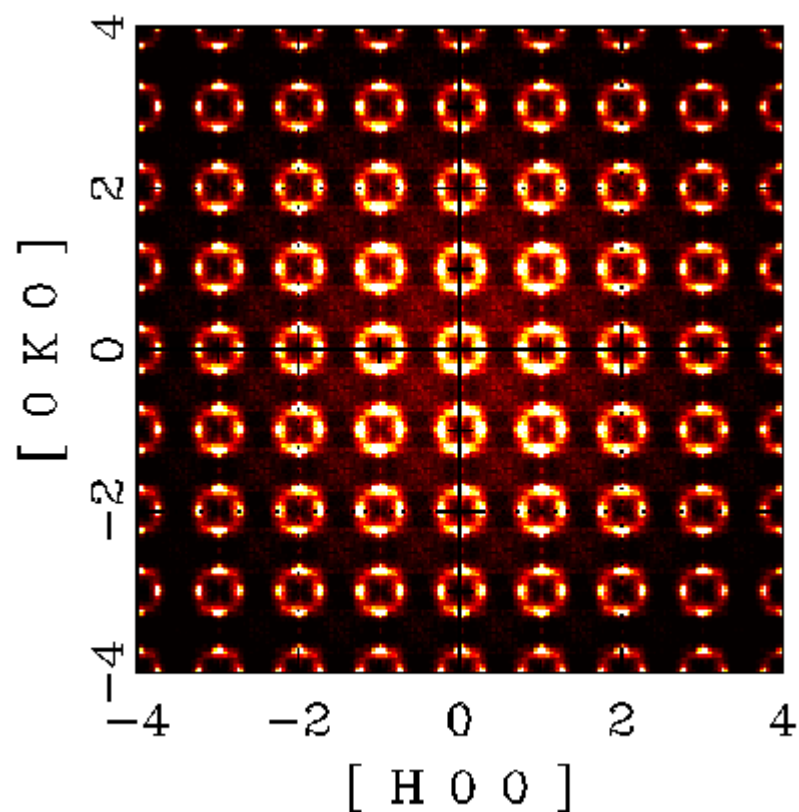


logarithmic
Erlangen

A word of CAUTION

Analyze single crystal refinement
Obtain complementary information:
HRTEM, SAXS, SANS, PDF, Chemistry

Helps to define correct starting model



logarithmic scale

University of Erlangen-Nürnberg



DISCUS_SUITE

Command line driven program

you type individual commands

main commands must be memorized

Extensive on-line help /Manual / Book

very flexibel

includes a programming language

allows simulation of any atom configuration

crystal, glass, nanoparticles, quasicrystals, ...

Structured into menus

combine several main tasks

read a unit cell / a structure

calculate diffraction pattern / PDF

plot the structure

extended defects

SRO, strain, stacking faults, domains, waves

nanoparticle builder

surface decoration

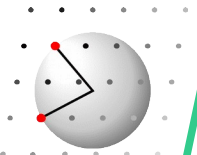
Refinement

Flexible global optimizer

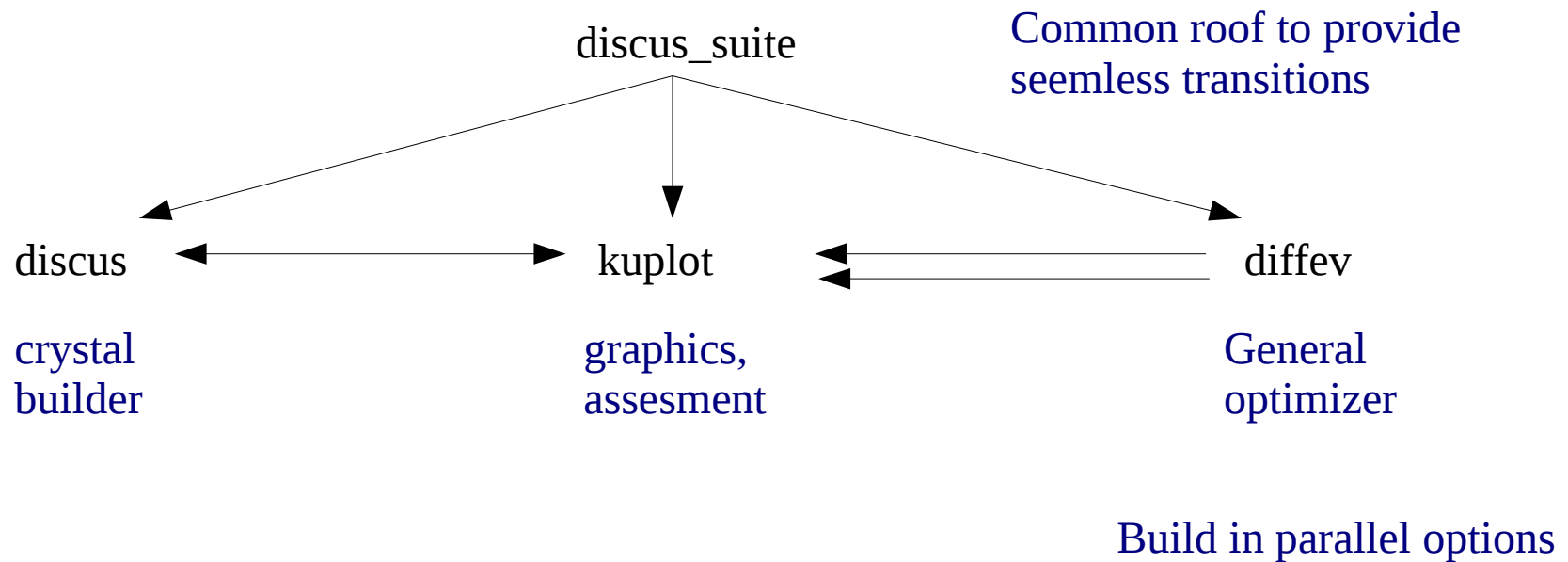
Define disorder model

Simulate structure

calculate diffraction pattern / PDF



DISCUS Program package

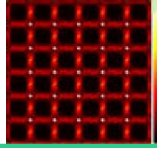
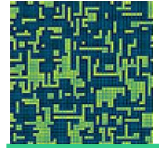


Availability:

Linux, Unix, Mac-OS, Windows

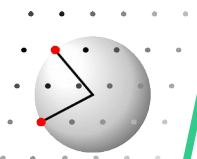
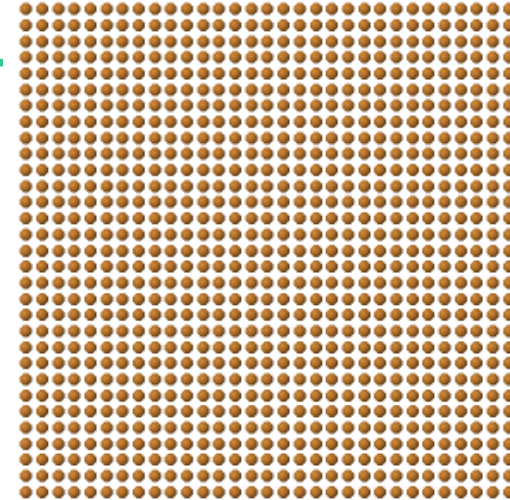
Single computer

Massive parallel farms



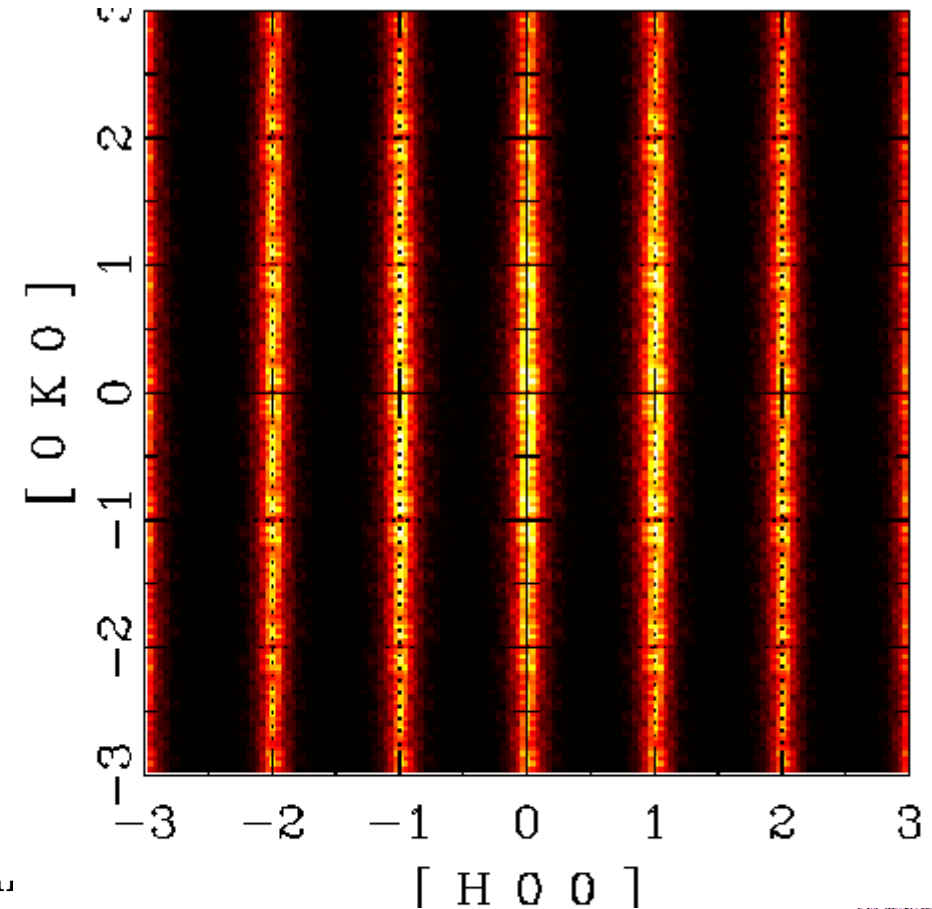
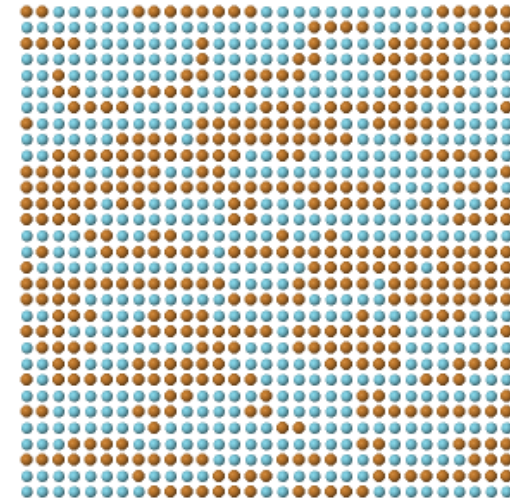
DISCUS / short range order menu

```
#      Example
read
      cell crystal.cell,200, 200,1
#
plot
      program jmol
      select all
      outfile crystal_plot.cif
      run plot:inter
exit
#
```



DISCUS / short range order menu

```
#
read
    cell crystal.cell,200, 200,1
#
replace cu,au,all,0.50
#
mmc
#
    set vec,1, 1,1, 1, 0, 0
    set vec,2, 1,1, -1, 0, 0
    set neig,vec,1,2
#
    set mode, 1.0, swchem,all
    set targ,1,corr,cu,au, 0.80, 0.0,CORR
    set cyc, 100*n[1]
    set feed, 5*n[1]
    set temp, 2.5
    run
exit
#
fourier
    xray
    ll -3.00, -3.00, 0.00
    ll -3.00, -3.00, 0.00
    ll -3.00, -3.00, 0.00
    na 151
    no 151
    run
exit
```



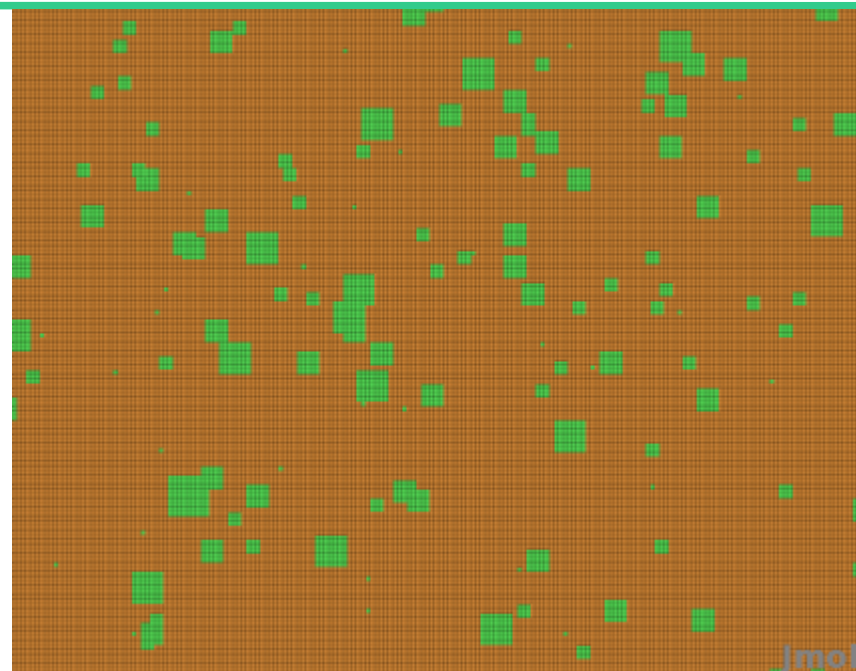
DISCUS / domain concept

```
#      Example
read
      cell host.cell,200, 200,1
#
domain
#
      mode pseudo
      input domain.list  !Domain origins
#
      assign character, SI, cube
      assign fuzzy,      SI, 1.5
      assign content,    SI, guest.stru
      assign orient,     SI,1,  1,0,0,  0
      assign orient,     SI,2,  0,1,0,  0
      assign orient,     SI,3,  0,0,1,  0
      assign shape,      SI,1,  2,0,0,  0, 0.5
      assign shape,      SI,2,  0,2,0,  0, 0.5
      assign shape,      SI,3,  0,0,1,  0, 0.0

      run
exit
#
plot . . .
```

A regular DISCUS file with
Domain coordinates

use any tool to distribute
the origins



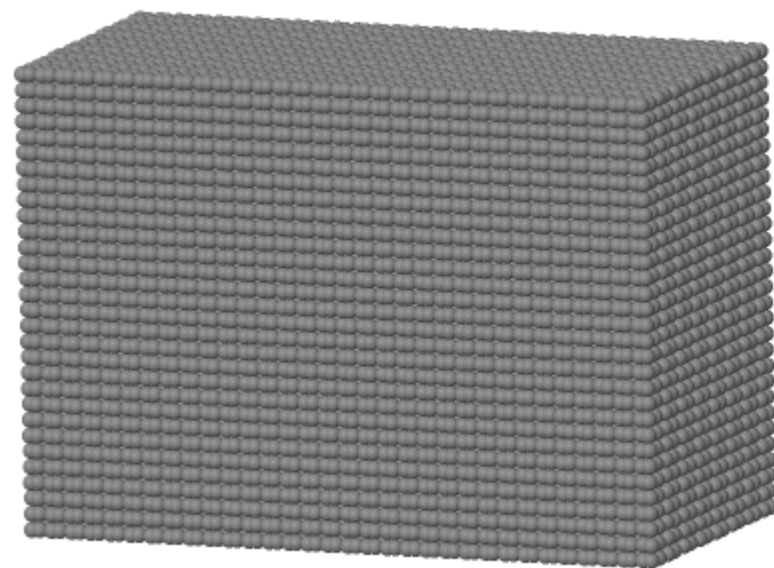
domain.list

```
title Primitive host structure
spcgr P4
cell 2.50, 2.50, 2.50, 90., 90., 90.
atoms
SI 71.0, -100.00, 0.00, 0.05, 1
SI -22.0, -98.00, 0.00, 0.05, 1
. . .
SI -19.0, 2.00, 0.00, 0.05, 1
SI 75.0, 2.00, 0.00, 0.05, 1
SI -20.0, 5.00, 0.00, 0.05, 1
. . .
```

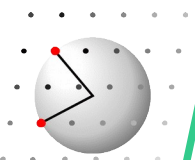



DISCUS / nanoparticle

```
variable real, diam_a
variable integer, ncellx
#
diam_a = $1
#
read
    free 2,3,4, 90,90,90, Pmmm
insert C, 0.3, 0.25,0.3, 0.5
#
ncellx = int(diam_a/lat[1]) + 2
#
save
    outf internal.cell
    run
exit
#
read
    cell internal.cell, ncellx, ncelly, ncellz
@plot.mac block
```



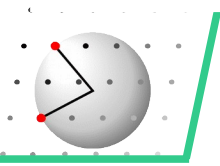
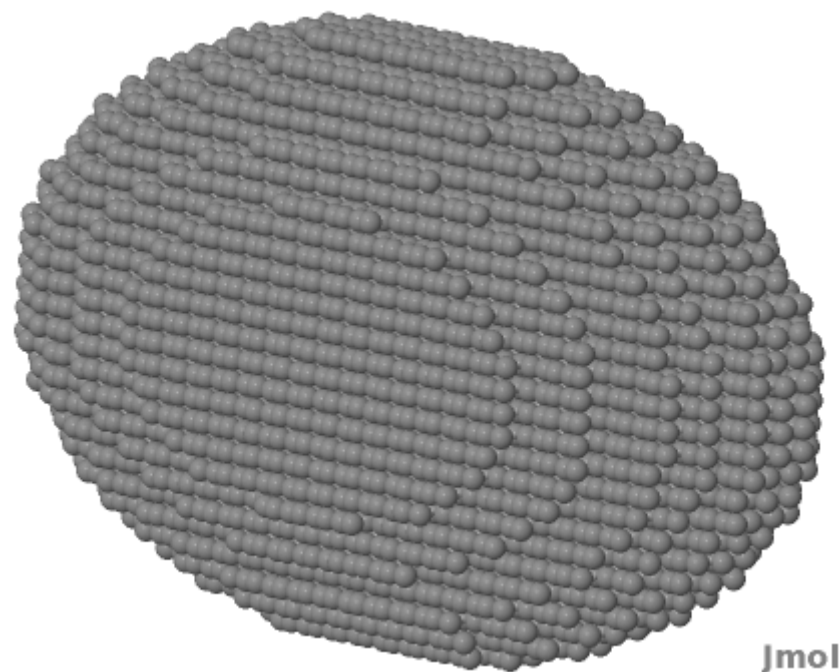
Jmol





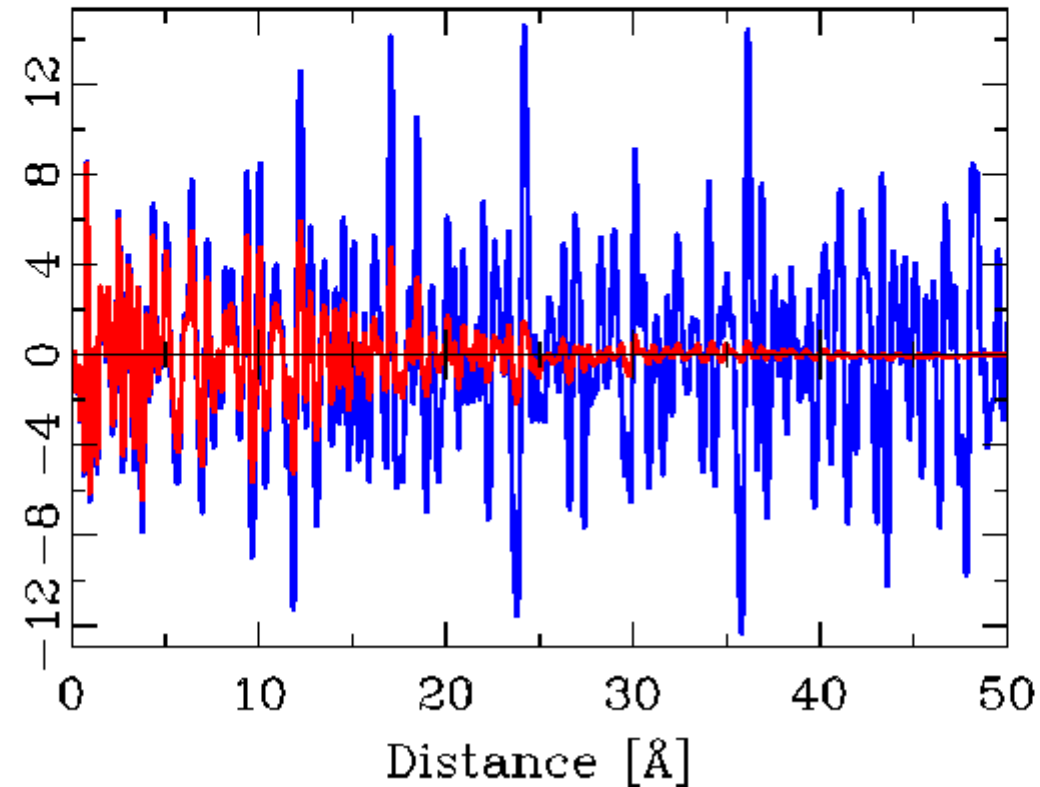
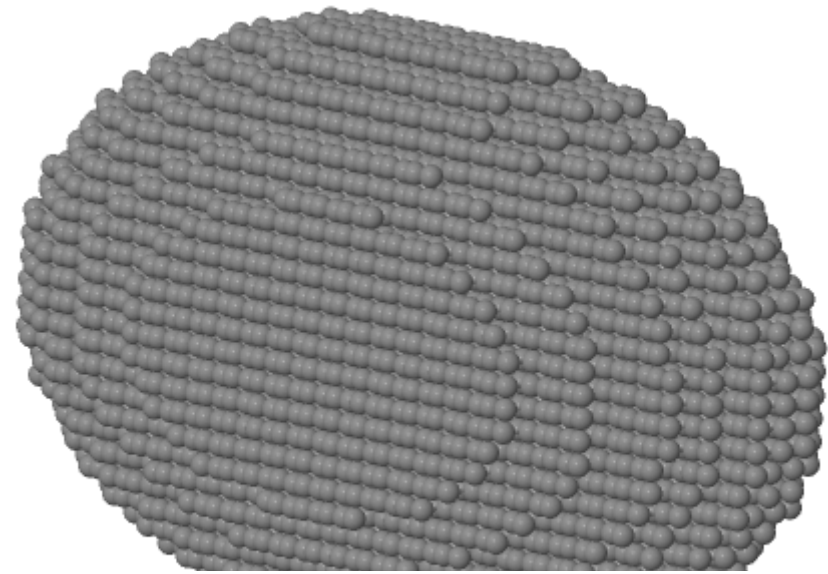
DISCUS / nanoparticle

```
variable real, diam_a
variable integer, ncellx
...
#
diam_a = $1
diam_c = $2
diam_b = $3
#
read
    free 2,3,4, 90,90,90, Pmmm
insert C, 0.3, 0.25,0.3, 0.5
#
ncellx = int(diam_a/lat[1]) + 2
...
#
save
    outf internal.cell
run
exit
#
read
    cell internal.cell, ncellx, ncelly, ncellz
@plot.mac block
surface
    boundary, ellip, diam_a, diam_b, diam_c
exit
purge
@plot.mac ellipsoid
```



DISCUS / nanoparticle

```
variable real, diam_a
variable integer, ncellx
...
#
diam_a = $1
diam_c = $2
diam_b = $3
#
read
    free 2,3,4, 90,90,90, Pmmm
insert C, 0.3, 0.25,0.3, 0.5
#
ncellx = int(diam_a/lat[1]) + 2
...
#
save
    outf internal.cell
run
exit
#
read
    cell internal.cell, ncellx, ncelly, ncellz
@plot.mac block
surface
    boundary, ellip, diam_q, diam_b, diam_c
exit
purge
@plot.mac ellipsoid
@pdf.mac ellipsoid
```



Neder & Proffen, (Oxford, 2008)

Diffuse Scattering and Defect Structure Simulation

Simulation and refinement of disordered structures, cook book

T.R. Welberry (Oxford, 2004)

Diffuse X-ray Scattering and Models of Disorder

Limited to X-ray diffraction, short experimental part,
otherwise extensive theory, many examples

V.M. Nield & D.A. Keen (Oxford, 2001)

Diffuse Neutron Scattering from Crystalline Materials

Limited to neutron diffraction, otherwise extensive theory,
experimental methods, some simulations, many examples

T. Egami & S.J.L. Billinge (Pergamon 2003)

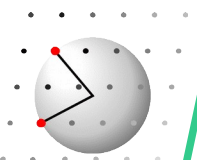
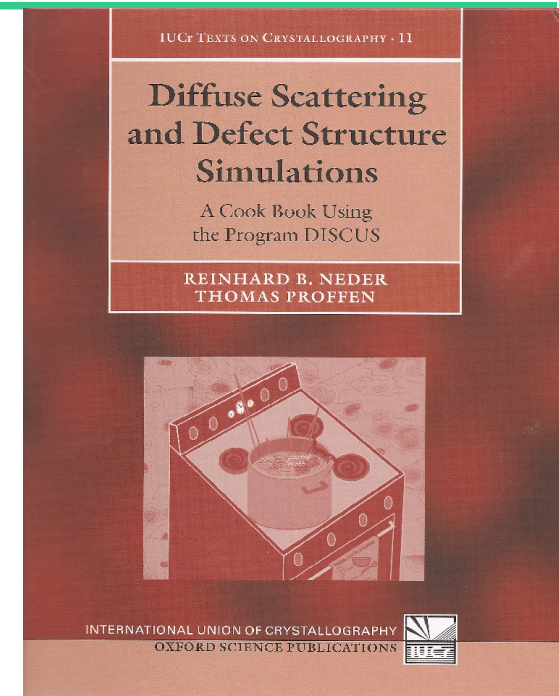
Underneath the Bragg Peaks

Extensive description of PDF Method

S.J.L. Billinge & M.F. Thorpe Eds. (Plenum, 1997)

Local Structure from Diffraction

Collection of papers on PDF and disorder in general



Interactive Tutorial on Diffraction

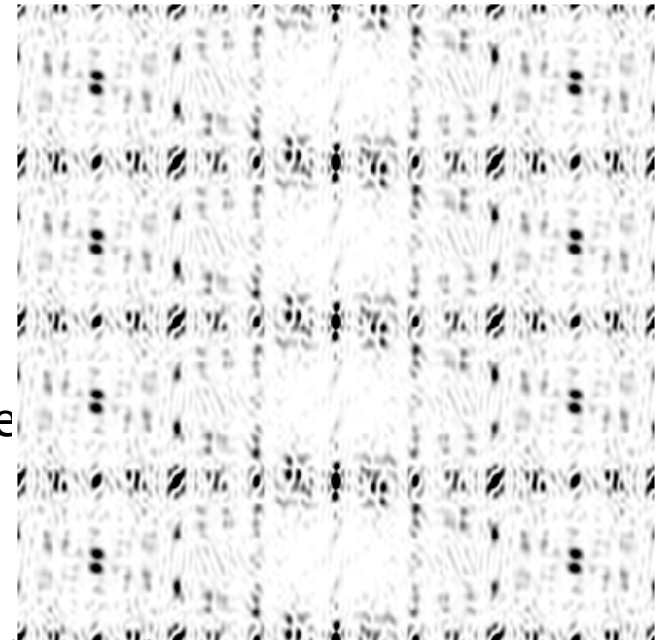
www.lks.physik.uni-erlangen.de/diffraction/index.html

DISCUS source code at:

Releases	github.com/tproffen/DiffuseCode/releases
Source Code	github.com/tproffen/DiffuseCode

DISCUS Workshop July 16 to July 20, 2018
Erlangen, Germany

<https://www.icsp.nat.fau.eu/neder-group/discus-home/>



F^{-1} THANK YOU



Interactive Tutorial on Diffraction

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