

Example 12.10: NiTe₂O₅

Last update 19.6.2024

Data	Topic	Level
Neutron single crystal	Simple antiferromagnetic structure ($k = 0$)	Intermediate

Input data

Data

Single crystal data collected on DEMAND at HFIR, ORNL neutron source at 4 K

Input files

NiTe2O5_nuc4K.int (single crystal data from FullProf)

NiTe2O5_powder X-ray.cif (nuclear structure)

Additional information

Chemical formula: NiTe₂O₅

Antiferromagnetic: $k=(0,0,0)$

References

J. H. Lee, M. Kratochvílová, H. Cao, Z. Yamani, J. S. Kim, J-G. Park, G. R. Stewart, Y. S. Oh (2019), Phys. Rev. B 100, 144441

Highlights

In this example, a single crystal data file from FullProf is imported. Corrections for $\lambda/2$ and different extinction models are performed.

Instructions

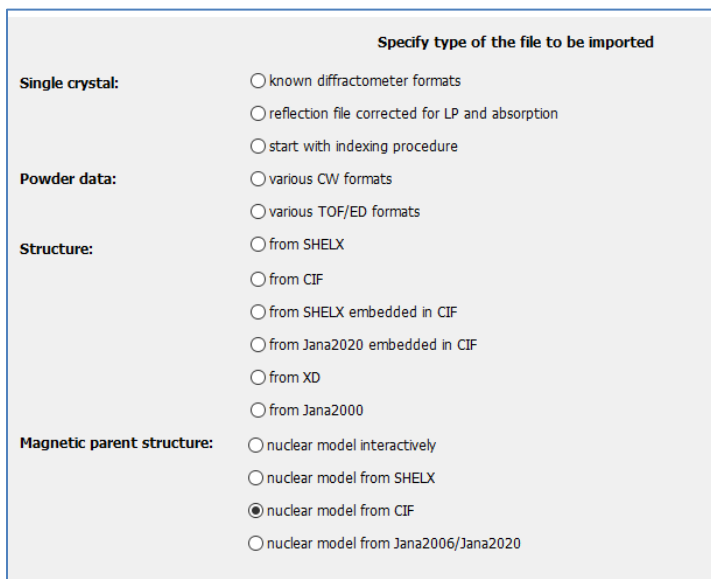
1. Data import

Start Jana2020

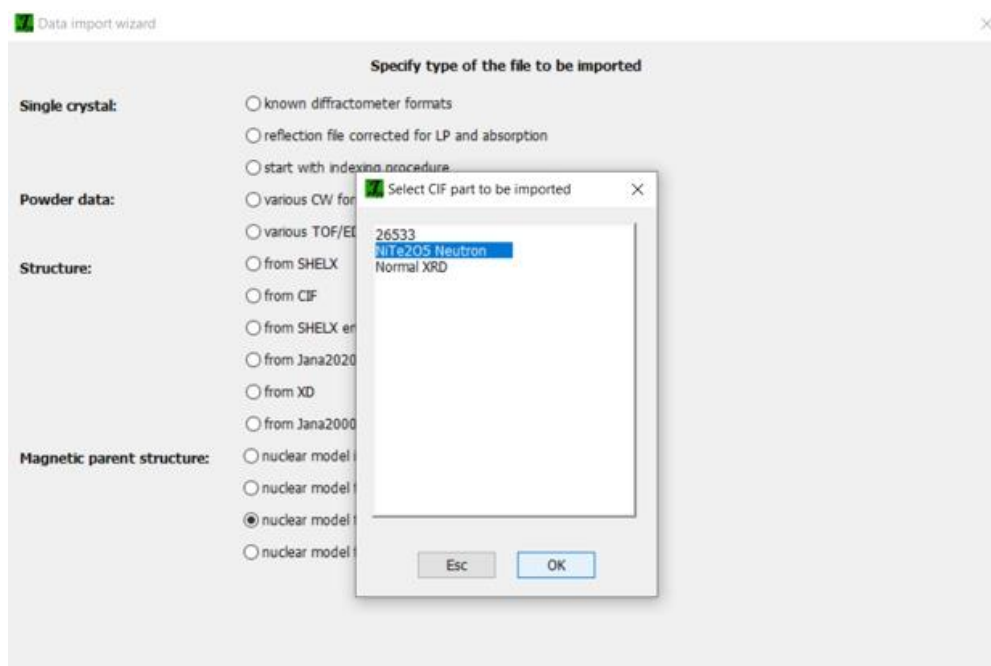
In the Main menu bar, use “Structure → New” and open new structure NiTe2O5 in the directory of Example 12.9

[On the screen: Specify type of the file to be imported]

Select “Magnetic parent structure: nuclear model from CIF”; NEXT



In Windows file explorer: select the input file NiTe2O5_powder X-ray.cif; Open
Choose the CIF part to be imported: NiTe2O5 Neutron; OK



2. Define the magnetic propagation vector and form factors

[On the screen: Define magnetic propagation vector(s) and form factors]

Set the “Number of superimposed IRs:” to ‘1’ and select the propagation magnetic vector to be “GM, k19 (0,0,0)” (if not there by default)
Select “Atom type” Ni; check “Use as a magnetic atom” and for “Magnetic formfactor <j0>” select “Ni2+”

Data import wizard

Define magnetic propagation vector(s) and form factors

Number of superimposed IRs: 1 GM, k19 (0,0,0) Qa 0 Qb 0 Qg 0

Formula - list of atomic types: Te16 Ni8 O40

Formula units: 1 Calculate density

Atom type: Ni Own scattering length for neutrons Use as a magnetic atom

Own form factors Edit

Magnetic formfactor <j0> Ni2+ + Magnetic formfactor <j2> + Magnetic formfactor <j4>

Magnetic formfactor <j0>+c<j2> Edit

NEXT;

[On the screen: Information]

The parent structure is now created.

NEXT;

3. Import of the reflection file

[On the screen: Specify type of the file to be imported]

Select “Single crystal: Reflection file corrected for LP and absorption”; NEXT
Select the format “From FullProf file”; Check that the name of the input file is
“NiTe2O5_nuc4K.int”; NEXT

[On the screen: Complete/correct experimental parameters]

Check that wavelength is 1.5503; Temperature is 4 K

Data import wizard

✕

Complete/correct experimental parameters

Cell parameters:

Number of input indices:

Radiation:

X-rays

Neutrons

Electrons

Kalpha1/Kalpha2 doublet

Wavelength:

Data collection details:

Temperature:

NEXT; NEXT; OK

[On the screen: Information]

FINISH

[On the screen: Data repository]

OK

4. Testing different irreps to get the best model

[On the screen: Information]

OK

[On the screen: List of irreps and corresponding kernel symmetries]

Representation analysis

✕

List of irreps and corresponding kernel symmetries:

Representation	Dimension	Shubnikov space group	Axes	Origin shift	
mGM1+	1	Pnma	(1,0,0 0,1,0 0,0,1)	(0,0,0)	<input type="button" value="Details"/>
mGM2+	1	Pn'm'a	(1,0,0 0,1,0 0,0,1)	(0,0,0)	<input type="button" value="Details"/>
mGM3+	1	Pnm'a'	(1,0,0 0,1,0 0,0,1)	(0,0,0)	<input type="button" value="Details"/>
mGM4+	1	Pn'ma'	(1,0,0 0,1,0 0,0,1)	(0,0,0)	<input type="button" value="Details"/>
mGM1-	1	Pn'm'a'	(1,0,0 0,1,0 0,0,1)	(0,0,0)	<input type="button" value="Details"/>
mGM2-	1	Pnma'	(1,0,0 0,1,0 0,0,1)	(0,0,0)	<input type="button" value="Details"/>
mGM3-	1	Pn'ma	(1,0,0 0,1,0 0,0,1)	(0,0,0)	<input type="button" value="Details"/>
mGM4-	1	Pnm'a	(1,0,0 0,1,0 0,0,1)	(0,0,0)	<input type="button" value="Details"/>

Notes

This window has an informative character. You can see here all irreps leading to magnetic ordering. The "Details" buttons provide more information about an individual irrep and its connection to the kernel magnetic symmetry.

NEXT

[On screen: List of kernels and epikernels]

The eight orthorhombic Shubnikov space groups are selected by default (if not, just click over one of them to select the block); NEXT

Notes

This will pass all the magnetic space groups to further testing.

[On the screen: Select Shubnikov space group]

Notes

For each space group, the components allowed for the magnetic moment of Ni are indicated in the column 'Moment'. Five of those groups have global moment zero corresponding to antiferromagnetic configurations.

Select the first Shubnikov group *Pnma*

Representation analysis

Select Shubnikov space group :

Shubnikov space group	Atom	Moment	Global	sin(2.Pi.x4)	cos(2.Pi.x4)
Pnma	Ni1	(M,M,M)	(0,0,0)		

Information: The symbol "M" indicates that the component can have non-zero value. For more details press the button "Show details". Magnetic moments are expressed in the parent cell.

Show details

Start graphic simulation

Start profile simulation

Continue with the selected Shubnikov space group

5. Refinement of the magnetic structure in *Pnma* (mGM1+) and testing of the different extinction models

Press 'Continue with the selected Shubnikov space group'

[On the screen: Select structure name]

Use the default name i.e. NiTe2O5_01; Save

Notes

The test runs under the new job name in a new window to keep the parent structure unchanged. The Jana window with the parent structure remains open.

Jana2020 makes averaging of the data automatically, contrary to FullProf that uses non-averaged data. To make a comparison, we can remake the refinement file.

[On the screen: Basic window of Jana2020]

In the Command tree, expand "Reflection file" and open "Create refinement reflection file"

[On the screen: Reflection file wizard]

NEXT

[On the screen: Import/Statistics-obs/all]

OK

[On the screen: Reflection file wizard: Process refinement file]

Choose "Use non-averaged data", NEXT; FINISH

[On the screen: Basic window of Jana2020]

In the Command tree, expand "Refinement" and open "Refinement commands"

[On the screen: Refinement commands]

On the page "Restrains/Constraints", select "Restrictions" and ensure that all atoms are restricted to have identical ADP parameters and not restricted occupancies ("restric *12"); OK

On the page "Restrains/Constraints", select "Fixed commands" and ensure that all atomic positions are fixed in their current values ("fixed xyz *")

OK

Notes

For this specific case we will see that the positions of atoms are not correct and that should be refined later.

On the page "Basic", check that the "Number of cycles" is set to 100 and the "Damping factor" to 0.1

OK; YES+START to start the refinement;

Notes

R factors : [582=564+18/5]

GOF(obs)~16.0 GOF(all)~15.8

R(obs)~36.6 wR2(obs)~62.9 R(all)~36.8 wR2(all)~63.2

R factors for nuclear reflections : [472=466+6]

R(obs)~35.5 wR2(obs)~62.4 R(all)~35.7 wR2(all)~62.7

R factors for magnetic reflections : [110=98+12]

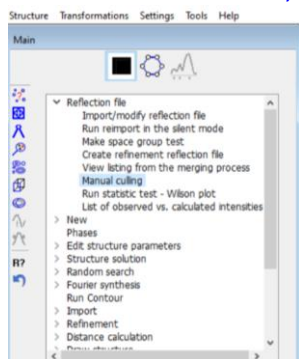
R(obs)~52.8 wR2(obs)~77.8 R(all)~52.8 wR2(all)~78.3

In the RF factors overview window, we can see that there is a line containing 'R factors for magnetic reflections'. In this case the propagation vector is zero, but because we have Pnma symmetry, we can create new reflections that were originally systematically absent for the nuclear structure. These reflections are very weak, leading to very high R values. To see better how this refinement works we can use manual culling.

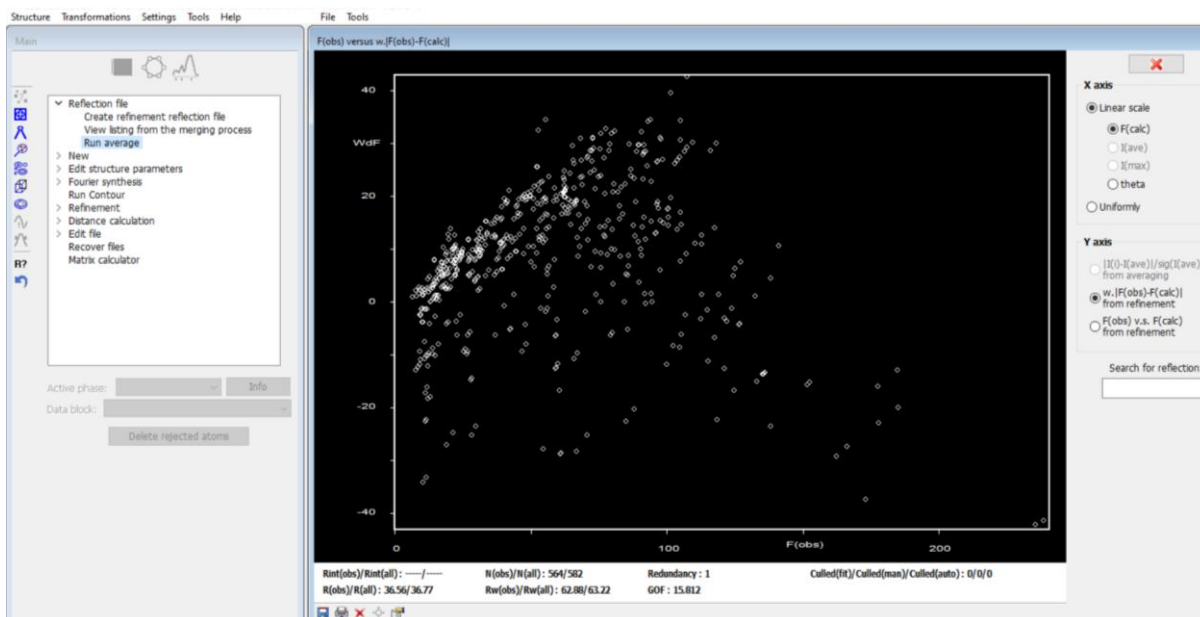
[There is a 'Serious warning' message concerning negative displacement parameters. We shall fix it later if needed.]

[On the screen: Basic window of Jana2020]

In the Command tree, expand "Reflection file" and open "Manual culling"



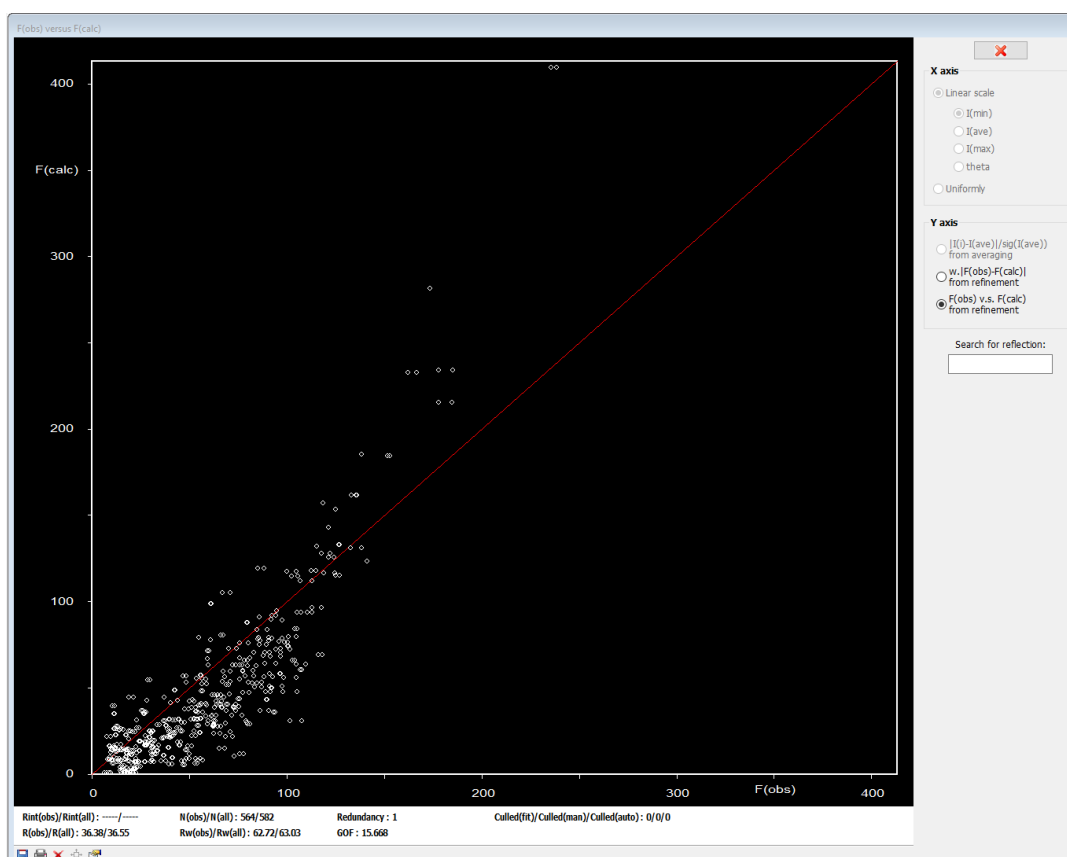
[On the screen: Fobs versus w|Fobs-Fcalc|]



Notes

In this window one can see the Fobs-Fcalc plot and follow its changes during the refinement.

On the right pane choose “Fobs v.s. Fcalc from refinement”



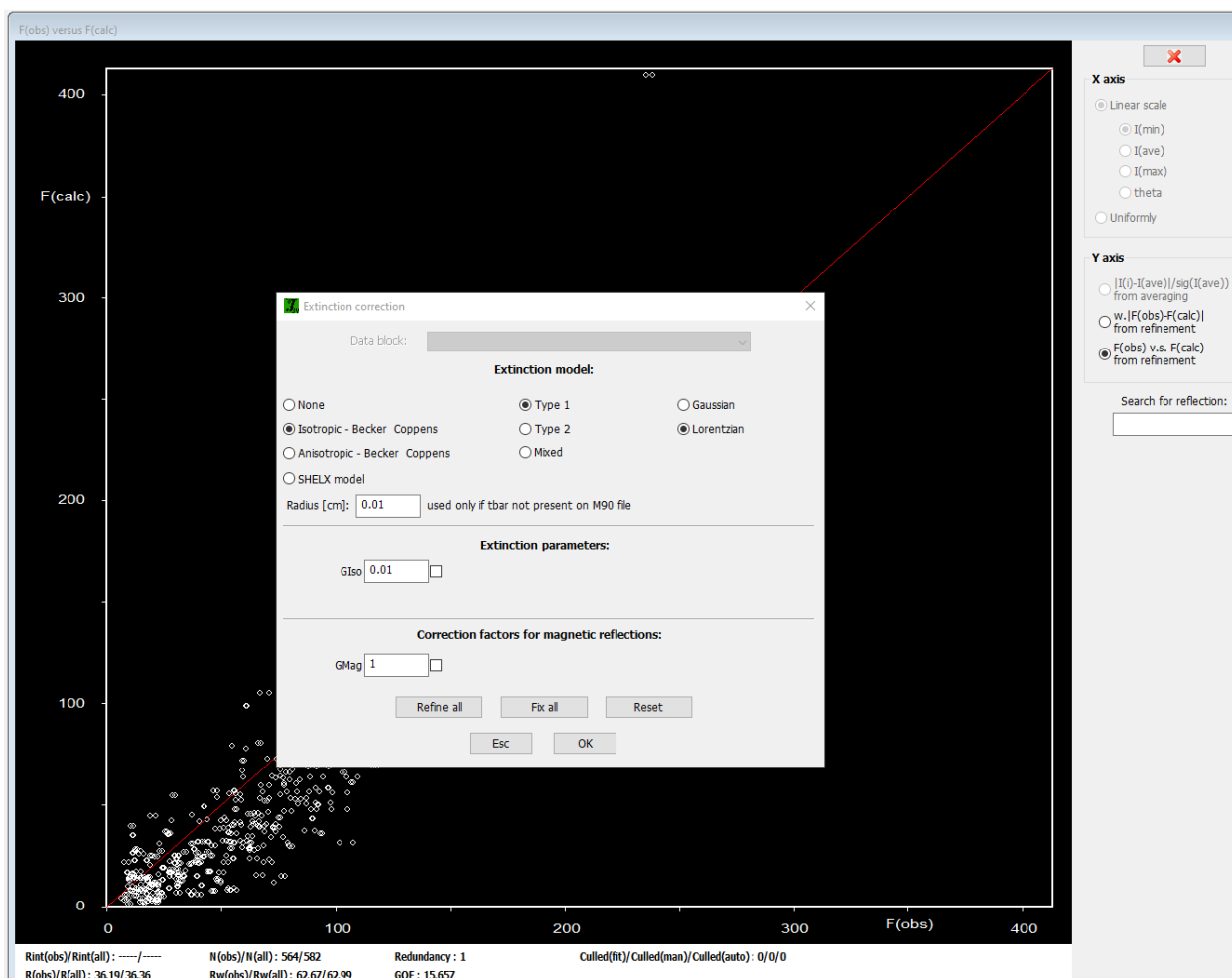
Notes

Here, it is clear that the observed intensity of strong reflections is much weaker than expected. An extinction correction is needed.

In the Command tree, expand “Edit structure parameters” and open “Edit extinction parameters”;

[On the screen: Extinction correction

Select “Isotropic”; “Lorentzian”; “Type 1”;



OK; YES; Run refinement or press 

Notes

R factors : [582=564+18/6]

$GOF(\text{obs}) \sim 13.7$ $GOF(\text{all}) \sim 13.8$

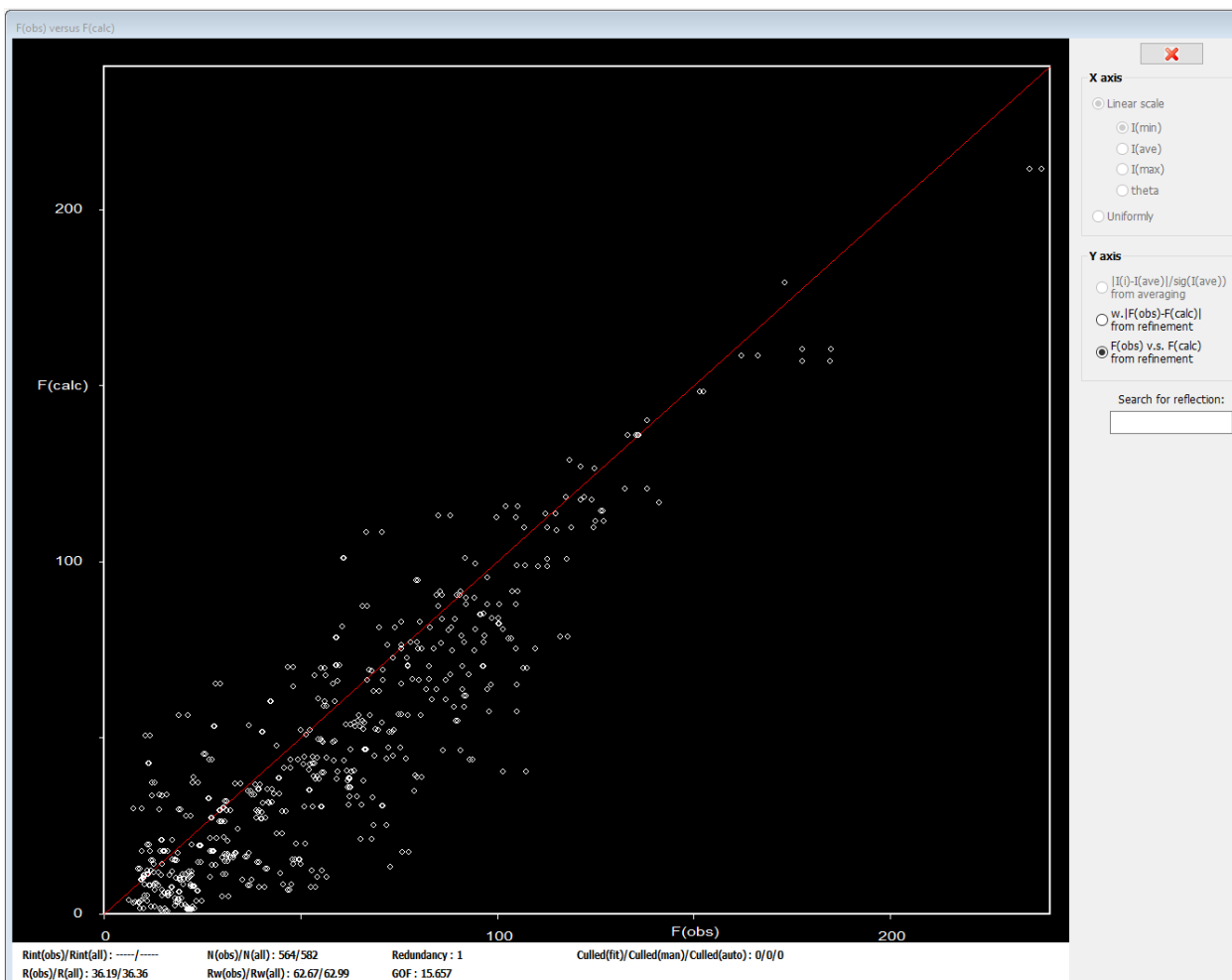
$R(\text{obs}) \sim 28.0$ $wR2(\text{obs}) \sim 53.8$ $R(\text{all}) \sim 28.4$ $wR2(\text{all}) \sim 54.7$

R factors for nuclear reflections : [472=466+6]

$R(\text{obs}) \sim 26.8$ $wR2(\text{obs}) \sim 53.6$ $R(\text{all}) \sim 27.1$ $wR2(\text{all}) \sim 54.5$

R factors for magnetic reflections : [110=98+12]

$R(\text{obs}) \sim 45.4$ $wR2(\text{obs}) \sim 60.7$ $R(\text{all}) \sim 45.8$ $wR2(\text{all}) \sim 61.0$



Notes

From the plot and R factors, it is obvious that the refinement looks better. Nevertheless, the R factors are yet very large. We'll now try to refine the positions of atoms.

In the Command tree, expand "Refinement" and open "Refinement commands"

[On the screen: Refinement commands]

On the page "Restraints/Constraints", select "Fixed commands". Click over the line "fixed xyz *", and press "Disable"; OK

OK; YES+START

Notes

R factors : [582=564+18/32]

GOF(obs)~4.0 GOF(all)~4.0

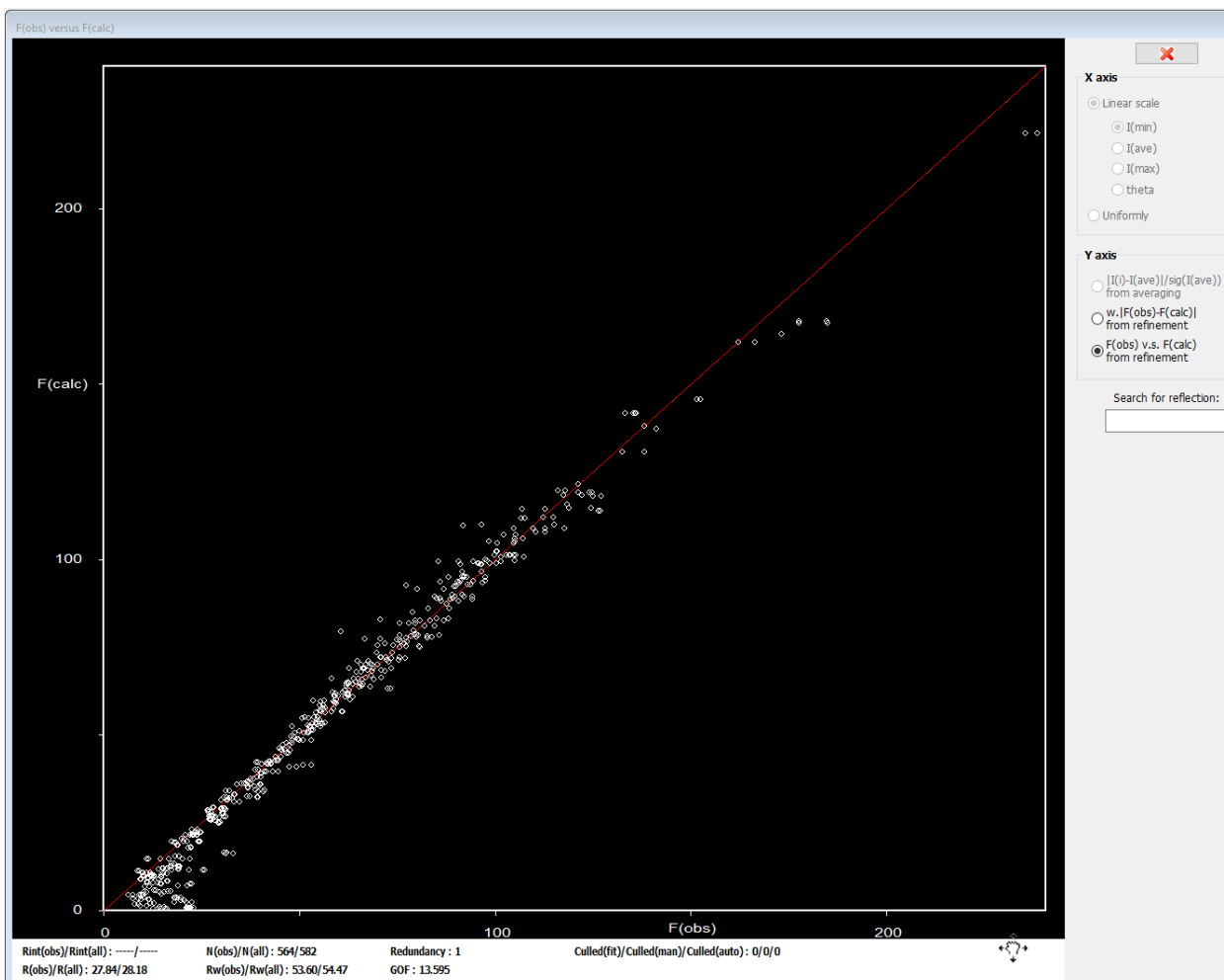
R(obs)~7.6 wR2(obs)~16.2 R(all)~7.9 wR2(all)~16.3

R factors for nuclear reflections : [472=466+6]

R(obs)~5.2 wR2(obs)~12.5 R(all)~5.2 wR2(all)~12.5

R factors for magnetic reflections : [110=98+12]

R(obs)~43.3 wR2(obs)~56.3 R(all)~43.7 wR2(all)~56.5



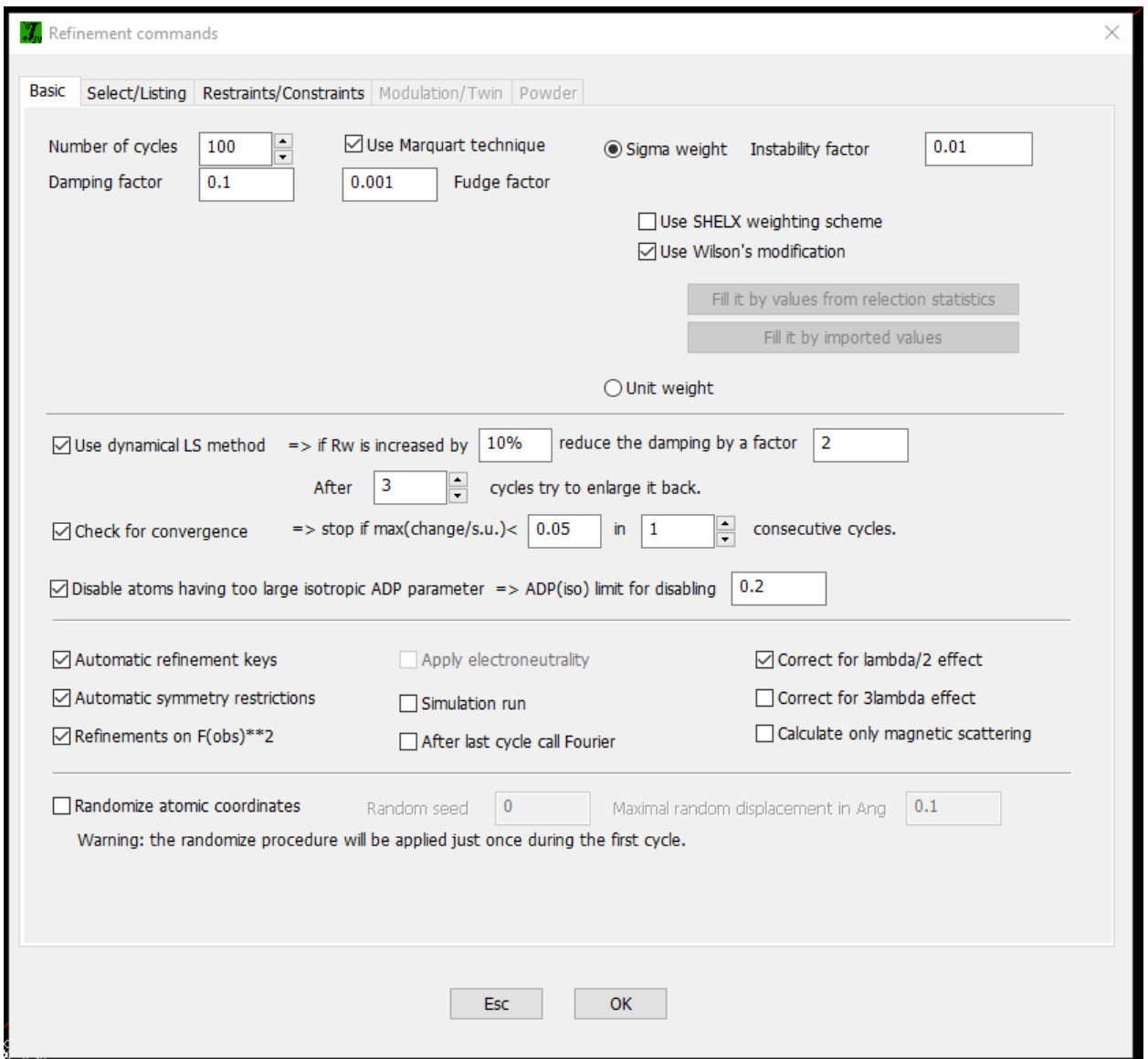
Notes

After refinement there is a great improvement in the model, especially for the nuclear reflections. Another important point for this data file is that it was measured on single crystal neutron diffraction for which there is a strong effect of $\lambda/2$ which needs to be accounted for.

In the Command tree, expand “Refinement” and open “Refinement commands”.

[On the screen: Refinement commands]

In the “Basic” tab, tick the option “Correct for lambda/2 effect”;



OK; YES

In the Command tree, expand “Edit Structure parameters” and open “Edit scale factors”.

[On the screen: Edit scale parameters]

Tick the parameter “sclam/2”. OK; YES; Start refinement

Notes

R factors : [582=564+18/33]

GOF(obs)~2.7 *GOF(all)*~2.7

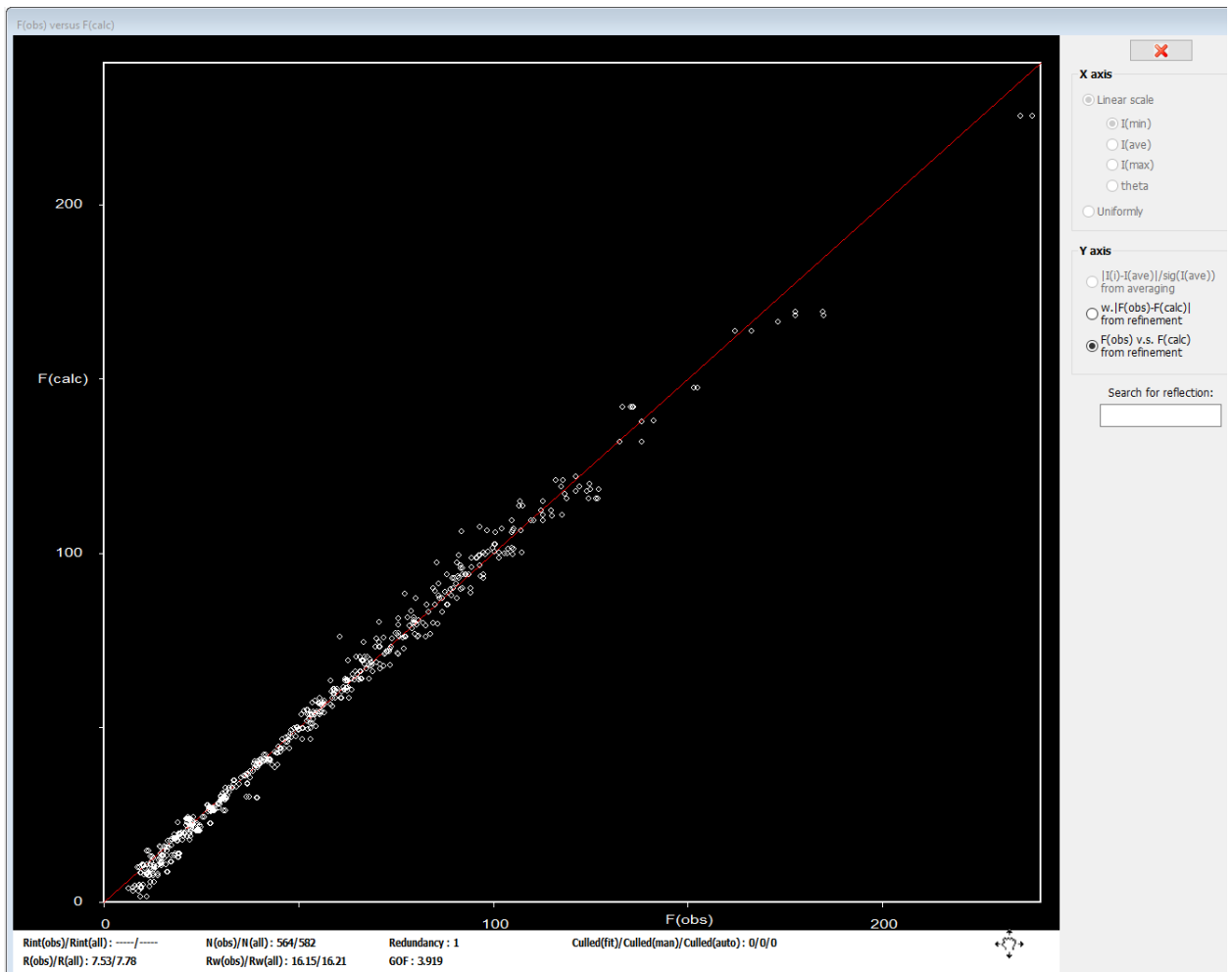
R(obs)~4.7 *wR2(obs)*~10.8 *R(all)*~4.9 *wR2(all)*~10.8

R factors for nuclear reflections : [472=466+6]

R(obs)~4.0 *wR2(obs)*~9.8 *R(all)*~4.1 *wR2(all)*~9.8

R factors for magnetic reflections : [110=98+12]

R(obs)~14.3 *wR2(obs)*~24.9 *R(all)*~15.5 *wR2(all)*~25.4



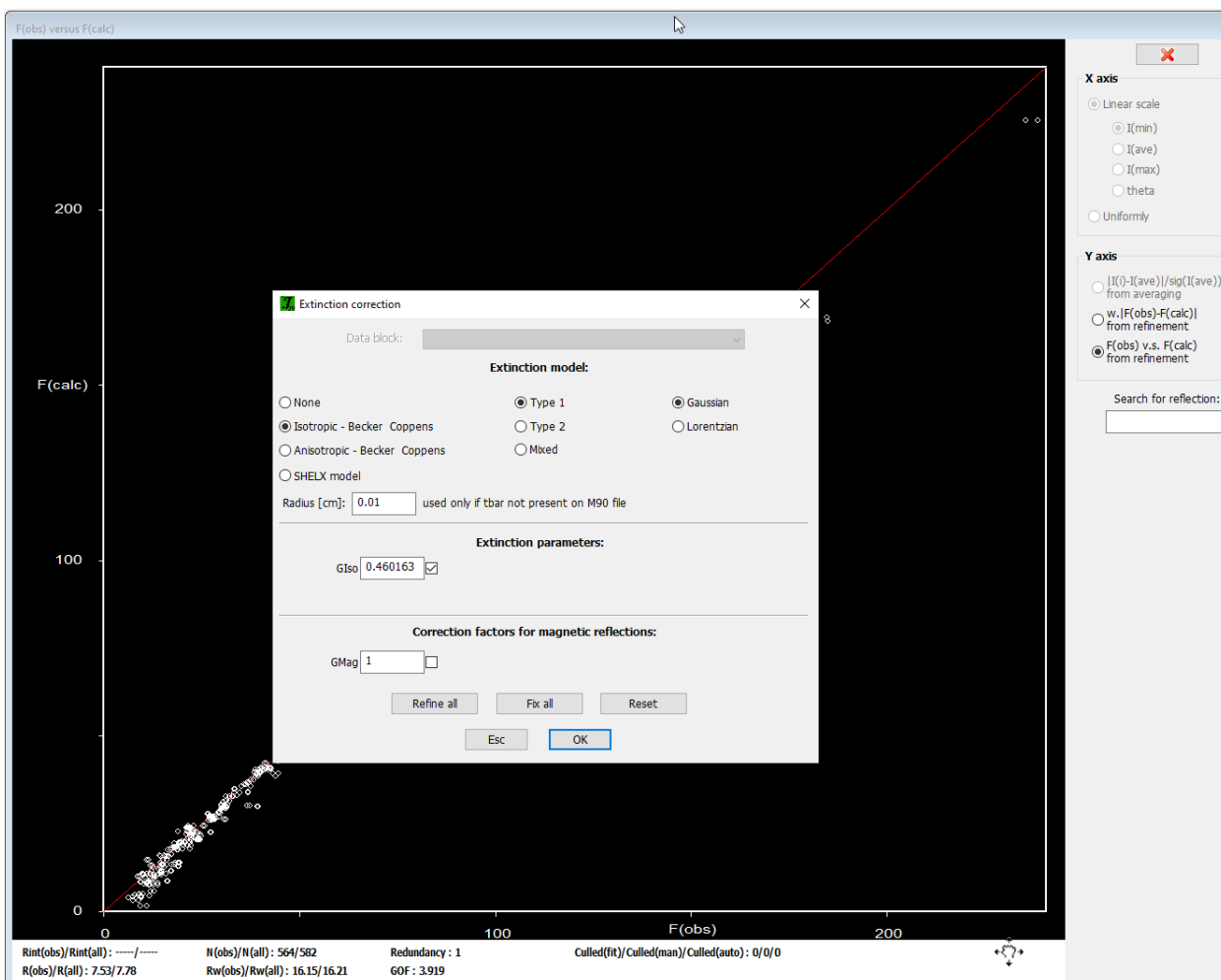
Notes

This result shows the importance of the extinction correction to reach good models. We shall now try another extinction model.

In the Command tree, expand “Edit structure parameters” and open “Edit extinction parameters”;

[On the screen: Extinction correction]

Select “Isotropic”; “Gaussian”; “Type 1”;



OK; YES; Start refinement

Notes

R factors : [582=564+18/33]

GOF(obs)~3.8 GOF(all)~3.8

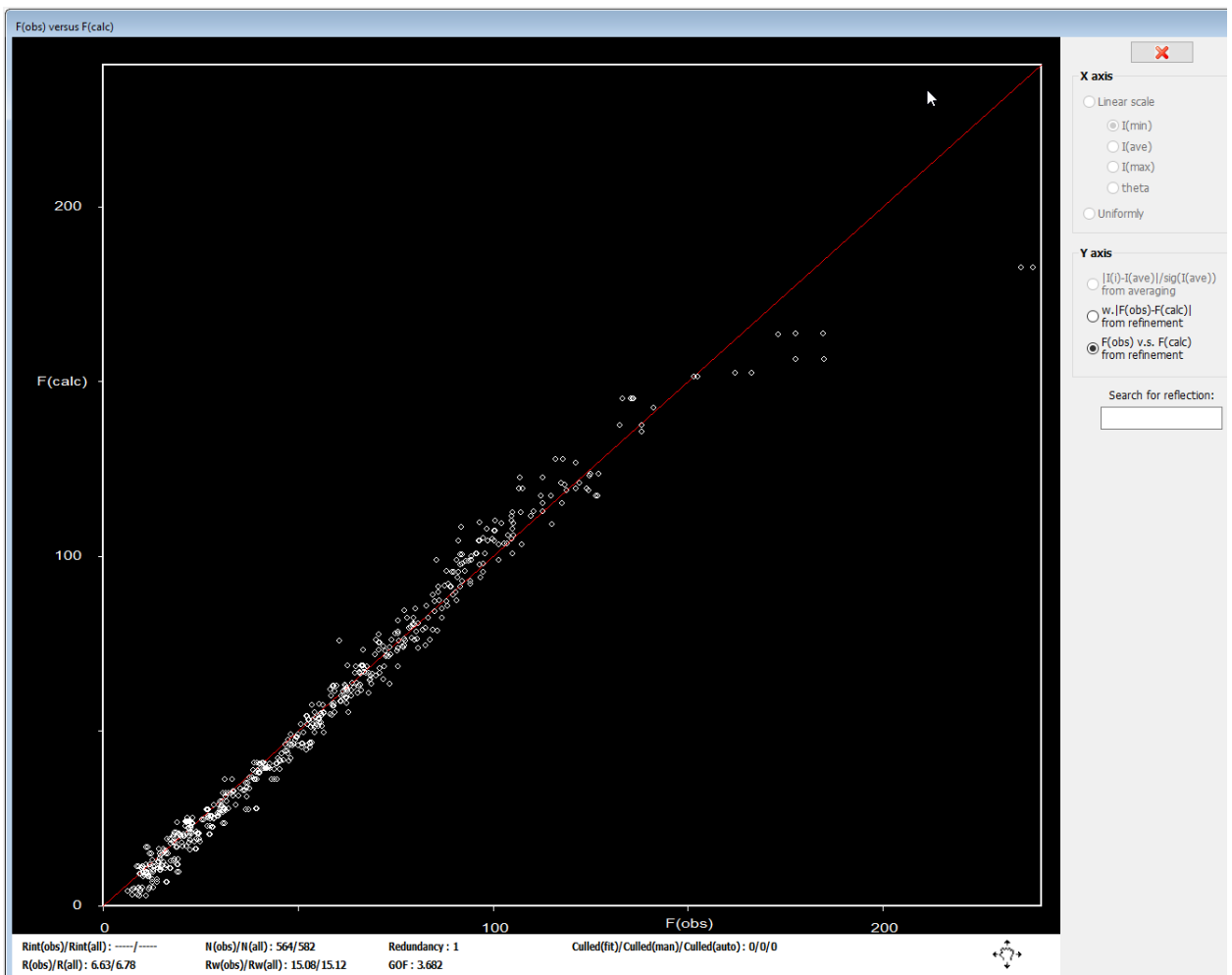
R(obs)~6.9 wR2(obs)~15.4 R(all)~7.0 wR2(all)~15.4

R factors for nuclear reflections : [472=466+6]

R(obs)~6.2 wR2(obs)~14.6 R(all)~6.2 wR2(all)~14.6

R factors for magnetic reflections : [110=98+12]

R(obs)~16.6 wR2(obs)~29.8 R(all)~17.7 wR2(all)~30.2



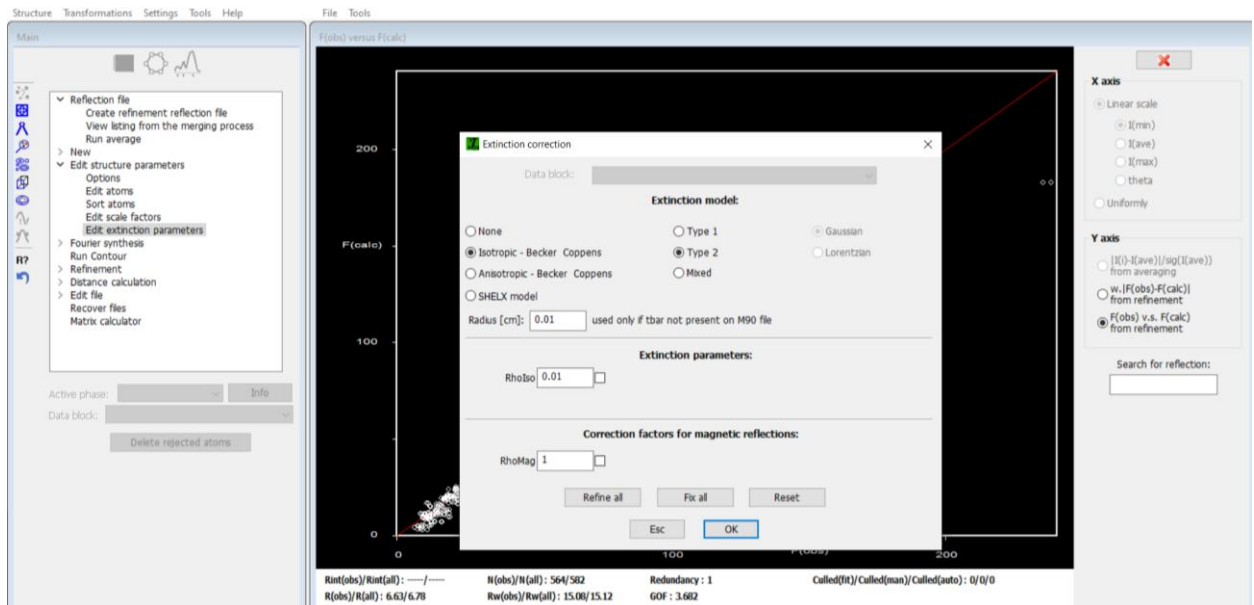
Notes

For the strong reflections, this extinction model gives much worse R factors than the previous one. We can try now also the extinction model type 2, which accounts for the dominating contribution of the crystal size to extinction.

In the Command tree, expand “Edit structure parameters” and open “Edit extinction parameters”;

[On the screen: Extinction correction]

Select “Isotropic”; “Type 2”;



OK; YES; Start refinement

Notes

R factors : [582=564+18/33]

$GOF(obs) \sim 3.0$ $GOF(all) \sim 3.0$

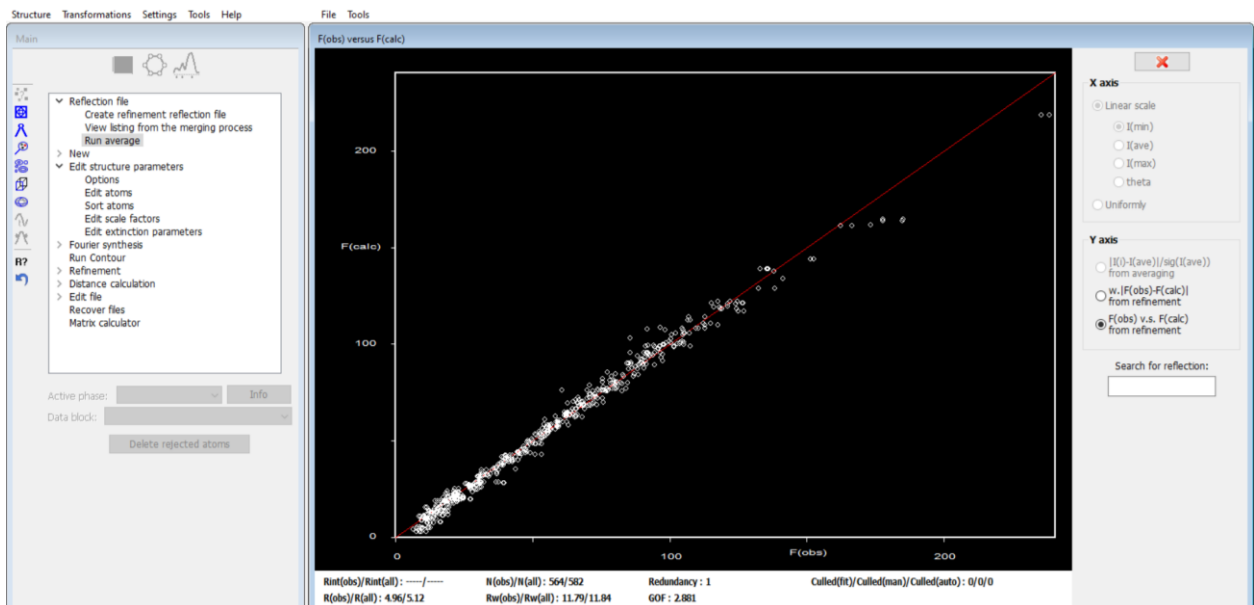
$R(obs) \sim 5.2$ $wR2(obs) \sim 12.1$ $R(all) \sim 5.3$ $wR2(all) \sim 12.1$

R factors for nuclear reflections : [472=466+6]

$R(obs) \sim 4.3$ $wR2(obs) \sim 10.6$ $R(all) \sim 4.4$ $wR2(all) \sim 10.6$

R factors for magnetic reflections : [110=98+12]

$R(obs) \sim 17.1$ $wR2(obs) \sim 32.1$ $R(all) \sim 18.0$ $wR2(all) \sim 32.4$



Notes

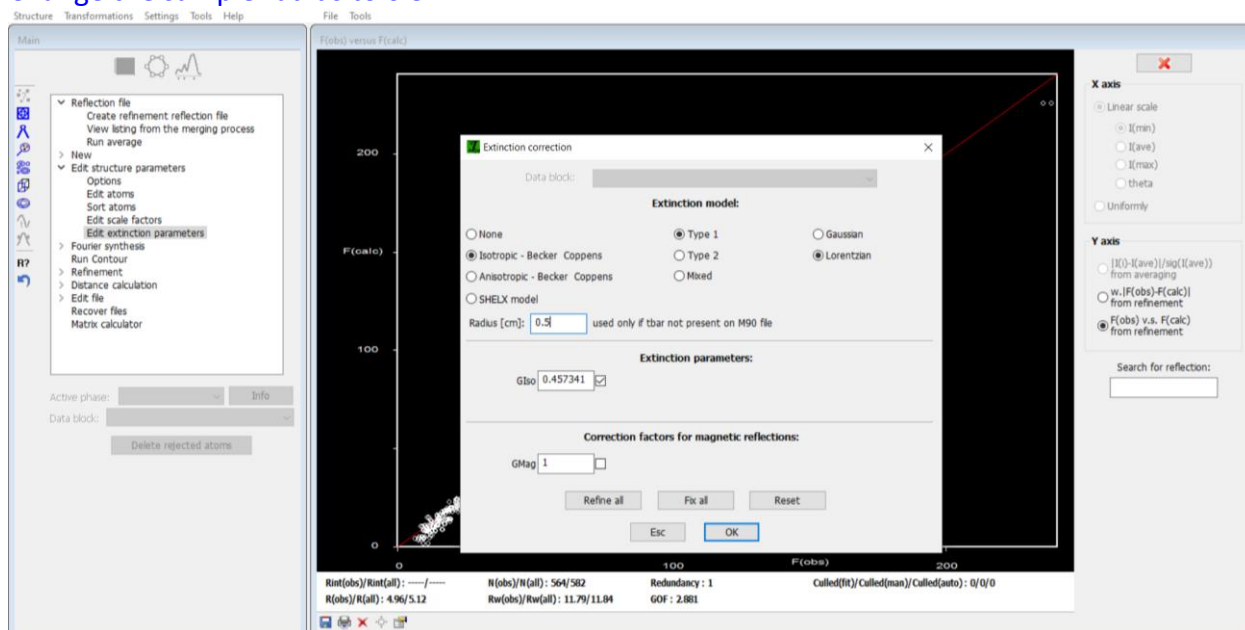
The best model for the extinction seems to be the Type 1 Lorentzian. We shall also change the sample radius to a larger value, as the samples for neutron diffraction are generally larger than for x-ray diffraction (the default value of 0.01).

In the Command tree, expand “Edit structure parameters” and open “Edit extinction parameters”;

[On the screen: Extinction correction]

Select “Isotropic”; “Lorentzian”; “Type 1”;

Change the sample radius to 0.5.



OK; YES; Start refinement

Notes

R factors : [582=564+18/33]

$GOF(obs) \sim 2.7$ $GOF(all) \sim 2.7$

$R(obs) \sim 4.7$ $wR2(obs) \sim 10.8$ $R(all) \sim 4.9$ $wR2(all) \sim 10.8$

R factors for nuclear reflections : [472=466+6]

$R(obs) \sim 4.0$ $wR2(obs) \sim 9.8$ $R(all) \sim 4.1$ $wR2(all) \sim 9.8$

R factors for magnetic reflections : [110=98+12]

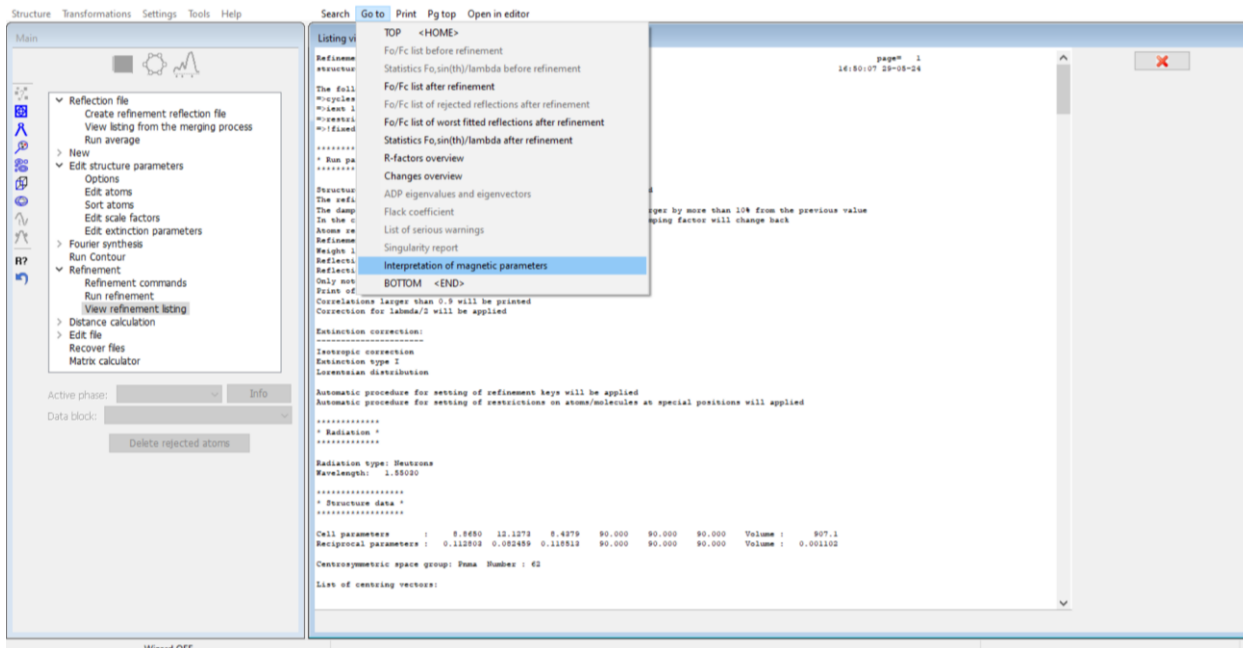
$R(obs) \sim 14.3$ $wR2(obs) \sim 25.0$ $R(all) \sim 15.5$ $wR2(all) \sim 25.4$

We can now check the magnetic moment of Ni after the best extinction correction.

In the Command tree, expand “Refinement” and open “View refinement listing”;

[On the screen: Listing viewer from refinement]

In the menu, select “Go to → Interpretation of magnetic parameters” and check the refined magnetic moments $Mx0$, $My0$, and $Mz0$ for the Ni atom.



```

*****
* Interpretation of magnetic parameters *
*****

                                Magnetic moments

Atom                along a        along b        along c        length
Ni1                  0.30 (5)        1.97 (5)        0.31 (4)        2.02 (8)

                                Sum of magnetic moments over the whole cell


Atom                along a        along b        along c
Ni1                  0.00 (15)       0.00 (13)       0.00 (10)
Summa                0.00 (15)       0.00 (13)       0.00 (10)

Program started at 15:33:41 ended   at 15:33:58 cpu time :    14.70

```

Notes
The refined components for the magnetic moment of Ni are similar to those refined in FullProf.

6. Visualization of the refined model with JanaDraw

Close the refinement list and the plot panel by 
 [On the screen: Basic window]

Change focus to "Janadraw"



[On the screen: JanaDraw]

Fill the unit cell by the button 

In the right pane, select “View along” c;

Adjust the figure by the button  and remove all bonds by the button 

