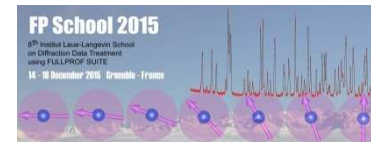


WinPLOTR :

a graphical tool for powder diffraction data analysis



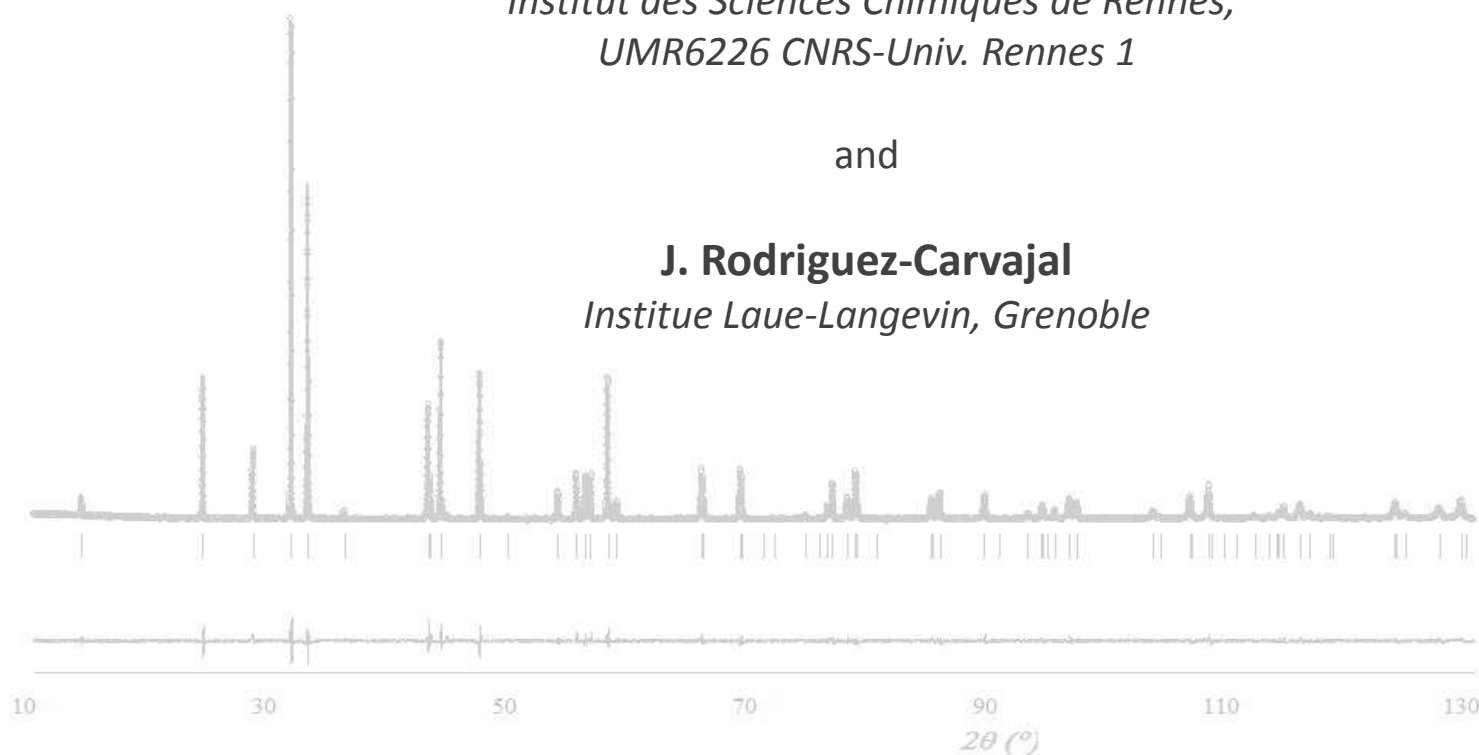
T. Roisnel

*Centre de Diffractométrie X (CDIFX)
Institut des Sciences Chimiques de Rennes,
UMR6226 CNRS-Univ. Rennes 1*

and

J. Rodriguez-Carvajal

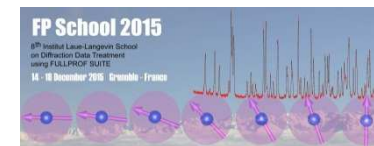
Institut Laue-Langevin, Grenoble



<http://www.cdifx.univ-rennes1.fr/winplotr/winplotr.htm>

(Tutorials and demos, download, what's new ?, links ...)

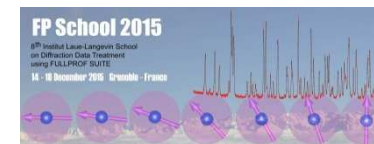
What can be done with WinPLOTR ?



WinPLOTR has been designed following two concepts :

- **WinPLOTR** is a program to plot and analyse powder diffraction patterns. It can be used to plot raw or normalized data files coming from different kinds of diffractometers (neutron and X-ray, conventional or synchrotron radiation) as well as profile refinement files (**.PRF**) created by the FullProf program.
- **WinPLOTR** can also be used as a Graphical User Interface for programs used frequently in powder diffraction data analysis (ex: FullProf, DicVol ...) or other external programs defined by the user.

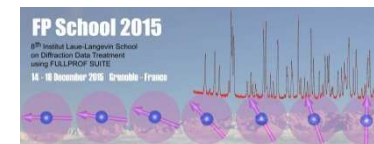
1. Plot and process powder diffraction data



Powder diffraction data features

- **Raw data, normalized ou refined data**
 - neutrons, X-Rays (conventional or synchrotron)
 - constant wavelength, dispersive energy, time of flight
- **Large number of data file formats (only ASCII) coming from several kinds of diffractometers** (commercial and home-made)
- **File conversion** : save data as **INSTRM0** FP format, multicolumns
- **Scattering space**
 - scattering variable : $2\theta(^{\circ})$ /time of flight ($\mu\text{sec.}$)/energy (KeV)
 - reciprocal space : $Q (\text{\AA}^{-1})$, $1/d (\text{\AA}^{-1})$, $\sin\theta/\lambda (\text{\AA}^{-1})$
 - direct space : $d (\text{\AA})$
- **Different Y scales** : linear, log, $\sqrt{}$, $I.Q^2$

1. Plot and process powder diffraction data

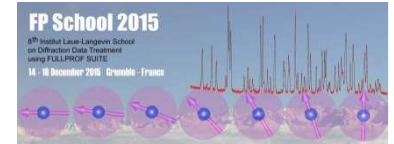


Graphical features

- **Graphics options (mouse handling)**
 - Cursor informations : X (including $2\theta/d$ values simultaneously) and Y values
 - Zoom, focus
 - Select and save points (countings, background)
- **Plot options**
 - Change colors, markers type and size, styles ...
 - Shift and offset the data (in X and/or Y) [pseudo 3D plots]
 - Hidden parts management
 - Error barrs
- **Graphics output files**
 - Bitmap
 - Postscript
 - WinPLOTTR output file (**.wp1**, **.pgf**)

...

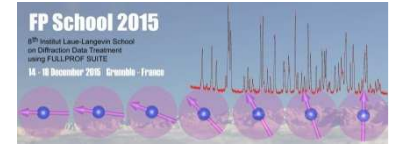
1. Plot and process powder diffraction data



Calculations features

- **Automatic search procedures:** background points, Bragg reflections
- **Background subtraction**
- **Calculations on diffraction patterns :**
 - patterns summation
 - pattern difference
 - smoothing
 - multiply X and Y
 - multiscan data normalisation
- **Profile fitting procedure** (interactif / automatic) (T.C.H. profile function)
- **Microstructural analysis** (Williamson Hall plots)

1. Plot and process powder diffraction data

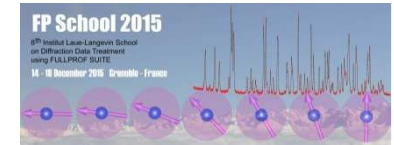


Calculations tools

- **FWHM calculation**
 - Cagliotti formula
 - T.C.H. formula
- **Crystallographic calculations :**
 - space groups
 - *hkl* list generation
 - unit cell volume
 - ...

...

2. GUI for external programs



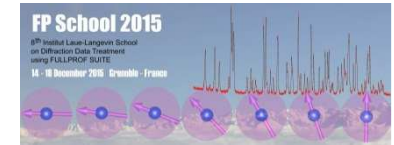
- **FullProf*** (JRC): profile refinement (Rietveld, whole pattern fitting)



- **.PCR** input file selection and edition
- FullProf launching : selection of input files (**.pcr** and diffraction data file)
 - **.PRF** file plotting (Rietveld plot) : Yobs, Ycalc, Yobs-Ycalc, Bragg positions
 - (hkl) information (Bragg peak position, indexation ...)
 - display (hkl) features
- Sequential FullProf refinement (ex: data=f(T)) and plot refined parameters

*included in the FPSuite package

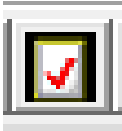
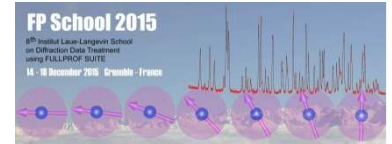
2. GUI for external programs



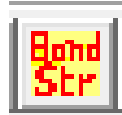
- Automatic indexation : **DICVOL** * (D. Louër), **TREOR** * (P.E. Werner), ...:
 - Peak search
 - Save Bragg peak positions as input files for indexation programs (dialog box)
 - **WinDICVOL/WinTREOR90** launching
 - If successful indexation : automatic creation of a **PCR** file for **FullProf** (« pattern matching » mode)
- **SuperCELL** * (JRC) : determination of a super unit cell
- **K-search** * (JRC) : search propagation wave vector components of a magnetic or structural modulated structure
 - Save Bragg peaks on difference pattern of a **PRF** file

*included in the FPSuite package

2. GUI for external programs



Check_Group* (JRC) : search of a space group from a list of integrated intensities



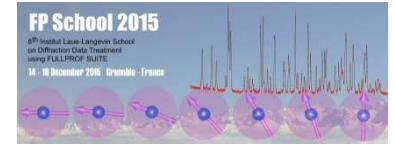
BOND_STR* (JRC) : distances, angles and bond valence calculations



EdPCR* (JGP-JRC):
 . editing and modifying **PCR** input file
 . Files conversion: **CIF** to **PCR**, **INS** to **PCR**

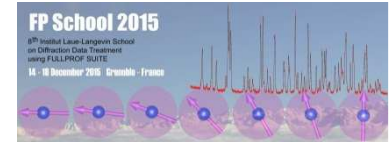
- Any program defined by the user (**winplotr.set** setting file)

winplotr.set setting file



- Located in the folder associated to the **WINPLOTR** environment variable
(ex: `c:\>fullprof_suite`)
- Definitions of :
 - arrays dimensions
 - external programs to execute through **WinPLOTR**
 - system applications : browser, editor ...
 - graphical plot options : colours, markers ...
 - instrumental parameters : λ , IRF
 - data files extensions
 - ...
- Accessible via the « **External applications / Edit a file / winplotr.set file** » menu option

winplotr.set setting file



```
*****
*
*      W i n P L O T R   s e t t i n g s
*
*****
```

[MAIN WINDOW POSITION AND SIZE]

```
0.100 0.100 0.800 0.800
```

```
.
.
.
```

[DIMENSIONS OF ARRAYS]

```
Max_Patterns = 20      ! max. number of patterns
Max_Points   = 60000   ! max. number of data points
Max_Refl     = 20000   ! max. number of Bragg reflections
```

```
.
.
```

[RUN PROGRAMS]

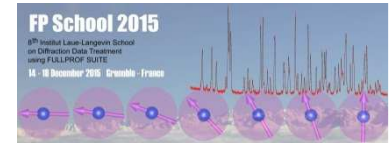
```
FullProf = wfp2k          ! FullProf program      ! WINDOWS VERSION
Edit = C:\Program Files\Keditw\KEDITW32.EXE ! My favorite editor ! WINDOWS VERSION
Dicvol = wdicvol06        ! Dicvol program      ! WINDOWS VERSION
SuperCell = SuperCell     ! SuperCell program   ! WINDOWS VERSION
Treor = wtoreo90          ! TREOR90 program     ! WINDOWS VERSION
ITO = wito15              ! ITO15 program       ! WINDOWS VERSION
MENDEL = mendel           ! MY PERIODIC TABLE ! WINDOWS VERSION
GFOURIER = gfourier       ! FOURIER PROGRAM     ! WINDOWS VERSION
@CALC_PDF = calc_pdf      ! PDF calculation     ! WINDOWS VERSION
@CRYSCALC = d:\progs\cryscalc ! Crystallographic calculator ! WINDOWS VERSION
@WINPLOTR2006 = D:\FullProf_Suite\WinPLOTR-2006.exe ! WinPLOTR_2006 ! Windows version
@K-search = k_search.exe  ! K_search            ! DOS
@VESTA = D:\progs\VESTA-win64\vesta.exe ! VESTA              ! W
@EXPO = D:\progs\EXPO2004\expo2004.exe ! EXPO               ! W
```

```
.
```

[SYSTEM APPLICATIONS]

```
Browser = "C:\PROGRAM FILES (X86)\MOZILLA FIREFOX\FIREFOX.EXE"
PostScriptViewer = "C:\Program Files\gs\gs9.14\bin\gs-win64.exe"
PDFreader = "C:\Program Files (x86)\Adobe\Reader 11.0\Reader\ACRORD32.EXE"
BMPviewer = "C:\WINDOWS\SYSTEM32\MSPAINTE.EXE"
CIFviewer = "D:\PROGS\CCDC\MERCURY 3.1\MERCURY.EXE"
```

winplotr.set setting file



```
[AFTER FULLPROF RUN]
plot_prf edit_pcr no_draw_fst
```

```
[MY WAVELENGTHS (A)]
1.54060 1.54060 1.00000 0.00000

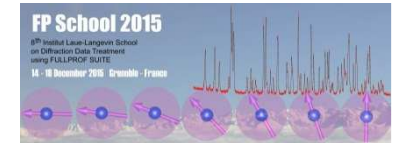
[MY RESOLUTION PARAMETERS (U,V,W,X,Y,Z)]
! HG**2= U.tan**2(theta) + V.tan(theta) + W
! HL = X.tan(theta) + Y/cos(theta) + Z
0.0100 -0.0100 0.0050 0.0000 0.0000 0.0000
```

```
[DATA FILES EXTENSIONS]
! format_number extension type
1 *.XY;*.DAT;*.BGR;*.EPF;*.XYN;*.XYS ! 1. X,Y data + INSTRM=10
2 *.DAT;*.SUB;*.SIM;*.BAC ! 2. INSTRM=0: Free F.(Ti,step,Tf)
3 *.DAT ! 3. INSTRM=1: Old D1A
4 *.DAT;*.D1B ! 4. INSTRM=3: D1B (ILL)
5 *.DAT ! 5. INSTRM=4: Brookhaven(Synchr.)
6 *.DAT ! 6. INSTRM=5: G4.1
7 *.DAT ! 7. INSTRM=6: D2B/3T2/G4.2
8 *.DAT ! 8. INSTRM=8: HRPT/DMC (PSI)
9 *.UXD ! 9. INSTRM=9: .UXD (D8 CSM)
10 *.DAT ! 10. INSTRM=11: Variable Time step
11 *.DAT;*.GSS;*.GSA ! 11. GSAS data
12 *.CPI ! 12. CPI (Xrays)
13 *.UDF;*.XRDML;*.JCP;*.CSV ! 13. PANalytical formats
14 *.DAT;*.GSS;*.GSA ! 14. INSTRM=14: ISIS normalized data
15 *.TXT ! 15. Rigaku RINT
16 *.SPC ! 16. DIFFAX (.spc)
```

```
[MY DEFAULT FORMATS (pattern, raw, Rietveld/profile)]
1 56 101

[MY COMMAND LINE DEFAULT FORMATS (.dat, .uxd, .prf)]
2 56 101
```

winplotr.set setting file



[BACKGROUND SCREEN]

```
background screen color: RGB(220,220,220)
background text color:   RGB( 60, 60, 60)
background plot color:   RGB(255,255,255)
plot frame color:        RGB( 0, 0, 0)
```

[EXCLUDED REGION COLOR]

```
RGB(192,192,192)
```

[PATTERNS PLOT OPTIONS]

! n	color	marker	size	style	pen_width
1	RGB(0, 0,255)	4	1.0	1	1
2	RGB(255, 0, 0)	4	1.0	1	1
3	RGB(0,128, 0)	4	1.0	1	1
4	RGB(255, 0,255)	4	1.0	1	1
5	RGB(128, 0, 0)	4	1.5	1	1

[PRF PLOT OPTIONS]

! n	color	marker	size	style	pen_width
1	RGB(255, 0, 0)	4	1.0	0	2
2	RGB(0, 0, 0)	4	0.0	1	2
3	RGB(0, 0,255)	4	0.0	1	1
4	RGB(0,128, 0)	8	3.0	0	1

```
write_Y_negative_graduations=0
```

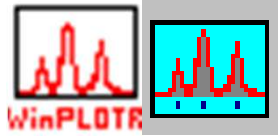
```
write_PRf_filename=0
```

```
exclude_CELL_from_title=1
```

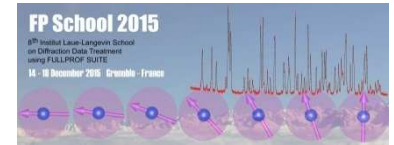
```
colored_tics=0
```

[TEXT FONTS]

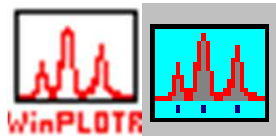
	Typeface	point	italic	underline	strikeout	bold
main title :	Arial	18	F	F	F	F
X legend :	Times New Roman	18	T	F	F	T
Y legend :	Courier New	18	T	F	F	T
X graduations:	Courier New	16	F	F	F	T
Y graduations:	Courier New	16	F	F	F	T
file name :	Courier	10	F	F	F	F
(hkl) indices:	Times New Roman	10	F	F	F	F
dialog boxes :	Times New Roman	10	F	F	F	F



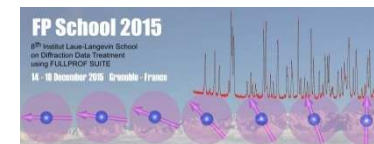
WinPLOTR demos



- Visualisation of (pseudo) 3D plots
- Run **FP** in sequential mode and plot refined parameters
- Profile fitting procedure
- Background points selection
- WinPLOTR tools
- Run WinPLOTR in command line



Visualisation of (pseudo) 3D plots



1. Powder diffraction data

- Create a buffer file containing the list of data files (common format) to plot, by using the

Tools / create buffer file menu option

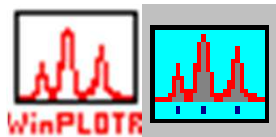
Or use the DOS command :

```
d:\data_3D>dir *.dat >> bufferfilename.buf /b
```

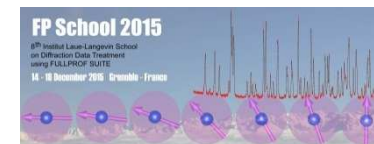
- Open a buffer file
- Select the common data file format



WinPLOTR

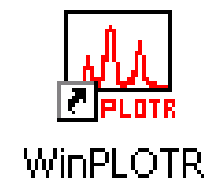
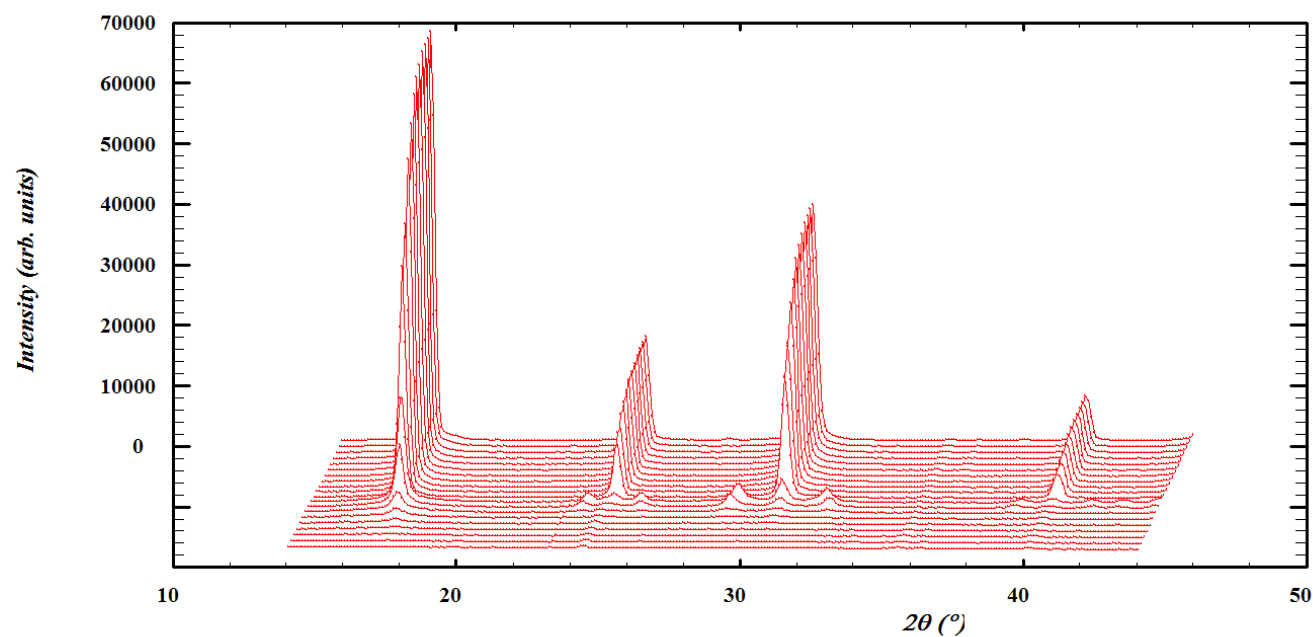


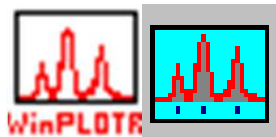
Visualisation of (pseudo) 3D plots



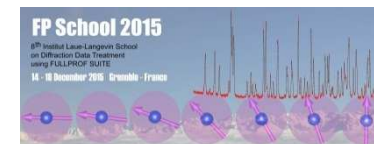
2. Graphical options in WinPLOTR :

- X and Y shifts
- Change colors, markers, style
- Hidden part

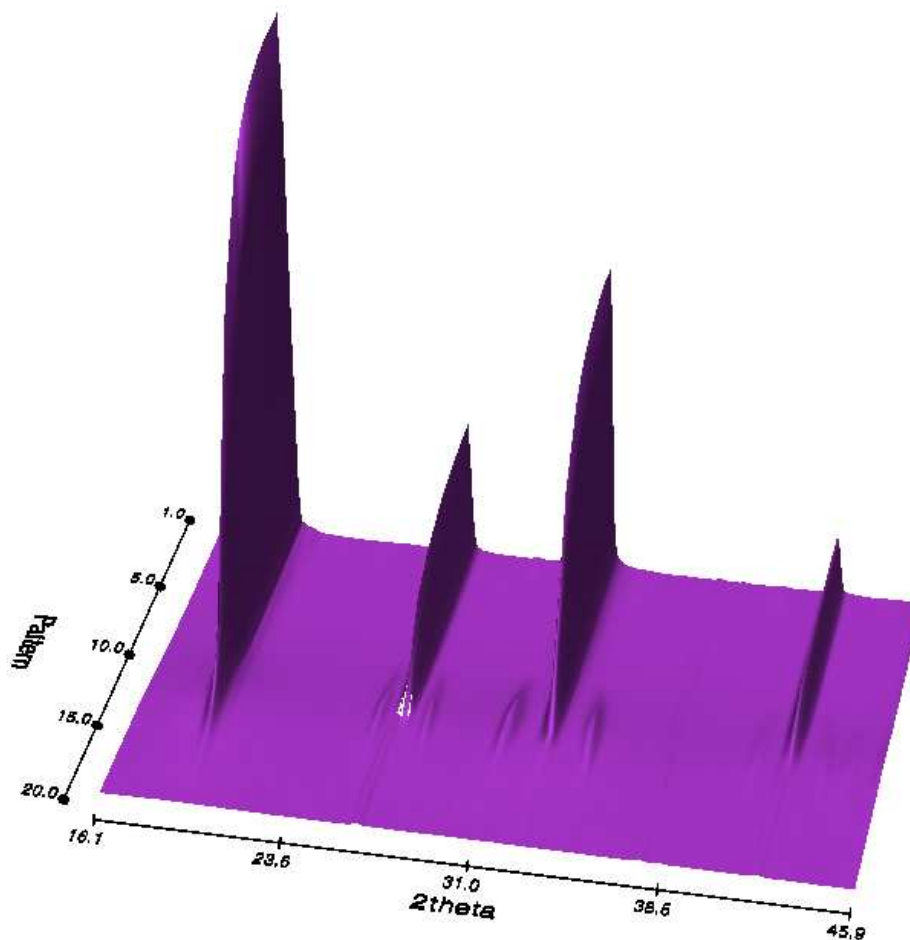


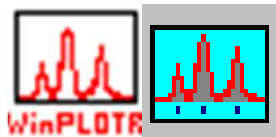


Or : 3D plots with WinPLOTR2006

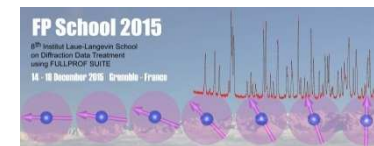


2. OpenGL 3D model





Run FullProf in sequential mode : plot of refined parameters



- Experimental requirements:

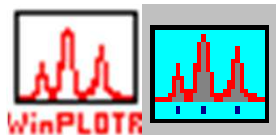
- . Diffraction experiment versus external parameter (T, P, time, ...)
- . Common format for all data files
- . Specific naming for data files

- FullProf specificities:

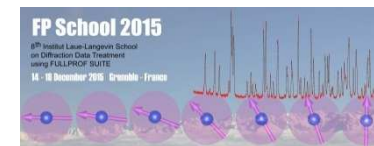
- . Only 1 starting PCR file (refine first temperature separately)
- . Profile matching, Rietveld, multiphases, microstructure ...
- . Distances calculations ...

All the patterns will be analysed in the same refinement conditions (background, physical model, number of phases, number of refined parameters...).

The evolution of the refined parameters has to be continuous ! Split temperature range if necessary.

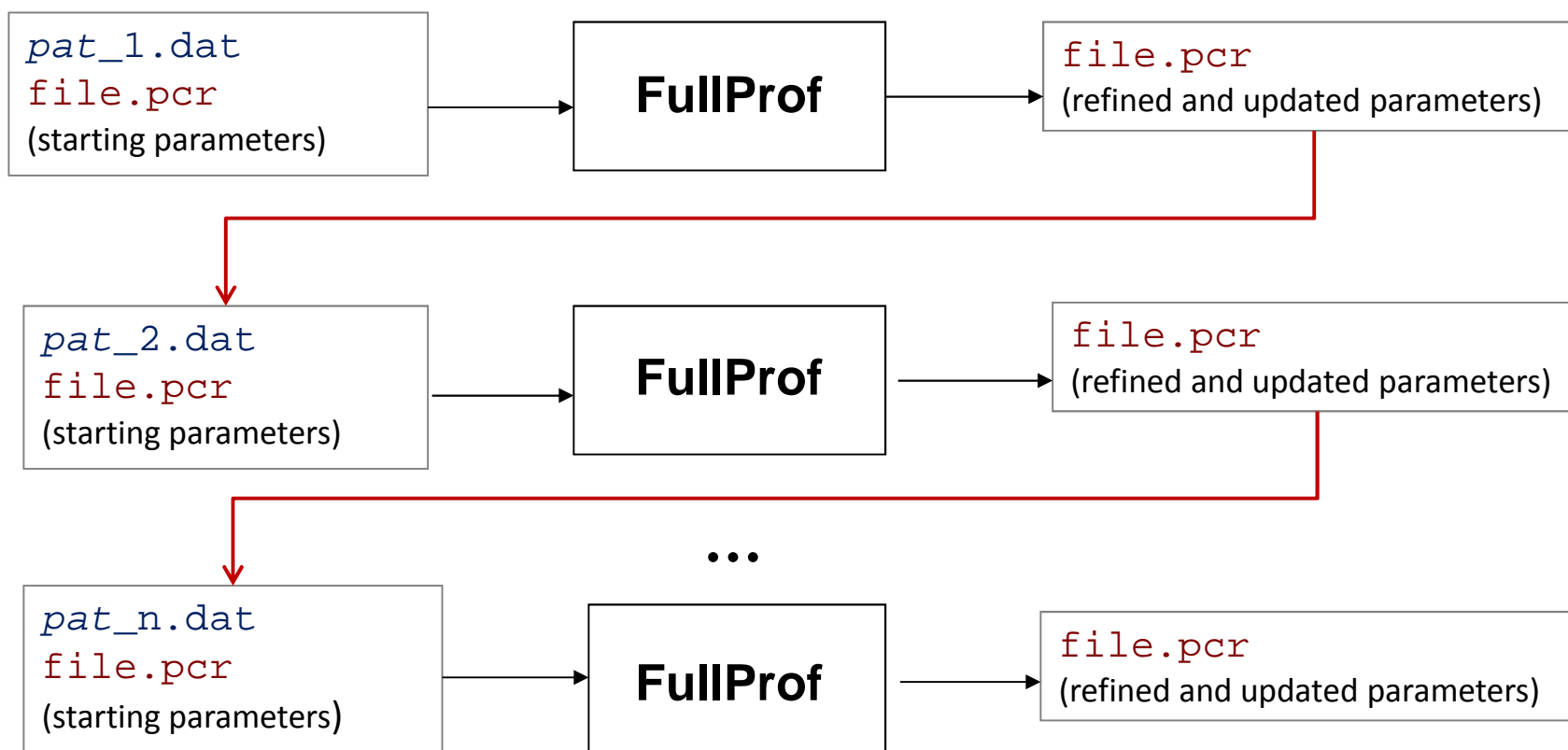


FullProf in sequential mode

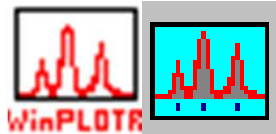


N diffraction data: *pat_1.dat, pat_2.dat ... pat_n.dat*

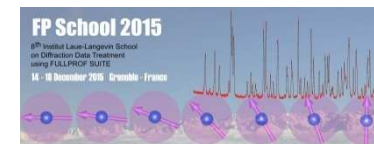
Only 1 pcr input file: *file.pcr* (starting parameters) **PCR=1**



All refined parameters during sequential run are stored in a **.SEQ** file



Prepare sequential FullProf launching setup



Sequential mode of FullProf:

Enter PCR file:

Enter data file code [.extension]:

Enter first numor :

Enter last numor :

☒ Save .PRF files

☐ Save .HKL files

☐ Save .MIC/SIZ/STR files

☒ Save .DIS files

☐ Generate .CALC files

☒ Plot sequential FullProf results

☐ Plot interatomic distances

☐ Plot HKL features

☐ Plot microstructural features

☒ Run Fullprof

☐ Create script for sequential mode

Variable external parameter (Temperature. ...):

Temperatures file:

PCR file

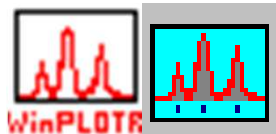
File name code

First and last numor

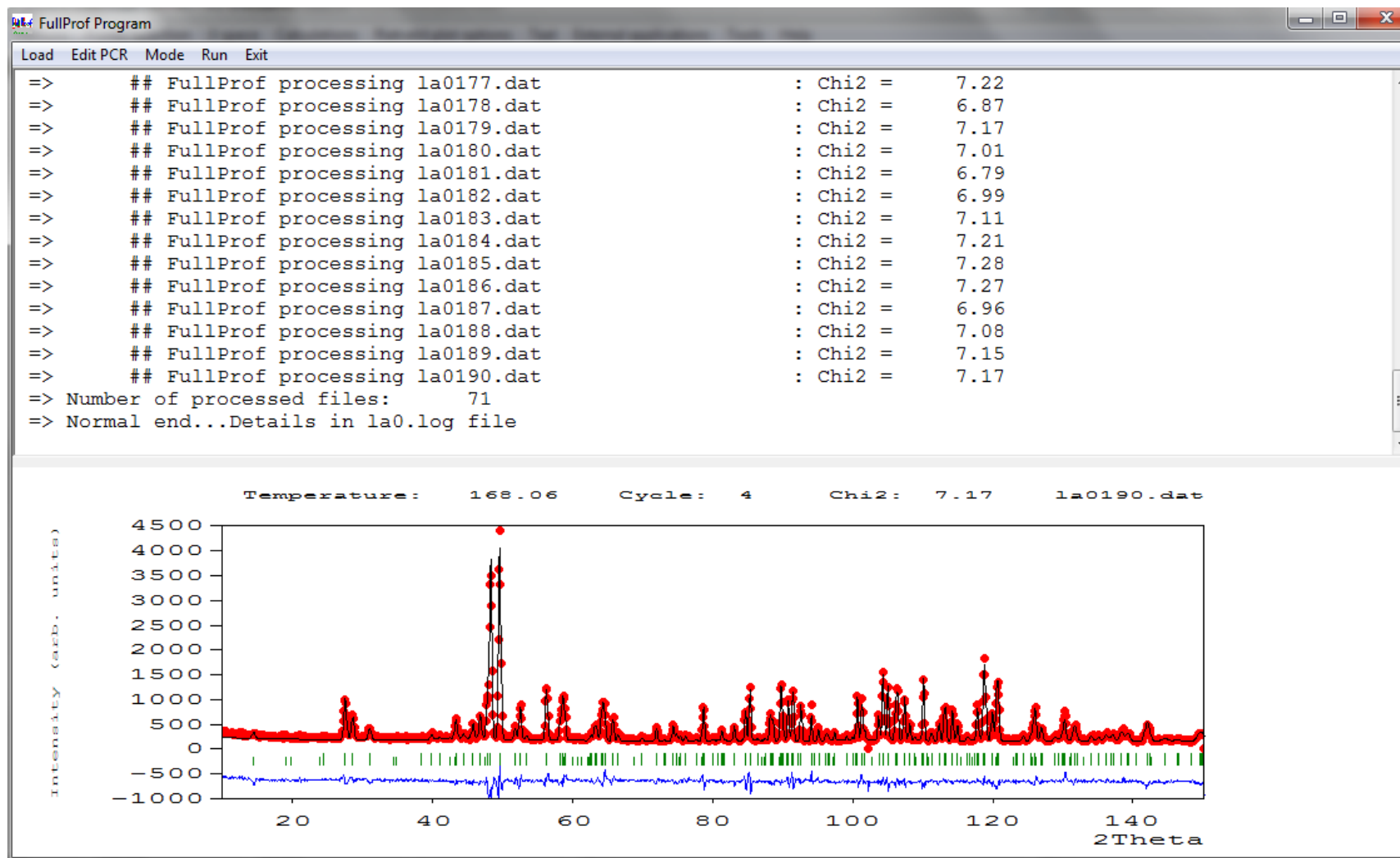
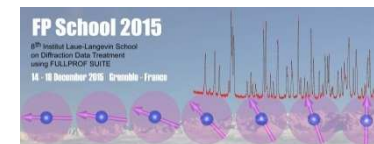
Output files

Results to plot

WinPLOTR @ FPSchool, ILL, 14-18 dec. 2015



Running sequential FullProf

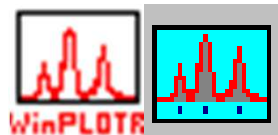


WinPLOTR @ FPSchool, ILL, 14-18 dec. 2015

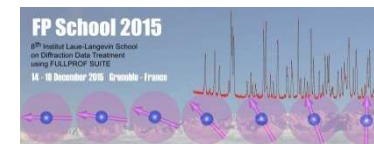
71 patterns (1535 points) / 21 refined parameters : 26 sec. of cpu time on my labtop !



WinPLOTR

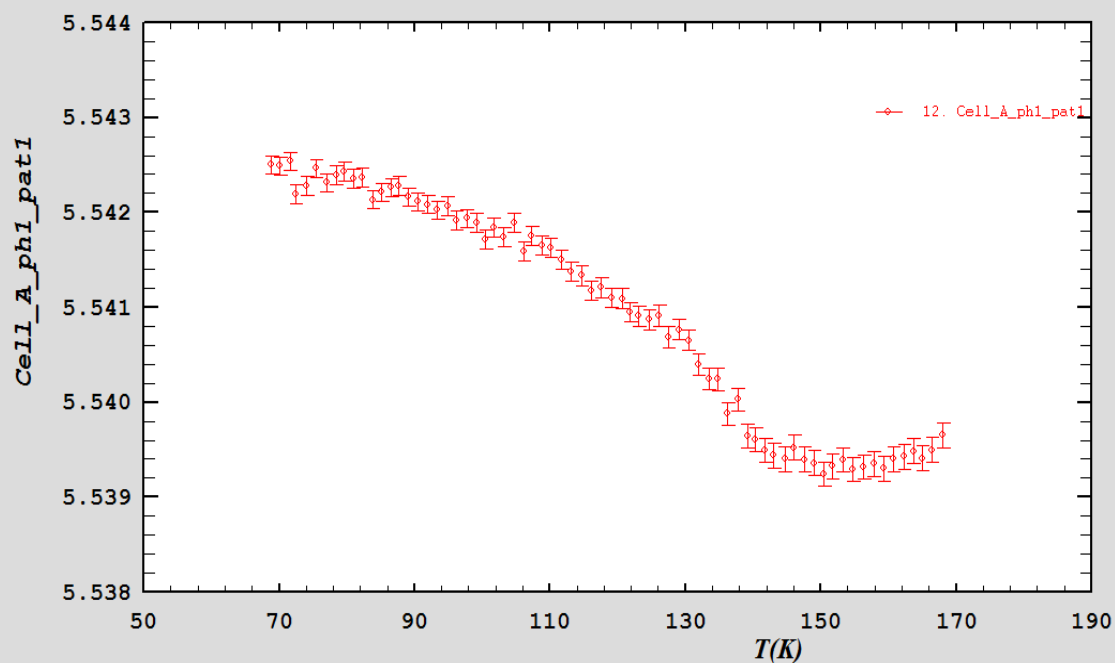


Plotting sequential FP results

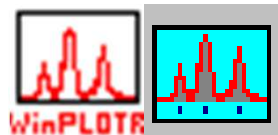


SEQ file: X

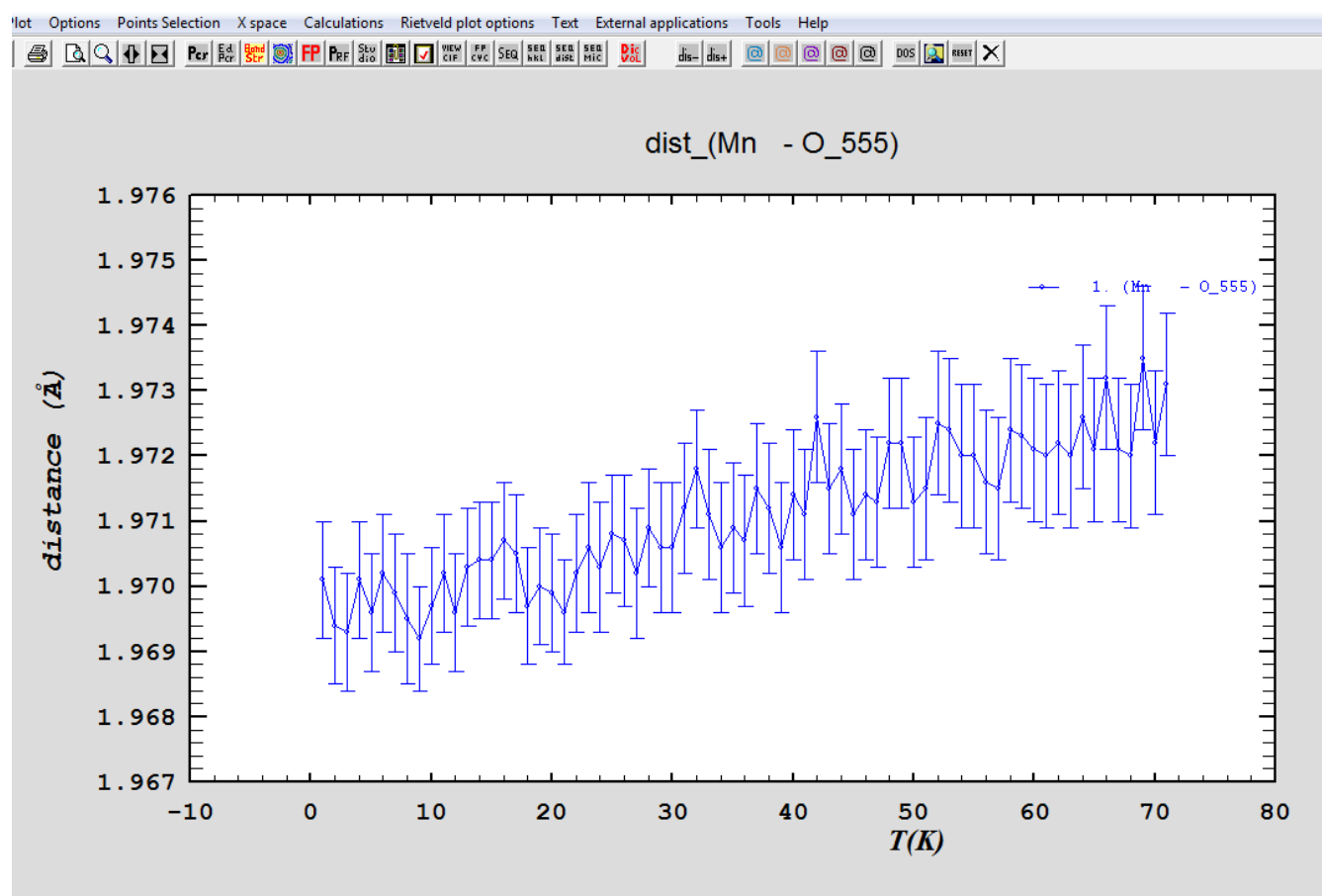
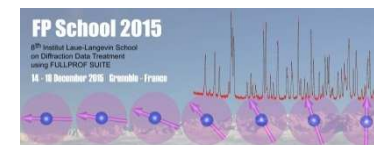
<input type="checkbox"/> 1- GLOBAL_CHI_2	<input type="checkbox"/> 21- Y_La_ph1
<input type="checkbox"/> 2- RB_1 (%)	<input type="checkbox"/> 22- X_O_ph1
<input type="checkbox"/> 3- RF_1 (%)	<input type="checkbox"/> 23- Y_O_ph1
<input type="checkbox"/> 4- Unit_Cell_Volume_1	<input type="checkbox"/> 24- Z_O_ph1
<input type="checkbox"/> 5- Weight_Fraction_1	<input type="checkbox"/> 25- Biso_Mn_ph1
<input type="checkbox"/> 6- RB_2 (%)	<input type="checkbox"/> 26- Biso_La_ph1
<input type="checkbox"/> 7- RF_2 (%)	<input type="checkbox"/> 27- Biso_O_ph1
<input type="checkbox"/> 8- Unit_Cell_Volume_2	<input type="checkbox"/> 28- Biso_O_ph1
<input type="checkbox"/> 9- Weight_Fraction_2	<input type="checkbox"/> 29- Asym1_ph1_pat1
<input type="checkbox"/> 10- Scale_ph1_pat1	<input type="checkbox"/> 30- Asym2_ph1_pat1
<input type="checkbox"/> 11- Zero_pat1	<input type="radio"/> File order
<input checked="" type="checkbox"/> 12- Cell_A_ph1_pat1	<input checked="" type="radio"/> External parameter
<input type="checkbox"/> 13- Cell_B_ph1_pat1	X legend: <input type="text" value="T (K)"/>
<input type="checkbox"/> 14- Cell_C_ph1_pat1	<input type="button" value="OK"/> <input type="button" value="Cancel"/>
<input type="checkbox"/> 15- U-Cagl_ph1_pat1	
<input type="checkbox"/> 16- V-Cagl_ph1_pat1	
<input type="checkbox"/> 17- W-Cagl_ph1_pat1	
<input type="checkbox"/> 18- Ry_Mn1_ph2	
<input type="checkbox"/> 19- X_La_ph1	
<input type="checkbox"/> 20- X_O_ph1	



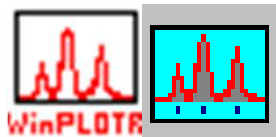
WinPLOTR @ FPSchool, ILL, 14-18 dec. 2015



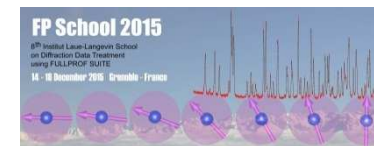
Plotting sequential FP results



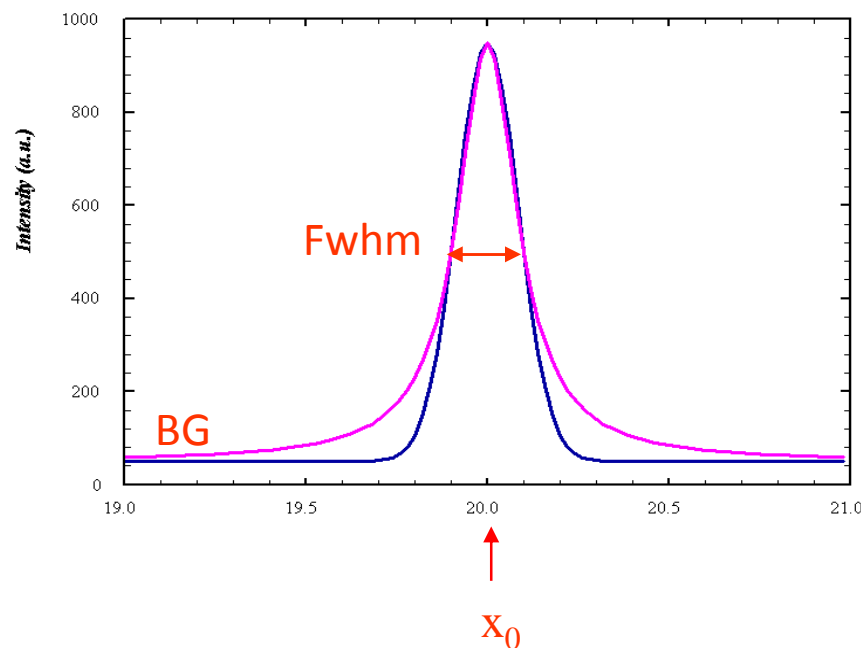
WinPLOTR @ FPSchool, ILL, 14-18 dec. 2015

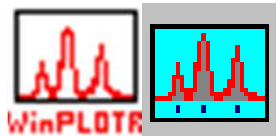


Profile fitting procedure in WinPLOTR

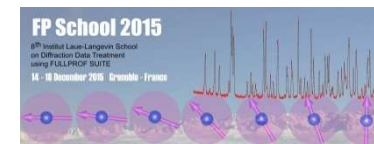


- Profile function : pseudo-Voigt $PV(x) = \eta \cdot L(x) + (1-\eta) \cdot G(x)$
T.C.H. formulation (idem **NPROF=7**) to mimic Voigt function ($L \otimes G$)
- Least-squares refinement of parameters :
 - linear background values
 - reflections parameters : intensity, position, FWHM, η

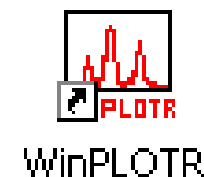




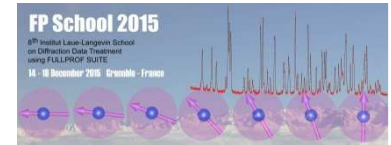
Profile fitting procedure



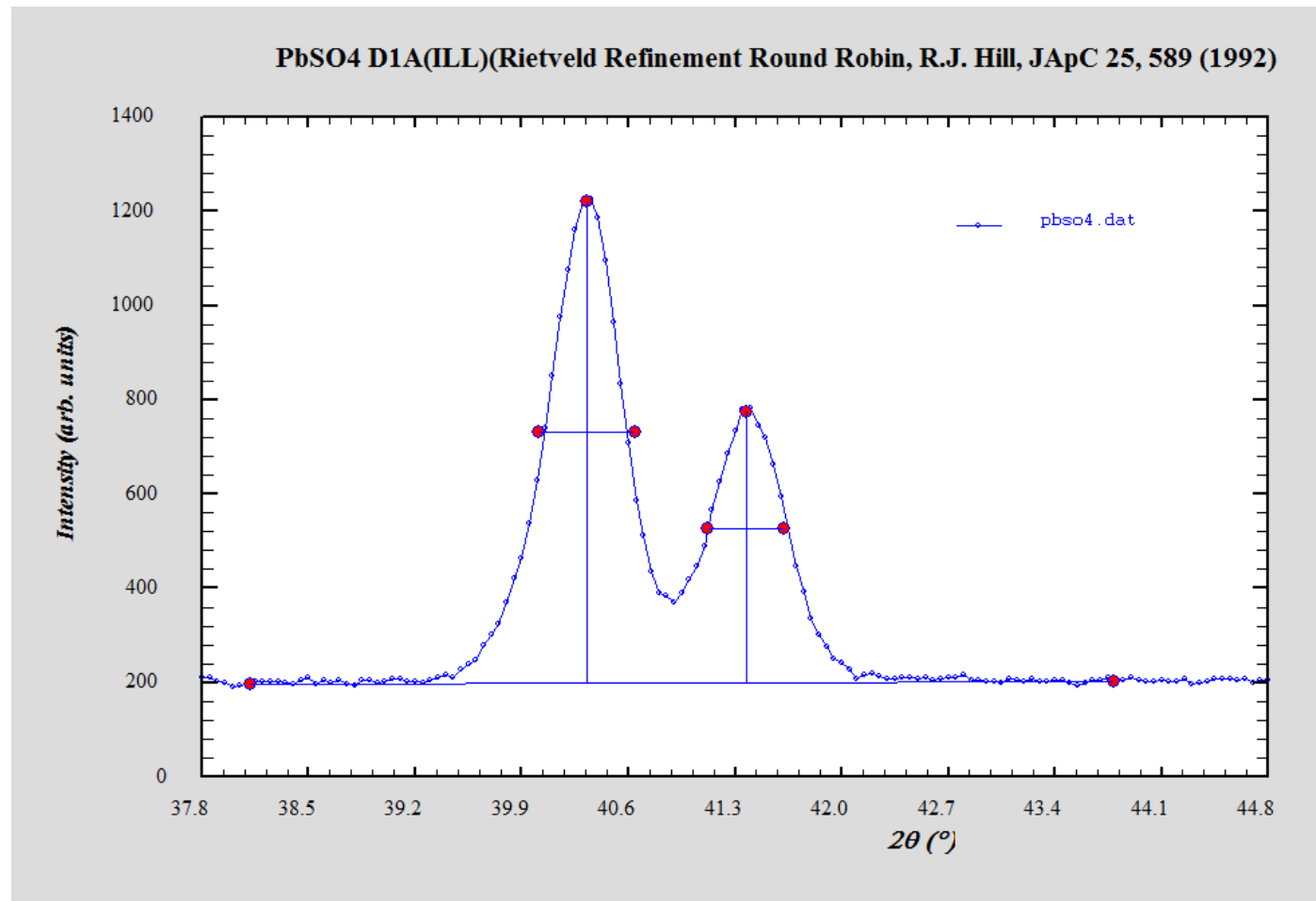
- Starting parameters of L.S. refinements ?
 1. Select parameters with the mouse
 2. Automatic search
 3. From a **.PIK** input file
 4. Single peak / doublet



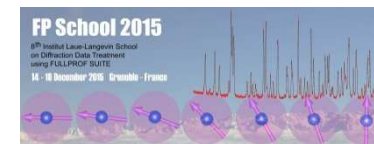
1. Profile fitting procedure : handling mode



- selection of starting parameters values



1. Profile fitting procedure : handling mode



Starting parameters for CWL data profile fitting procedure

Xmin:	<input type="text" value="38.4138832"/>	Xmax:	<input type="text" value="43.19758220"/>	
lambda1 (Å):	<input type="text" value="0"/>	lambda2 (Å):	<input type="text" value="0"/>	I(Ka2)/I(Ka1) ratio: <input type="text" value="0"/> <input type="checkbox"/>
Left background:	<input type="text" value="195.02140800"/> <input checked="" type="checkbox"/>	Right background:	<input type="text" value="197.86633300"/> <input checked="" type="checkbox"/>	
Asymmetry (S_L):	<input type="text" value="0"/> <input type="checkbox"/>	Asymmetry (D_L):	<input type="text" value="0"/> <input type="checkbox"/>	
U:	<input type="text" value="0"/> <input type="checkbox"/>	V:	<input type="text" value="0"/> <input type="checkbox"/>	W: <input type="text" value="0.31175855"/> <input type="checkbox"/>
Z:	<input type="text" value="0"/> <input type="checkbox"/>			
Eta0:	<input type="text" value="0.0001"/> <input checked="" type="checkbox"/>	X:	<input type="text" value="0"/> <input type="checkbox"/>	

Peak		Position		Intensity		Shift FWHM		Shift eta	
1	<input checked="" type="checkbox"/>	<input type="text" value="40.30774310"/> <input checked="" type="checkbox"/>		<input type="text" value="498.15774500"/> <input checked="" type="checkbox"/>		<input type="text" value="0"/> <input type="checkbox"/>		<input type="text" value="0"/> <input type="checkbox"/>	
2	<input checked="" type="checkbox"/>	<input type="text" value="41.39426420"/> <input checked="" type="checkbox"/>		<input type="text" value="363.11123700"/> <input checked="" type="checkbox"/>		<input type="text" value="0"/> <input type="checkbox"/>		<input type="text" value="0"/> <input type="checkbox"/>	

Number of cycles:

☒ no .IRF/.IPC output file ☐ Create .IRF file ☐ Create .IPC file

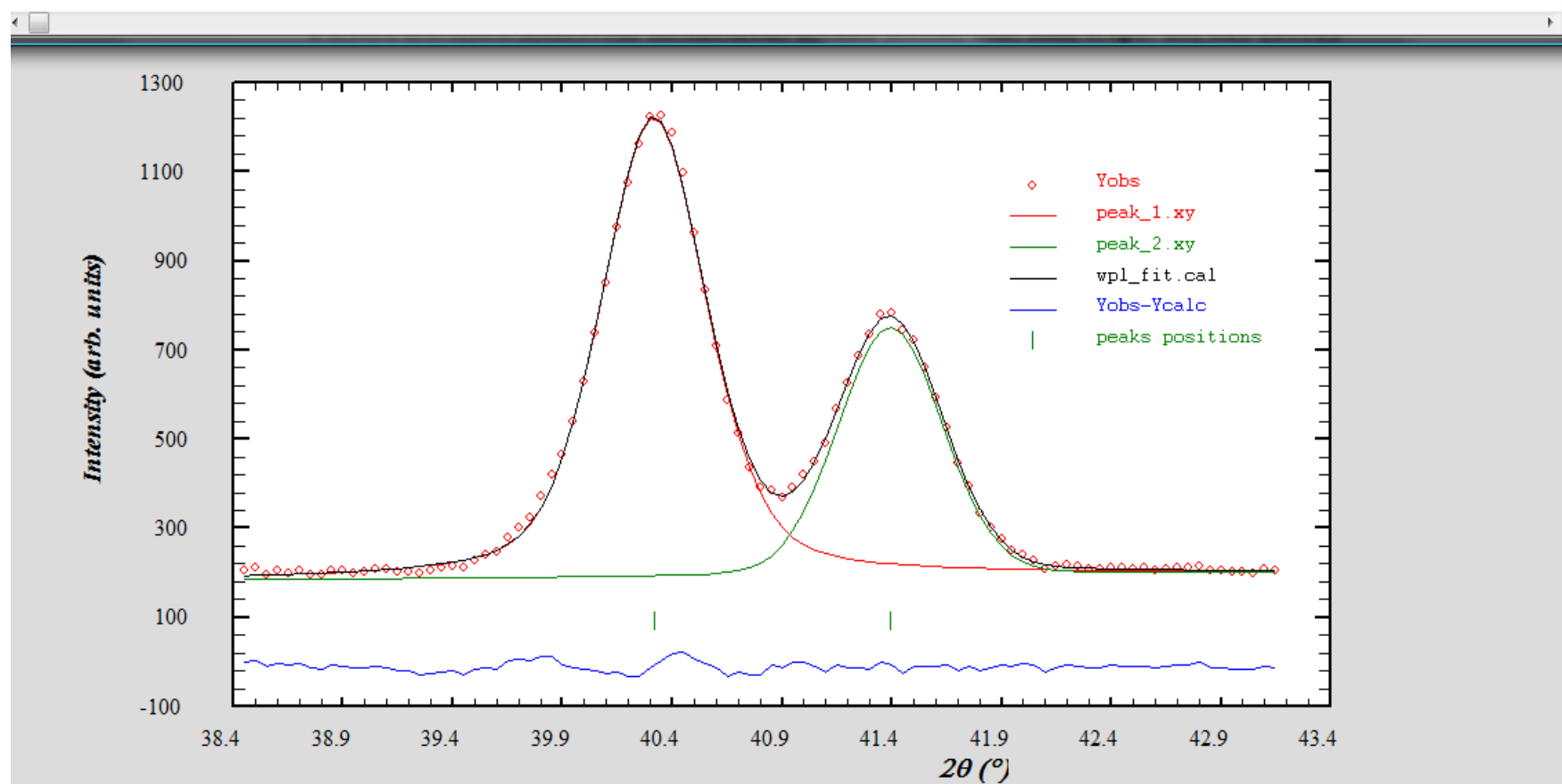
Constant wavelength data profile refinement:

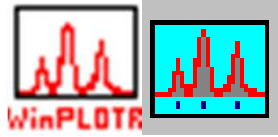
```
=> Parameter X      :      0.000000      0.000000
      Scatt. Variable Background      Sigma
           38.4500      184.1887      2.1392
           43.1500      200.8363      2.1189
```

Position	Sigma	Intensity	Sigma	Shf_FWHM	Sigma	Shf_Eta	Sigma	FWHM	Sigma	eta
40.315678	0.001408	737.35	8.67	0.006986	0.004230	0.488205	0.028404	0.565339	0.004230	0.488305
41.393456	0.002117	342.73	8.13	-0.000860	0.005647	0.114697	0.062580	0.557493	0.005647	0.114797
Rp (%)	Rwp (%)	Rexp (%)	Chi2							
1.8	6.1	4.6516	1.73565							

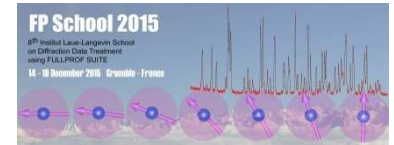
Output files:

```
. pbso4_PF.new: input file with refined parameters
. pbso4_PF.out: summary of the fitting procedure
```

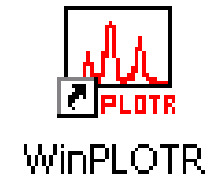




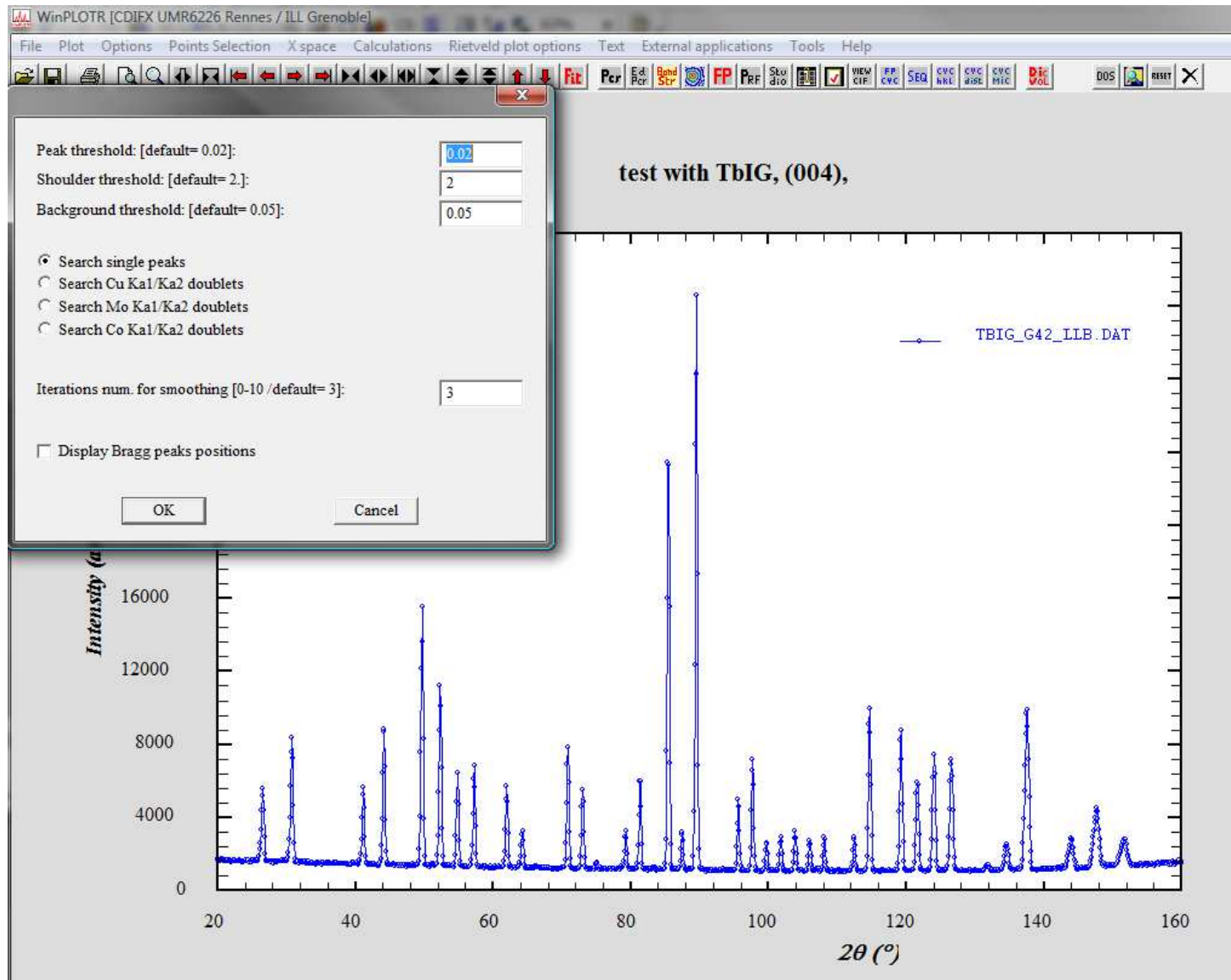
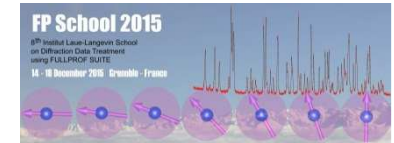
Profile fitting procedure



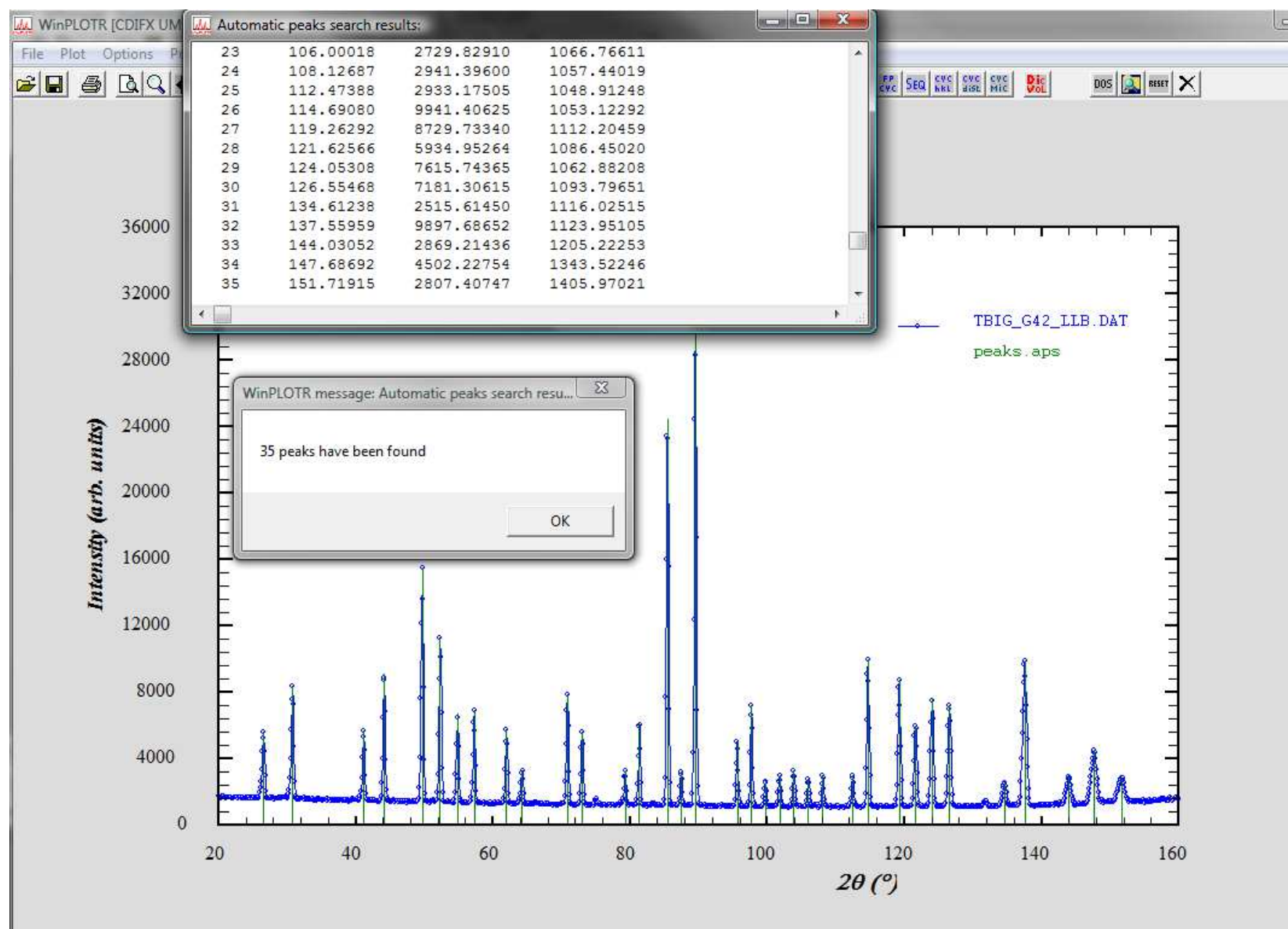
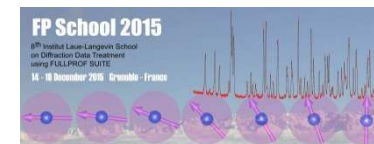
- Starting parameters of L.S. refinements ?
 1. Select parameters with the mouse
 2. Automatic search
 3. From a **.PIK** input file
 4. Single peak / doublet



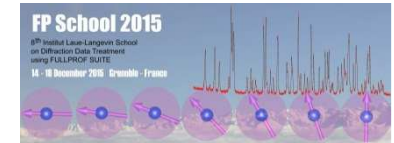
2. Fitting procedure : automatic peak search



2. Fitting procedure : automatic peak search



2. Fitting procedure : automatic peak search



Select your diffractometer:

X-Rays powder diffractometer:

- ☐ conventionnal diffractometer ($U=0.008$; $V=-0.008$; $W=0.009$)
- ☐ INEL CPS120 ($U=0.016$; $V=-0.01$; $W=0.008$)
- ☐ synchrotron radiation ($U=0.0$; $V=0.0$; $W=0.002$)

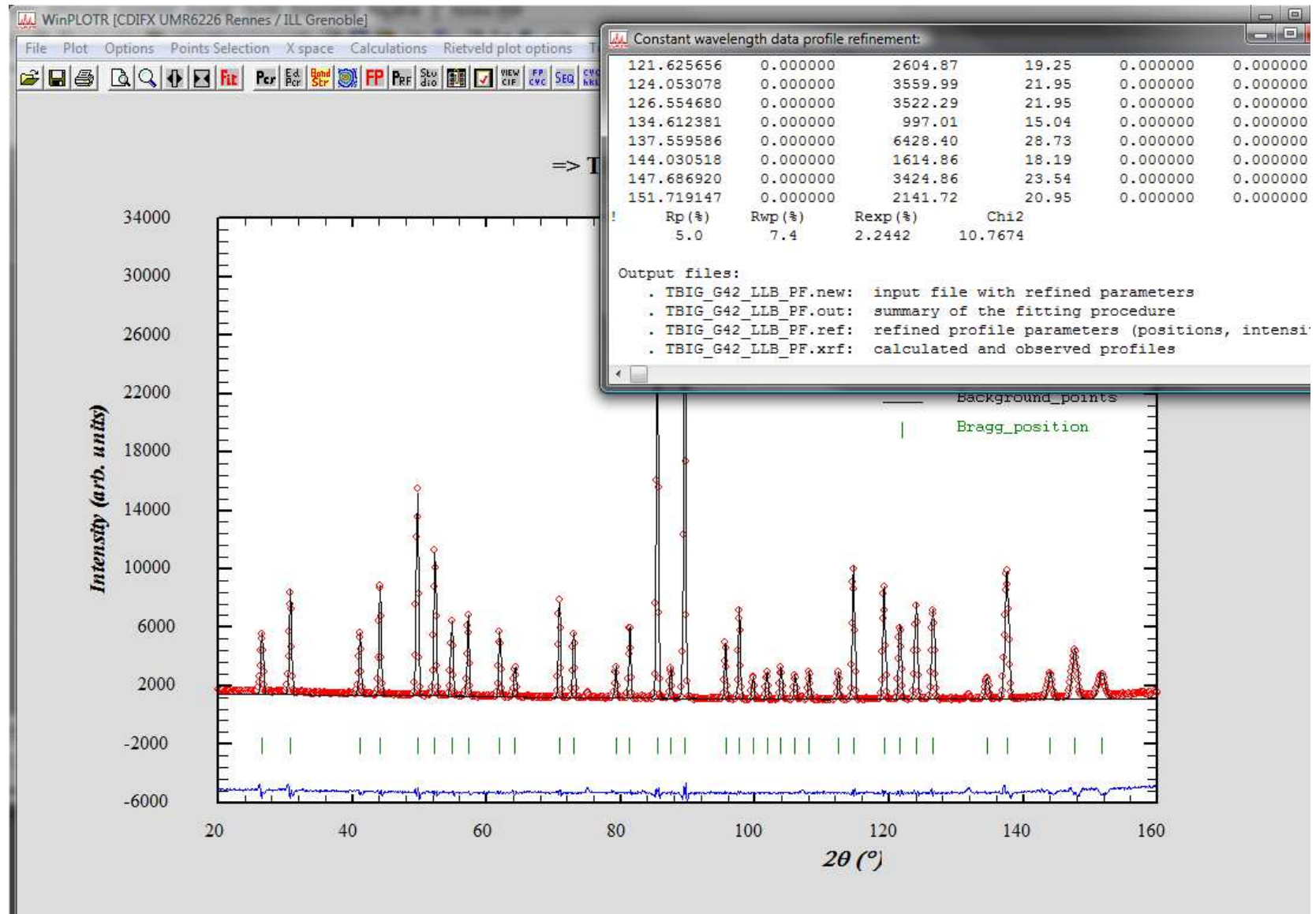
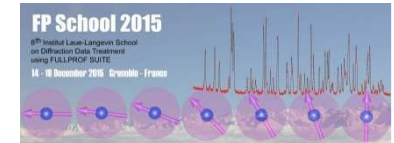
Neutron powder diffractometer:

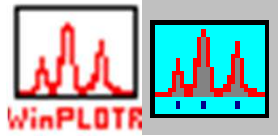
- ☒ G42 (LLB) $\lambda=2.343\text{\AA}$ ($U=0.113$; $V=-0.208$; $W=0.223$)
- ☐ G41 (LLB) $\lambda=2.42\text{\AA}$ ($U=0.81$; $V=-0.28$; $W=0.08$)
- ☐ 3T2 (LLB) $\lambda=1.225\text{\AA}$ ($U=0.306$; $V=-0.468$; $W=0.236$)
- ☐ D2B (ILL) $\lambda=1.59\text{\AA}$ ($U=0.086$; $V=-0.229$; $W=0.254$)
- ☐ other diffractometer (enter your own U,V,W values)

OK

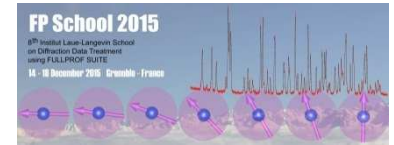
Cancel

2. Fitting procedure : automatic peak search

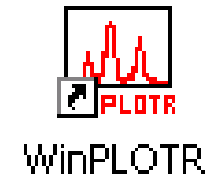




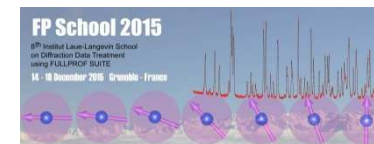
Profile fitting procedure



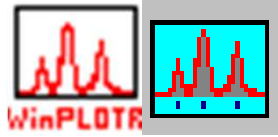
- Starting parameters of L.S. refinements ?
 1. Select parameters with the mouse
 2. Automatic search
 3. From a **.PIK** input file
 4. Single peak / doublet



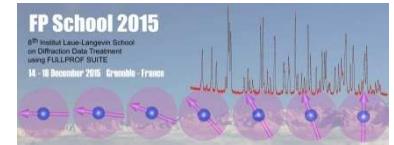
3. Fitting procedure : from .PIK input file



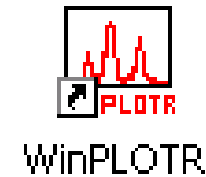
```
KEDIT - [D:\WinPLOTR\examples\pbso4_n.new]
File Edit Actions Options Window Help
continue
====>
*** Top of File ***
=> TITLE: pbso4_n.dat
! Ang_init   Ang_fin   Nbac   Npeak   Ncyc   Inte   Inst   Jobt   Cont   Weight   Corr   Constr
  44.0000    62.0000     2     11     10     0     6     2     0     0     0.     0
  1.540560    1.544330  <= Lambda1 & Lambda2
! Global Profile Parameters:
  0.0000     0     <= Kalph2/Kalph1 ratio & Flag
  0.0000     0     <= Asymmetry parameter1 & "
  0.0000     0     <= Asymmetry parameter2 & "
  0.0860     1     <= Parameter U & "
 -0.2290     1     <= Parameter V & "
  0.2540     1     <= Parameter W & "
  0.0000     0     <= Parameter Z & "
  0.1000     0     <= Parameter Eta0 & "
  0.0000     0     <= Parameter X & "
! Background Parameters:
! 2Theta/TOF/Energy   Background   Flag
  44.0000    200.0000     1
  62.0000    215.3333     1]
! Reflection Parameters:
! 2Theta/TOF/Energy   Intensity   Shift-FWHM   Shift-Eta   Flags
  46.5785    450.5725    0.0000    0.0000    1 1 1 0
  47.7482    129.3865    0.0000    0.0000    1 1 1 0
  49.5211    342.3236    0.0000    0.0000    1 1 1 0
  50.4567    248.8786    0.0000    0.0000    1 1 1 0
  51.5505    548.4024    0.0000    0.0000    1 1 1 0
  52.2982    848.1331    0.0000    0.0000    1 1 1 0
  53.1669    320.1445    0.0000    0.0000    1 1 1 0
  54.9535    1134.3667    0.0000    0.0000    1 1 1 0
  56.0510    929.1931    0.0000    0.0000    1 1 1 0
  57.8286    370.5429    0.0000    0.0000    1 1 1 0
  60.1728    73.4670    0.0000    0.0000    1 1 1 0
! Chi2 = 149.170
*** End of File ***
```



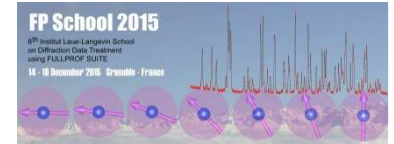
Profile fitting procedure



- Starting parameters of L.S. refinements ?
 1. Select parameters with the mouse
 2. Automatic search
 3. From a **.PIK** input file
 4. Single peak / doublet



4. Profile fitting procedure : single peak



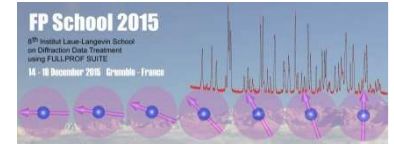
- . Starting parameters are detected automatically :
 - . background points
 - . single peak / doublet parameters : position, FWHM

Example: $\text{NdSrNi}_{0.8}\text{Cu}_{0.2}\text{O}_4$ / RX-D500



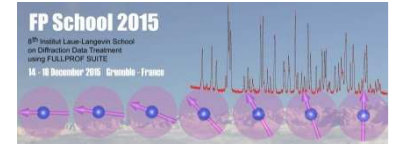
WinPLOTR

Background points selection



- . Automatic mode (**Points selection / Automatic background**):
 - . Created from the pattern file
 - . Number of points $\sim \text{SQRT}(n)$, with a quasi uniform repartition on 2θ
 - . Modify background (**Points selection / Add /remove background points**)
 - . Save background points to paste into .PCR file for **FullProf**
- >> Raw background for raw refinements (not recommended for advanced analysis)

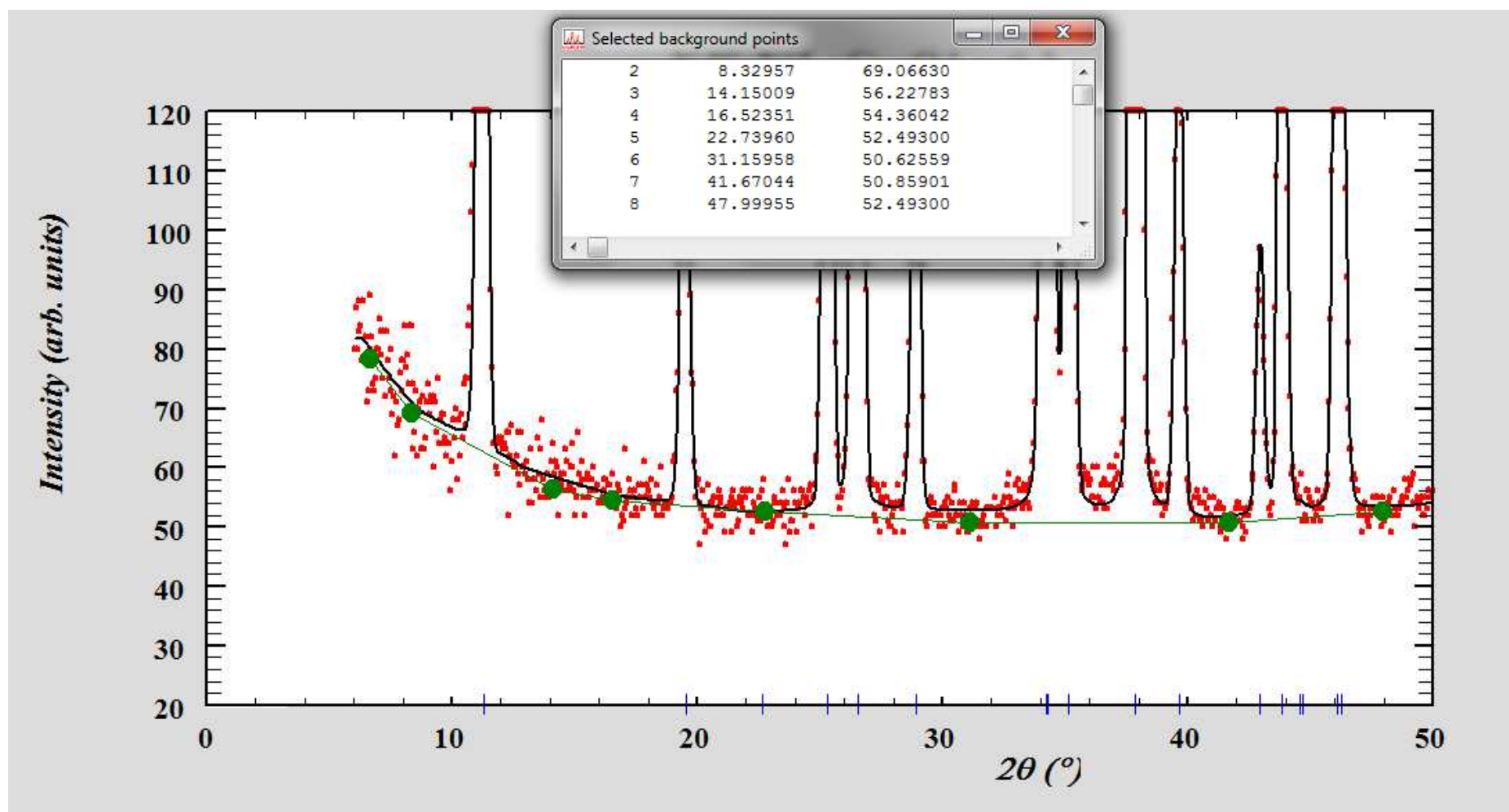
Background points selection

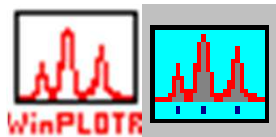


- . Manual mode (**Points selection / Select background points**):
 - . Created from the data file or preferentially from the **.PRF** file
 - . Number of points = depends on peaks overlap, background modulation, ...
 - . Modify backgrounds (**Points selection / Add / background points**)
 - . Save background points to include into **.PCR** file for **FullProf**

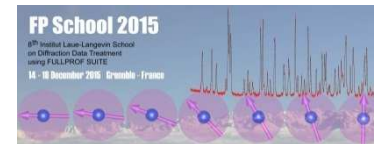
>> Good starting points for **FullProf** refinements (Rietveld / profile matching (strongly recommended !))

Background points selection





WinPLOTR tools (**T**ools menu option)



- **Space group informations :**
 - . features
 - . symmetry operators
 - . Wyckoff positions
 - . extinctions

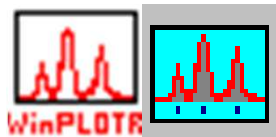
- **(hkl) list generation in a given 2θ range, from :**
 - . space group
 - . unit cell parameters
 - . wavelength

- **unit cell volume calculation**

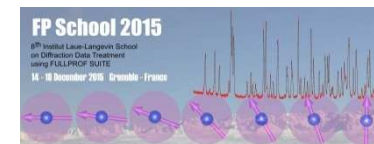
- **molecular weight calculation**



WinPLOTR



Advanced use of WinPLOTR

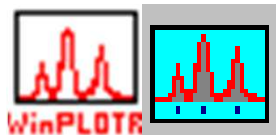


for repeatable actions :

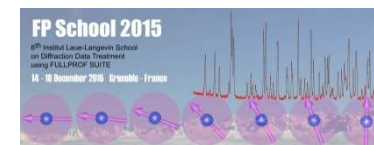
- . Files conversion
- . Multi-scans **UXD** or **XRDML** file normalization
- . Change X space
- . Automatic single peaks/doublet profile fitting
- . Save graphics as bitmap or PostScript files

Use a command file, containing keywords and arguments:

- . **FILE** file%name file%format
- . **SAVE_AS_XY**, **SAVE_AS_INSTRM_0**
- . **UXD_NORMA/XRDML_NORMA**
- . **SHIFT_X/SHIFT_Y** shift_x/shift_y
- . **OFFSET_X/OFFSET_Y** offset_x/offset_y
- . **MULTIPLY_X/MULTIPLY_Y** mult_x/mult_y

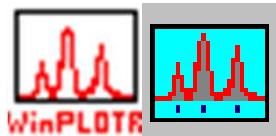


Advanced use of WinPLOTTR

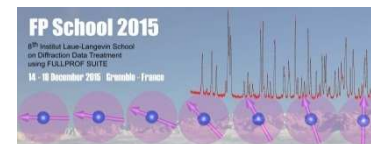


Use a command file, containing keywords and arguments:

```
. SUBTRACT file_1 file_2 format
. WAVE
. TRANSF_X1_TO_X2 (2THETA / Q / STL / D / S)
. FIT_SINGLE_PEAK xmin xmax
. FIT_DOUBLET_CU/MO/CO xmin xmax
. BITMAP / EPS
```



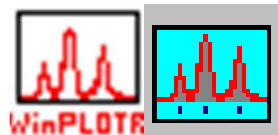
Launching **WinPLOT** from the command line, using a command input file



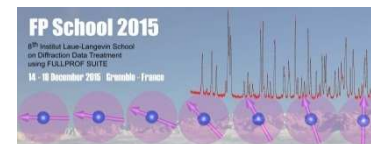
```
d:\data>winplotr command_file.cmd
```

Example 1 : conversion of files

```
FILE data_file_001.uxd 9  
SAVE_AS_XY  
FILE data_file_002.uxd 9  
SAVE_AS_XY  
...  
FILE data_file_455.uxd 9  
SAVE_AS_XY
```

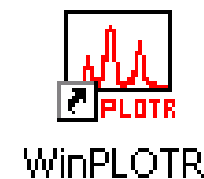


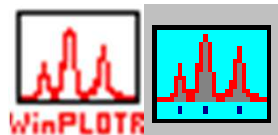
Launching **WinPLOTR** from the command line, using a command input file



Example 2 : profile fitting

```
FILE ndsrni_rx.dat 1
FIT_SINGLE_PEAK 24. 25.
FILE ndsrni_rx_pf.xrf 100
BITMAP
PLOT
```





WinPLOTR : what's new ?

