

# **First FullProf School 2025:**

## **Diffraction data analysis of energy materials**

# **Introduction to the FullProf Suite**

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# Outline

**1. History of the FullProf Suite Development.**

**2. Programs of the FullProf Suite.**

**3. Live Presentation of the FullProf Suite**

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**1. History of the FullProf Suite Development.**

2. Programs of the FullProf Suite.

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# ANCESTORS OF FULLPROF: FROM 1967 TO 1986

## Hugo Rietveld family of programs

**PREF & PROFLS**  
Hugo Rietveld - 1967, 1969 Petten

Language: Algol, Fortran

**PREF & PROFLS**  
Alan Hewat - 1973 Harwell

Uij added, Fortran

**PREF & PROFLS**  
Von Dreele - 1973 Oxford

Bigger structures & X-rays (didn't work!)

**DBW2.9 - 1981-1994**  
Young & Wiles - 1973 Oxford

X-ray profiles, Young & Wiles  
rewrite a single program,  
Sakthivel (DBWS)

**TOFPREF & TOFLS**  
Von Dreele - 1982 RAL

Neutron TOF, IPNS  
adopted by Rotella for VAX

**LPHM- Hunter % Howard -Australia**  
**Phillips – Instrument package**  
Might be other descendants from DBW/DBWS

**GSAS**  
Larson & Von Dreele - 1982 -1986

**FullProf – J. Rodriguez-Carvajal –France**

THE EUROPEAN NEUTRON SOURCE

# THE EARLY DEVELOPMENT OF FULLPROF: 1988 - 1992

Reorganization of the Fortran 77 code of DBW (include files, more modular subroutines)

**Anisotropic broadening** due to strains and size effects

Introduction of **magnetic structures** and the formalism of propagation vectors.

Refinement without structural model as replacement of the Pawley program (**Le Bail fit**).

# HISTORY OF FULLPROF: FROM 1988 TO 1992 @ILL

## Still @ILL ... but moving to the LLB

The first public version of FullProf was presented in a Satellite of XV IUCr congress in Bordeaux on powder diffraction (1990), in which Le Bail fits and conventional treatment of commensurate magnetic structures were implemented.

J. Rodriguez-Carvajal, "FullProf: A Program for Rietveld Refinement and Pattern Matching Analysis," Abstract of the Satellite Meeting on Powder Diffraction of the XV Congress of the IUCr, Toulouse, France, 1990, p. 127

Later I implemented in the program **FullProf** the **formalism of propagation vectors** that was able to treat all kind of magnetic structures including incommensurate structures. This was presented in the **WORKSHOP ON THE USE OF NEUTRONS AND X-RAYS IN THE STUDY OF MAGNETISM** (Grenoble, **January 21-23, 1993** )

Published in **Physica B 192, 55-69 (1993): Recent advances in magnetic structure determination by neutron powder diffraction**

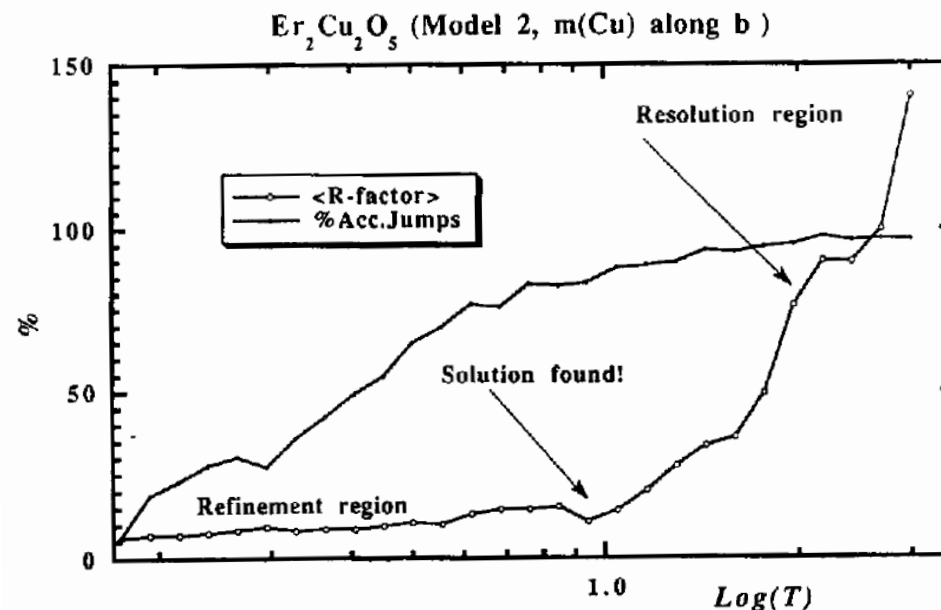
# THE REFERENCE PAPER OF FULLPROF

J. Rodriguez-Carvajal, Physica B 192, 55 (1993)

First description of some features of the program **FullProf**

Description of the program **MagSan** for determining commensurate magnetic structures using Simulated Annealing (later included in **FullProf** for general structures)

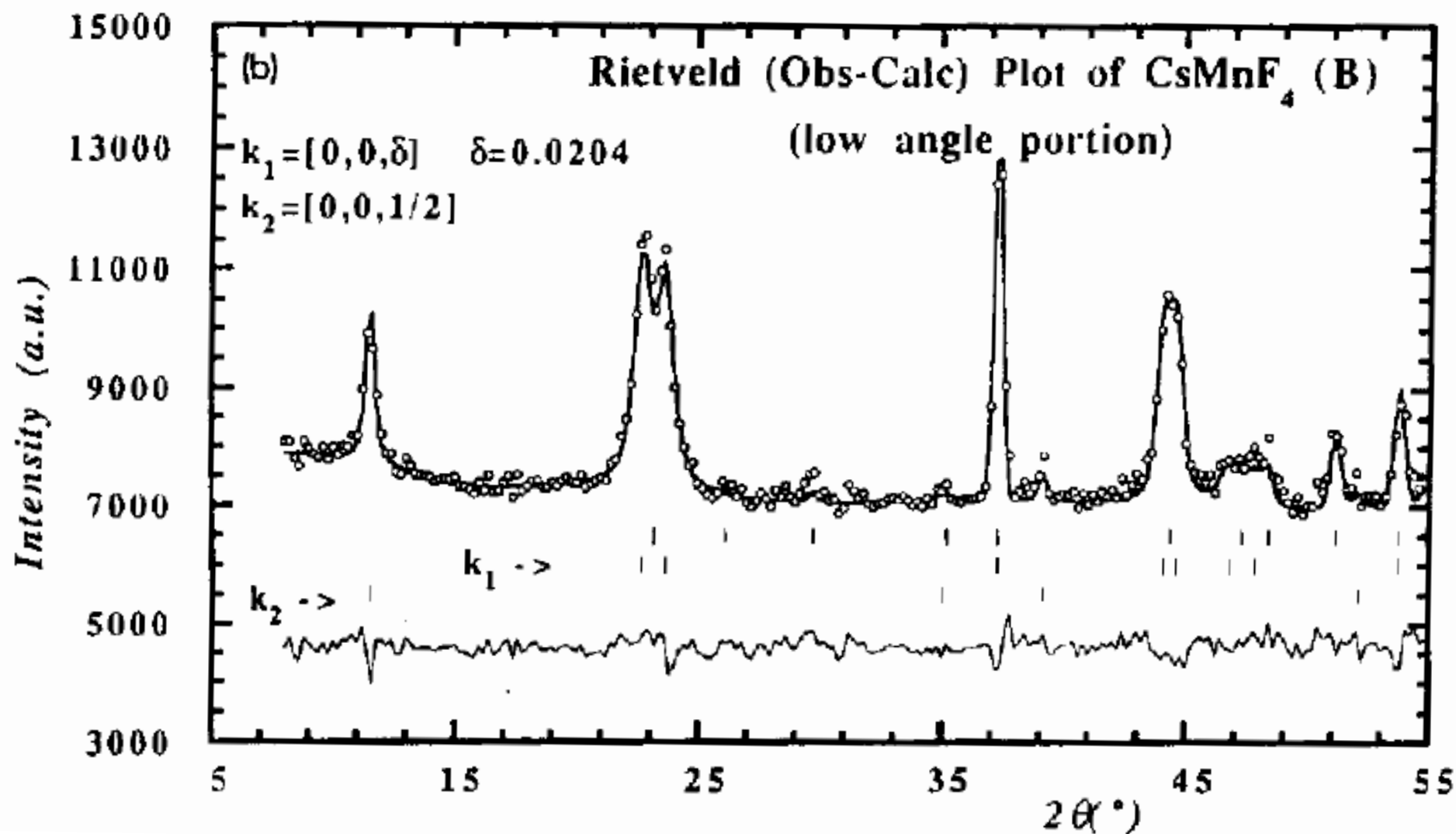
```
begin
  Initialize (set to zero useful quantities, do preliminary
             calculations)
   $\tau = 1$ 
  do
    do
      Perturb the system:
       $\Omega_{old} \rightarrow \Omega_{new}, \Delta = E(\Omega_{new}) - E(\Omega_{old})$ 
      if  $\Delta \leq 0$  then accept, else
        if  $\exp(-\Delta/T_\tau) > \text{random}[0,1]$  then accept
        if accept then Update (replace  $\Omega_{old}$  by  $\Omega_{new}$ )
      until equilibrium is approached closely enough (Ncyc)
       $T_{\tau+1} = f(T_\tau)$  (decrease temperature, usually  $T_{\tau+1} = qT_\tau$ ,
                         $q \approx 0.9$ )
       $\tau = \tau + 1$ 
    until stop criterion is true (maximum  $\tau$ , convergence, low
      % accepted. . .)
end
```



SOURCE

# THE REFERENCE PAPER OF FULLPROF

J. Rodriguez-Carvajal, Physica B 192, 55 (1993)





# THE EARLY DEVELOPMENT OF FULLPROF

## THE MIDDLE OF THE NINETIES

Rigid bodies, special form factors ... (1992 ... 1995)

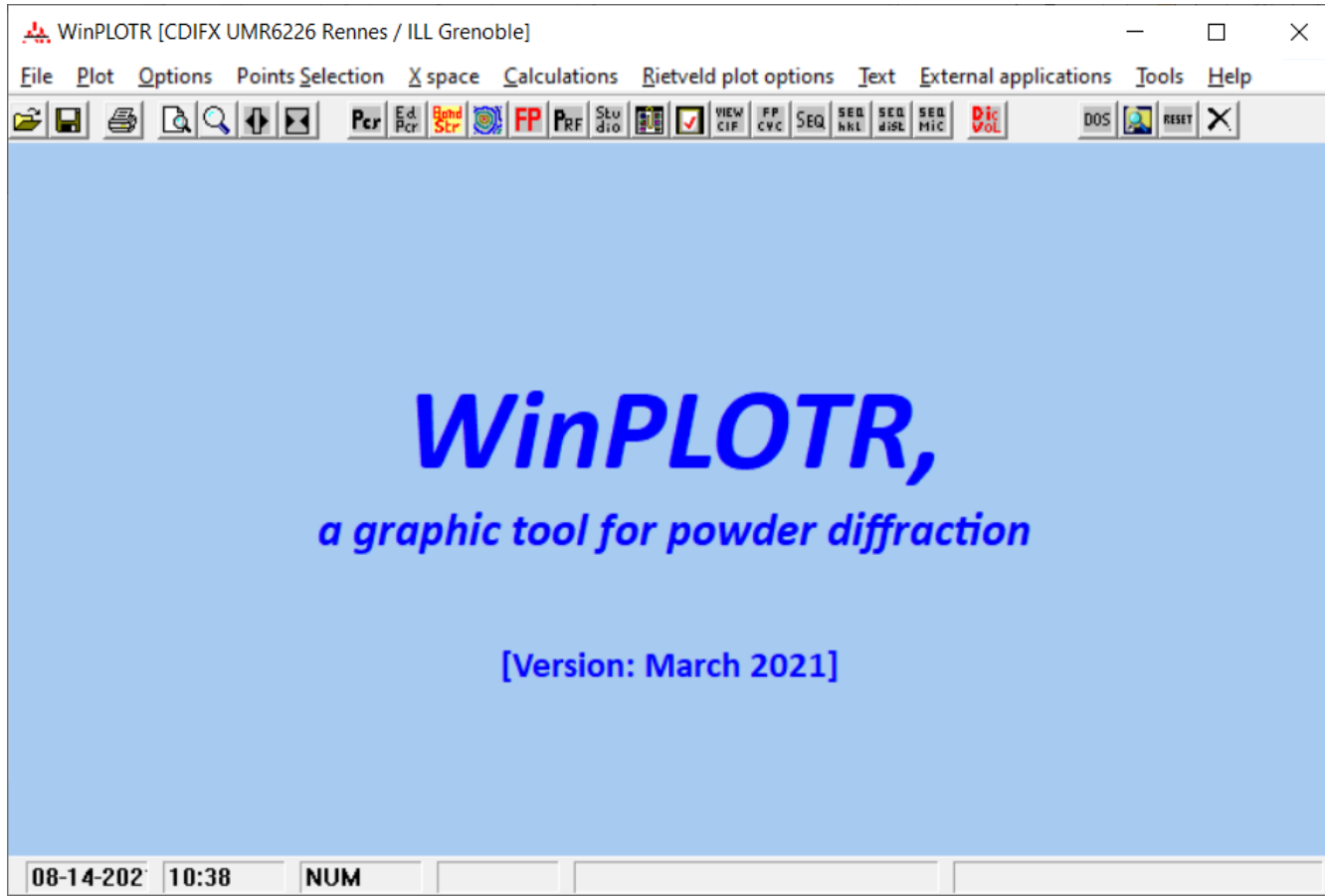
Rewriting the code (C++ or Fortran 90?), 1995-1996 ... **Finally a subset (ELF) of Fortran 90** for FullProf.

Suppression of old code, commons, creation of modules, starting the development of CrysFML (subset of Fortran 90: F), time of flight neutron diffraction, multiple patterns, single crystal refinements.

# THE EARLY DEVELOPMENT OF FULLPROF

## THE MIDDLE OF THE NINETIES

Development of WinPLOTR with Thierry Roisnel at LLB (Fortran 90 + RealWin): *a high impact in the distribution of FullProf*

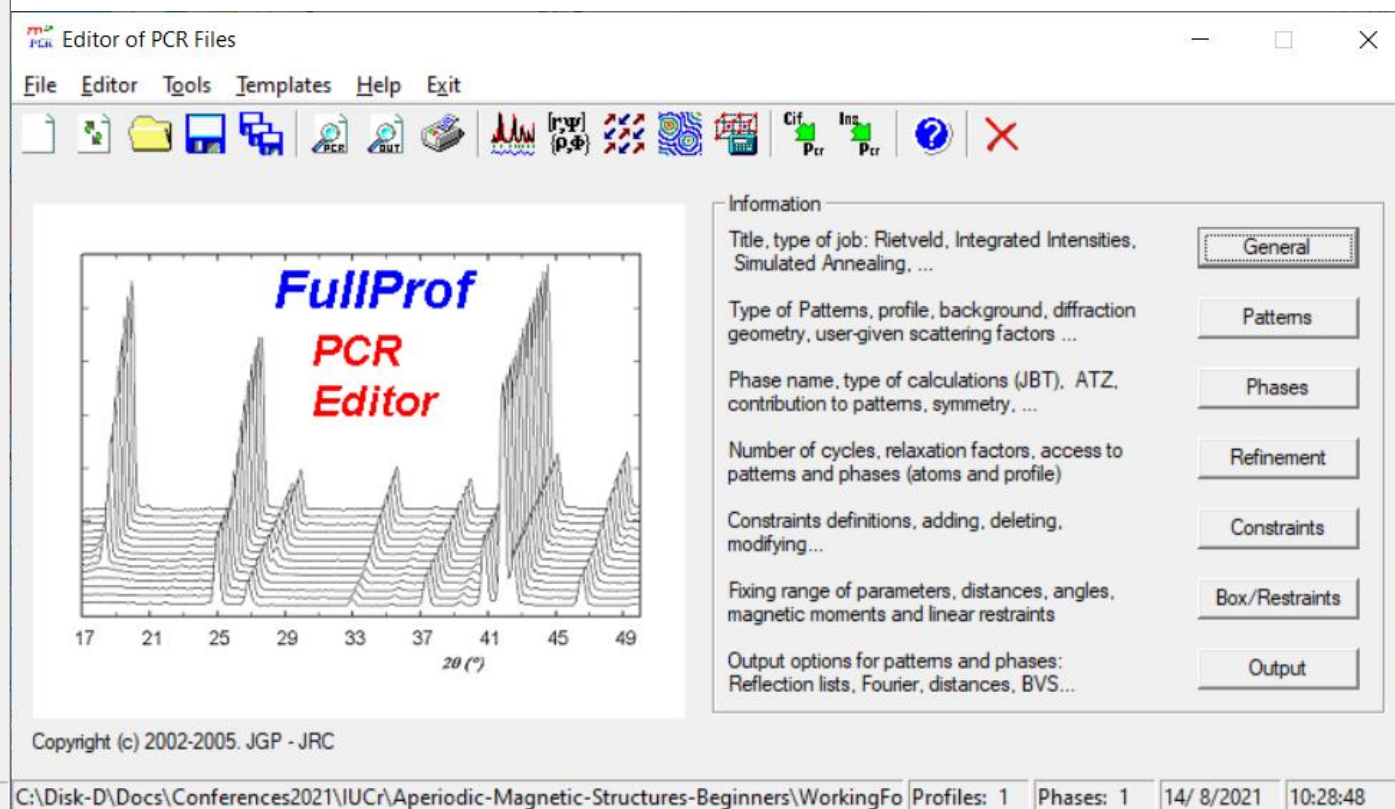
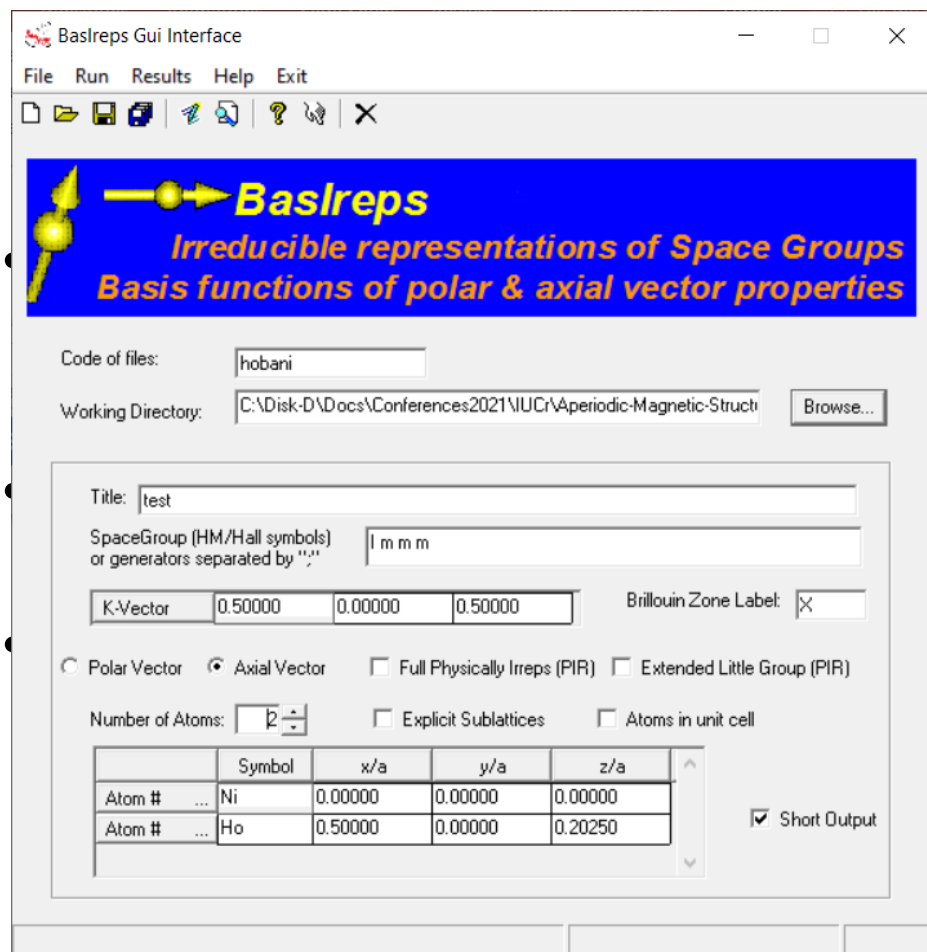


- Visualization of powder patterns
- Peak search, background generation
- Fitting capabilities
- Run FullProf from WinPLOTR
- Interoperability with other programs: BondStr, DICVOL, TREOR, etc.

A recent version of WinPLOTR that works only in Windows and it is a 32 bit application. It will not be working in the future.

# THE EARLY DEVELOPMENT OF FULLPROF

## THE END OF THE NINETIES TO THE NEW CENTURY



Recent developments of the program FullProf

Juan Rodríguez-Carvajal

*Commission for Powder Diffraction, IUCr, Newsletter 26*, 12-19 (2001).

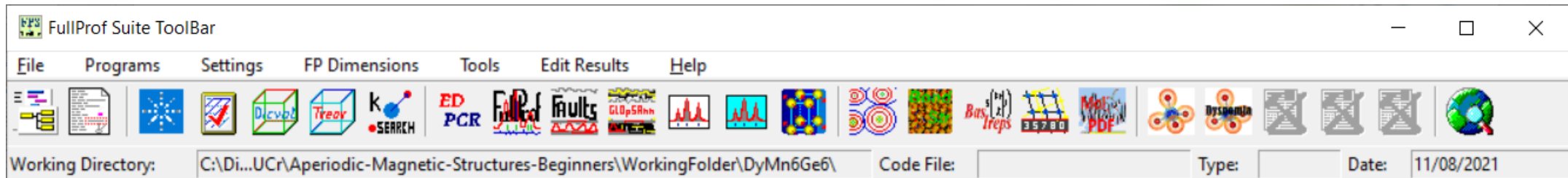
WinPLOTR: a Windows tool for powder diffraction patterns analysis

T. Roisnel and J. Rodríguez-Carvajal.

*Materials Science Forum* **378-381**, 118-123 (2001).

# THE EARLY DEVELOPMENT OF FULLPROF

## THE NEW CENTURY: CRYSFML AND FULLPROF 2K



- Development of **FullProf Studio** (L.C. Chapon)
- Development of **GBondStr/BondStr** (J. González-Platas)
- Development of the new **WinPLOTR-2006** (J. González-Platas, O. Baltuano)
- During the first decade of the new century the **FullProf Suite** was continuing to be developed and largely distributed through the Internet

**Crystallographic Fortran Modules Library (CrysFML): A simple toolbox for crystallographic computing programs**

Juan Rodríguez-Carvajal and Javier González-Platas

Computing Commission of the International Union of Crystallography.

*Compcomm Newsletter* **1**, 50-58 (2003).

**The Once and Everliving FORTRAN : Why Fortran still goes onward and upward while many of its "replacement" languages have already died**

Juan Rodríguez-Carvajal

Computing Commission of the International Union of Crystallography.

*Compcomm Newsletter* **3**, 32-40 (2004).

# Outline



1. History of the FullProf Suite Development.

**2. Programs of the FullProf Suite.**

3. Live Presentation of the FullProf Suite

# PRESENTATION OF THE FULLPROF SUITE

**FullProf** (may be run in console mode using scripts for thousands of patterns)

- A program for :
  - Simulation of powder diffraction patterns
  - Pattern decomposition  $\Rightarrow$  integrated intensities
  - Structure refinement
  - Powder and single crystal data
- Crystal and magnetic structures: special form factors, rigid bodies, symmetry adapted modes, incommensurate magnetic structures, restraints, microstructure effects, superspace for magnetic structures, etc.
- Multiple data sets: simultaneous treatment of several powder diffraction patterns (CW X-rays & neutrons, Energy dispersive X-rays, TOF neutron diffraction)
- Combined treatment of single crystal and powder data
- Crystal and magnetic Structure determination capabilities:
  - simulated annealing on integrated intensity and profile data
- Polarized neutrons: Flipping ratio analysis
- Single crystal electron diffraction (kinematic only)



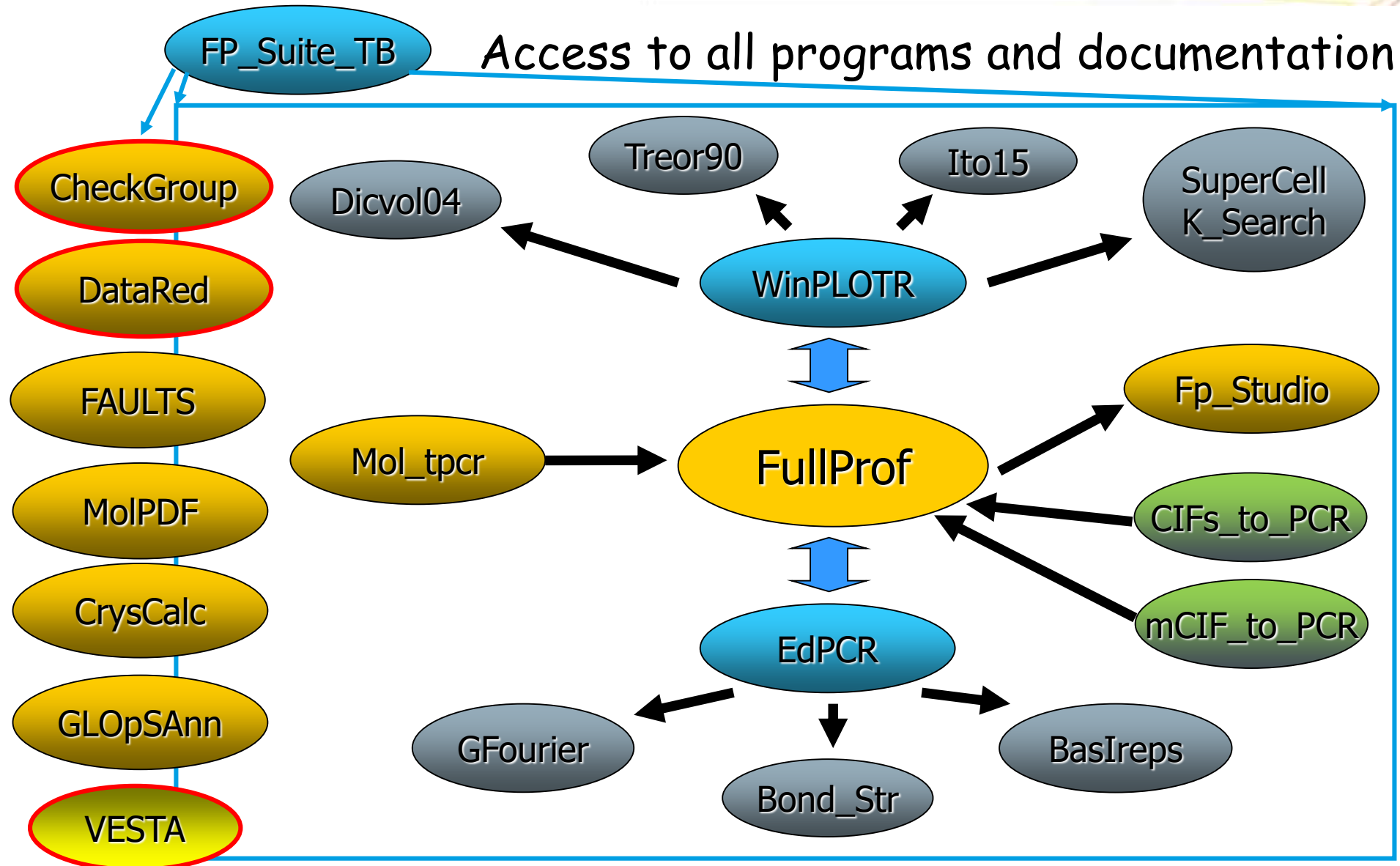
## The PCR file

It is a text file containing all instructions for running **FullProf**. A good text editor is needed for using the most advanced options

The PCR file may be produced automatically for combining different patterns and phases using the utilities:  
**CIFs\_to\_PCR** or **mCIF\_to\_PCR**

It may be produced from the scratch by using **EdPCR**

# PRESENTATION OF THE FULLPROF SUITE



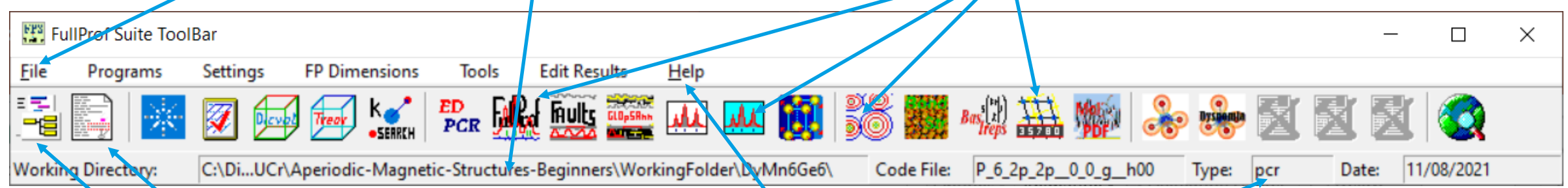
Other utilities:  
**MHall**  
**TOF\_fit\_LM**  
...



# PRESENTATION OF THE FULLPROF SUITE

File: Select working directory

Run Programs



The screenshot shows the FullProf Suite toolbar with various icons and a status bar. Annotations with blue arrows point to specific features:

- File: Select working directory** points to the 'File' menu.
- Run Programs** points to the 'Programs' menu.
- Edit current file** points to the 'Edit Results' menu.
- Load PCR or recognised files** points to the 'Tools' menu.
- Access to documentation** points to the 'Help' menu.

Working Directory: C:\Di...UCr\Aperiodic-Magnetic-Structures-Beginners\WorkingFolder\DiMn6Ge6\

Code File: P\_6\_2p\_2p\_0\_0\_g\_h00

Type: pcr

Date: 11/08/2021

# PRESENTATION OF THE FULLPROF SUITE

Settings for FullProf Suite Toolbar

FullProf Suite Programs: control the automatic run of a program after running FullProf

☐ Run automatically Fourier ☐ Run automatically WinPLOT-2006 ☐ Run automatically WinPLOT

☐ Run automatically FP\_Studio ☐ Run automatically VESTA ☐ Run FullProf Console (fp2k) instead of wfp2k

OK Cancel Save settings

General Programs

Editor: C:\Program Files\Notepad++\notepad++.exe Browse

Browser: C:\Program Files\Mozilla Firefox\firefox.exe Browse

PDF Viewer: Browse

PS Viewer: Browse

External Programs

VESTA C:\Program Files\VESTA\VESTA.exe Browse

Dysnomia C:\Program Files\VESTA\Dysnomia64.exe Browse

Program 1: Label Program: Browse

Program 2: Label Program: Browse

Program 3: Label Program: Browse

☐ Toolbar always on TOP Prefix for console programs cmd /t:0e /c

Select a text Editor  
Select a Browser

Locate VESTA exe

Locate External programs

# PRESENTATION OF THE FULLPROF SUITE

## WinPLOTR

- 0: Visualizing multiple powder patterns (superimposed or 3D)
- 1: Automatic peak search for indexing
- 2: Saving peaks as DICVOL04, Treor90, etc
- 3: Running indexing programs
- 4: Automatic generation of PCR file for cell refinement and integrated intensity extraction (Le Bail fit)
- 5: Making individual peak fits
- 6: Exporting background files
- 7: Invoking other programs

## EdPCR

GUI for editing PCR files  
Only limited number of options  
Shubnikov and superspace groups are not currently available within EdPCR

# PRESENTATION OF THE FULLPROF SUITE: CIFs\_to\_PCR

Select input files with information of the different phases: e.g. a list of CIF files

Select data files containing diffraction patterns and the corresponding features

Select the IRF file of each pattern

Program CIFs\_to\_PCR

Dialog for running the program CIFs\_to\_PCR

Buffer File (or single CIF, PCR, CFL, RES file)

Browse

Edit Buffer File ☐ Include Path ☒ No Path Edit Final PCR File

Run CIFs\_to\_PCR

Cancel / Exit

Save File C2PCR

Pattern file #1: Browse ☒ X-rays ☐ Neutrons CW ☐ Neutrons TOF Instm. 0

IRF file #1: Browse ☒ Profile ☐ Int. Intensities 0 IRF-type

Pattern file #2: Browse ☒ X-rays ☐ Neutrons CW ☐ Neutrons TOF Instm. 0

IRF file #2: Browse ☒ Profile ☐ Int. Intensities 0 IRF-type

Pattern file #3: Browse ☒ X-rays ☐ Neutrons CW ☐ Neutrons TOF Instm. 0

IRF file #3: Browse ☒ Profile ☐ Int. Intensities 0 IRF-type

Pattern file #4: Browse ☒ X-rays ☐ Neutrons CW ☐ Neutrons TOF Instm. 0

IRF file #4: Browse ☒ Profile ☐ Int. Intensities 0 IRF-type

Pattern file #5: Browse ☒ X-rays ☐ Neutrons CW ☐ Neutrons TOF Instm. 0

IRF file #5: Browse ☒ Profile ☐ Int. Intensities 0 IRF-type

# PRESENTATION OF THE FULLPROF SUITE

## **FAULTS**

Program to refine powder diffraction patterns (X-rays and neutrons) of crystal systems with any type of coherent planar defect. Based on DIFFaX and CrysFML

## **MoIPDF**

Program to refine Pair Distribution Function specialized for Molecular systems

## **GLOpSAnn**

Global optimization by simulated annealing of crystal structures using a variety of cost functions: R-factors, BVS, distances, torsion angles, etc.

# PRESENTATION OF THE **FULLPROF SUITE**

The programs of the **FullProf Suite** are not black-box-like with a single way of doing the things.

The GUIs are mere help utilities for handling the input control files.

The learning curve for the use of the programs may be steeper than for other programs, however understanding the content of the input files and making manual changes is much faster, once the user is becoming expert, than using the GUIs.

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