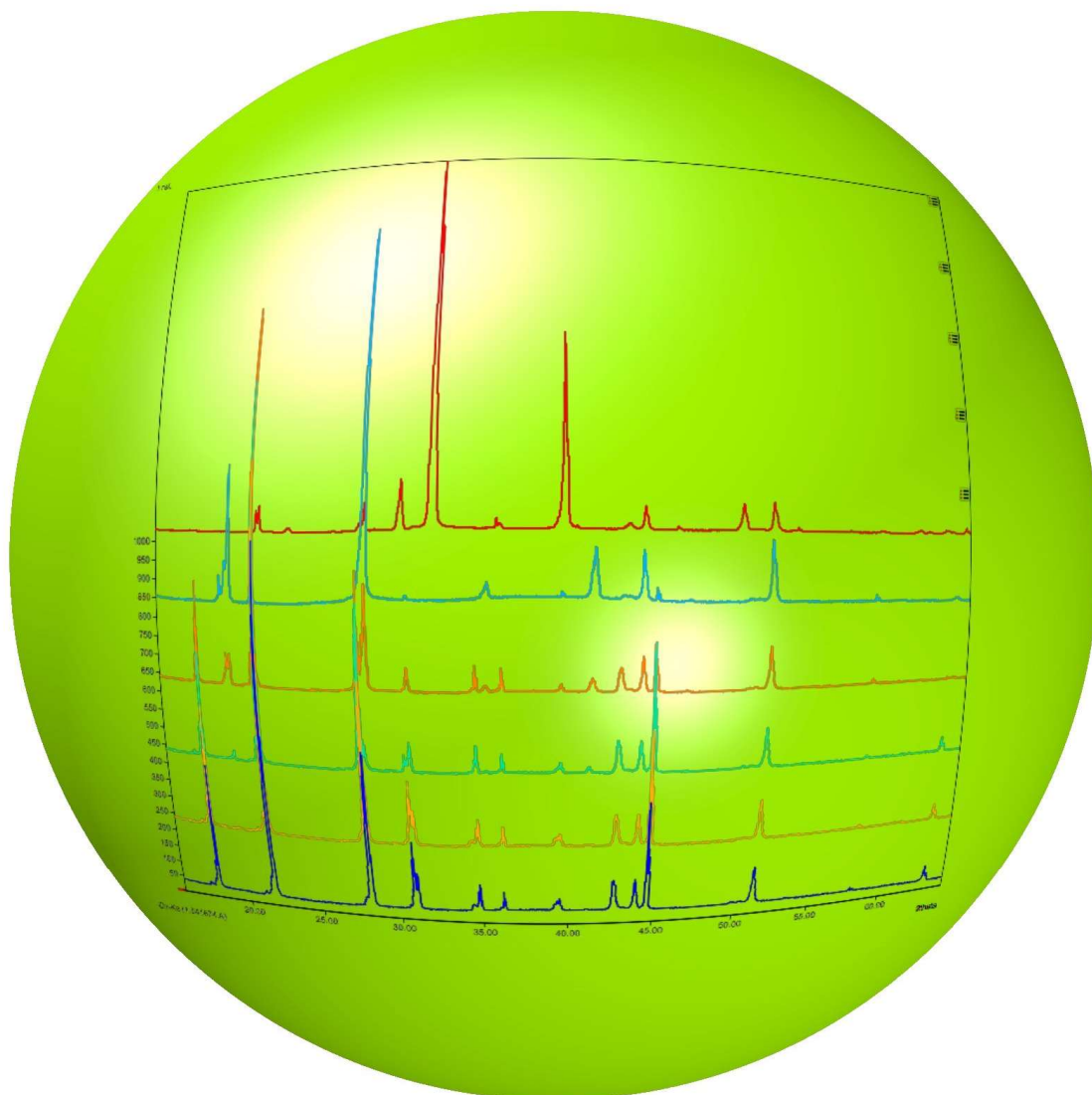


# MATCH!

Phase Identification from Powder Diffraction



**Progress in Phase Identification from Powder**

**Match!** is an easy-to-use software for phase identification from powder diffraction data. It compares the diffraction pattern of your sample to a database containing reference patterns, in order to identify the phases that are present. Additional knowledge about the sample like known phases, elements or density can be applied easily. In addition to this **qualitative analysis**, a **quantitative analysis** (using Rietveld refinement) can be performed as well.

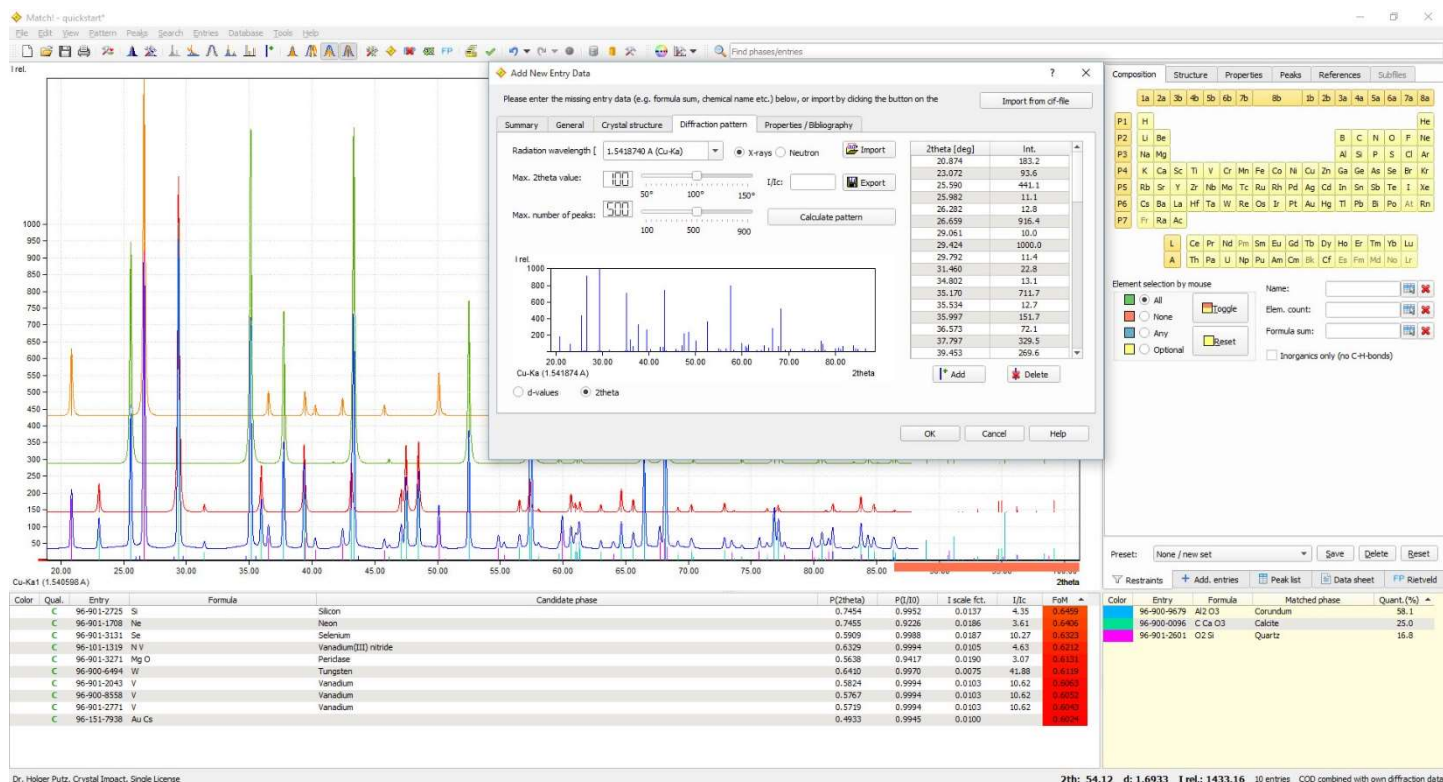
As reference database, you can apply the included **free-of-charge COD database**, use any **ICDD PDF database**, and/or **add your own powder diffraction**

**patterns** or crystal structure data.

You can quickly setup and run **Rietveld refinements** from within Match!, with the actual calculations being performed automatically, using the well-known program **FullProf** (by J. Rodriguez-Carvajal) in the background. Match! provides a **gentle introduction into Rietveld refinement**, from fully automatic operation (user-configurable) to the "Expert" mode.

More information and free-of-charge test version:

<https://www.crystalimpact.de/match>



## Feature

- Fast multiple phase identification from powder diffraction data
- Runs on Windows, Mac OS X and Linux
- Use free-of-charge reference patterns calculated from the COD (incl. I/I<sub>c</sub>), any ICDD PDF database product, and/or your own diffraction or crystal structure data in phase identification
- Indexing (using Treor or Dicvol)
- Crystal structure solution (using Endeavour)
- Rietveld refinement calculations (e.g. for quantitative analysis) using the well-known FullProf in the background
- Instant usage of additional information (known phases, elements, density, color etc.) using perpetual restraining
- Automatic residual searching with respect to identified phases
- Automatic raw data processing
- Comfortable background definition/modification using the mouse
- Convenient editing of peaks (add/shift/delete/fit) using the mouse
- Improved zooming and tracking using mouse or dialog
- Display several piled experimental patterns e.g. for comparison
- Semi-quantitative analysis (Reference Intensity Ratio method)
- Crystallite size estimation based on Scherrer approach

## System requirements

- **Windows** XP, Vista, Windows 7, Windows 8/8.1 or Windows 10 (32- or 64-bit); **macOS** 10.10 "Yosemite" or higher; **Linux** (Intel 32- or 64-bit), e.g. openSUSE, Ubuntu
- 2 GB of RAM
- 500 MB of free disc space
- **Supported diffraction data file formats (automatic recognition):** Bruker/Siemens, DBWS, DRON-3, G670, Inel, GNR (formerly Ital Structures), Jade/MDI/SCINTAG, JEOL Export, PANalytical/Philips, Rigaku, SCINTAG, Seifert, Shimadzu, various text files (profile or peak list data), Siemens, Sietronics, Stoe, TXRD export, X Powder

## Prices for new licenses\*

	non-profit org.	profit org.
Single licence	599 €	1,198 €
Site licence**	1,198 €	2,396 €
Campus licence***	2,396 €	4,792 €

\* Prices include a 3-year subscription for software updates; they do not include taxes that may be due.

\*\* Unlimited number of installations within one institute/dept.

\*\*\* Unlimited number of installations within one university/company.



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