

Chapter 3.7

LAZY-PULVERIX

**A program to calculate theoretical
X-ray and neutron diffraction
powder patterns**

KLAUS YVON, WOLFGANG JEITSCHKO and (the late) ERWIN PARTHE
LABORATOIRE DE CRISTALLOGRAPHIE AUX RAYONS-X
UNIVERSITE DE GENEVE
24 QUAI ERNEST ANSERMET
CH 1211 GENEVA 4 SWITZERLAND

For a further description of the program see *J.Appl.Cryst.* (1977), 10, 73-74

LAZY-PULVERIX consists of two programs

- LAZY - decodes the input data and prepares the data file for PULVERIX.
- PULVERIX - reads the file from LAZY and calculates the powder pattern.

3.6.1 Instructions Summary

3.6.1.1 General remarks

Calculations for several structures may be done in one run, the order in which the data instructions are given within a set is not important. When reading the data instructions, the program identifies the kind of data instruction by labels in columns 1-6 (e.g. TITLE, CONDIR, CELL, SPCGRP etc.). One END instruction must terminate each set and one FINISH instruction must follow the last END instruction.

The format of the TITLE, CELL, LATTICE, SYMTRY, SPCGRP, ATOM, END and FINISH instructions are compatible with the corresponding format of the X-RAY 76 program system, except small differences in certain prescriptions (see CELL and ATOM instructions). The minimum of input must contain TITLE, CELL, SPCGRP, ATOM, END and FINISH instructions. The SPCGRP instruction may be replaced by LATTICE and SYMTRY instructions and vice versa.

TITLE instruction - title

```
format(a2,a3,1x,17a4)
  cols
  1- 5  title      punch instruction label TITLE
  7-74  compnd    any alphanumeric information (for instance the
                    name of the substance)
```

CONDIT instruction - experimental conditions

```
format(3a2,4x,a4,f6.0,2f5.0,1x,a1,i2,a2,1x,a1)
  cols
  1- 6  condit    punch instruction label  CONDIR
  11-14 symwl    symbol for wavelength
                    adjust to the left of the field.
                    example
                    CUA1 = Copper K $\alpha$ 1 radiation.
                    the list of allowed symbols is given at
                    the end of this description.
                    wavelengths for which no symbol exist
```

```

                                must be given explicitly in cols 15-20.
                                if left blank Cuk $\alpha$  X-radiation is assumed.

                                neutron diffraction
                                leave columns 11-14 blank and give
                                value of wavelength in columns 15-20.
15-20  wl          wavelength in Angstrom
                                need not be given if symwl is specified
21-25  tl          lower theta -limit of calculation
26-30  th          upper theta -limit of calculation
                                if left blank tl=0 and th=89 degrees.
                                for Guinier cameras th is 45 degrees
      32  norm      tabular representation of the powder pattern.
                                blank intensities normalized to 1000
                                A intensities not normalized
                                N no tabular representation of
                                the powder pattern
33-34  image      graphic representation of the powder pattern.
                                blank no graphic output
                                integer graphic output of intensities
                                in steps of 1/(2*integer) of
                                theta
35-36  symlp      experimental techniques
                                blank Debye-Scherrer or powder-diffractometer
                                NE neutron diffraction
                                GN Guinier-de Wolff camera
                                GH Guinier-Haegg camera
                                the formulae for the lorentz -
                                polarisation factors are given
                                at the end of this description.
                                1 no lp-factor correction applied
      38  iano      correction for anomalous dispersion (x rays only)
                                blank correction is made
                                N no correction is made.

```

Notes

- It is recommended to compute all structures with neutral atoms. If form-factors for ionized atoms are used, the program will not make dispersion corrections. No dispersion correction will be made for neutron diffraction.
- If no CONDIT instruction is given, Cu-K α radiation and Debye-Scherrer technique is assumed. A correction for anomalous dispersion will be made and a complete powder pattern will be calculated.

CELL instruction - lattice constants

format (3a2, 7x, 3f8.0, 3f9.0)

```

cols
  1- 4  cell      punch instruction label CELL
14-21  a          lattice parameters in Angstrom and degrees
22-29  b          cubic          omit b, c, alpha, beta, gamma
30-37  c          hexagonal     omit b, alpha, beta,
                                and set gamma=120.
                                rhombohedral see note below
38-46  alpha      tetragonal   omit b, alpha, beta, gamma
47-55  beta       orthorhombic omit alpha, beta, gamma
56-64  gamma      monoclinic  omit alpha, gamma

```

- Rhombohedral should be calculated with hexagonal axes. If rhombohedral axes are desired the structure must be described in the triclinic system.

LATICE instruction - symmetry center and Bravais lattice (this instruction may be replaced by a SPCGRP instruction)

```
format (3a2, 2x, a1, 2x, a1)
cols
1- 6  lattice      punch instruction label  LATICE
    9  isymce      center of symmetry at origin
                        C   yes (centric)
                        A   no (acentric)

12  symbr         bravais lattice indicator
                        P   primitive
                        I   body centered
                        R   rhombohedral
                        F   face centered
                        A   a centered
                        B   b centered
                        C   c centered
blank primitive
```

For trigonal case

- P space groups give hexagonal lattice constants and set symbr=P
- R space groups :
 1. hexagonal cell set symbr=R. The program assumes the standard (obverse) setting.
 2. rhombohedral cell set symbr=P and give a,b, c and angles explicitly on CELL instruction (triclinic description).

If no LATICE instruction is given, a non centrosymmetric structure and a primitive lattice is assumed.

SYMTRY instruction - equivalent point positions (these instructions may be replaced by a SPCGRP instruction)

```
format (3a2, 45a1)
cols
1- 6  symtry      punch instruction label  SYMTRY
7-51 ipos         equivalent position coded in verbatim form.
                        include one instruction for each position.

rules for coding
coordinates are separated by commas,
fractions are written as two integers
                        separated by a slash.
blank spaces are ignored.

example      space group p 21/c
SYMTRY      X, Y, Z
SYMTRY      X, 1/2-Y, 1/2+Z
```

- If isymce=C then only one of any two centrosymmetric positions need to be given. If symbr= I,R,F,A,B or C, only one of the positions related by centering needs to be given
- If no SYMTRY instruction is given, x,y,z is automatically assumed, however if SYMTRY instructions are given then the x,y,z position MUST be included.

SPCGRP instruction - space group (this instruction may replace LATICE and SYMTRY instructions)

```
format (3a2,35a1)
  cols
  1- 6  spcgrp      punch instruction label SPCGRP
  8-17  ipos       Hermann-Mauguin symbol for the space group.
                        adjust to the left of the field

                        rules for coding

                        symmetry operators are separated by a
                        slash or by a blank.
                        the bar operation is coded as minus *-*
                        screw axes are given by two integers
                        that are not separated by a blank.
                        examples
                        P B C N, P 21/C, P -3

                        the list of allowed symbols is given at
                        the end of this description.
```

- For all other symbols the program may generate wrong equipoints without error messages.
- For non standard space group settings SYMTRY and LATICE instructions MUST be used.
- For centrosymmetric groups, the program assumes the setting having the centre at the origin.
- For R-space groups the hexagonal setting is assumed. R-space groups with rhombohedral axes must be simulated using LATICE and SYMTRY instructions corresponding to a triclinic description.

ATOM instruction - atom identifier and coordinates (use one atom instruction for each atom in the asymmetric unit).

```
format (3a2,1x,a4,a2,3f8.0,f6.0,f5.0)
  cols
  1- 4  atom      punch instruction label ATOM
  8-11  elemnt   symbol of element and ionisation state
                        adjust to the left of the field.
                        examples
                        CA  symbol for calcium(neutral)
                        CA2+ symbol for calcium(ionized)
                        only symbols listed in the table for
```

```

                                atom identification at the end of
                                this description can be given.
12-13 ide      sequence number or atom label. (may be left blank)
                                it is convenient to number or label
                                atoms of the same type.

14-21 x        x coordinate
22-29 y        y coordinate
30-37 z        z coordinate
                                only coordinates between -1.and+1. are
                                allowed
                                fractions may be given as integers
                                separated by a slash
                                example
                                atom H    1/3    2/3    0.512
                                is equivalent to
                                atom H    .33333 .666667 0.512
38-43 btemp    Webye-Waller factor
                                If left blank no temperature factor correction
                                will be made.
44-48 foccu    occupation factor
                                This factor is usually 1 (=full occupancy of the
                                site) but it may be smaller in disordered struc-
                                tures. If left blank full occupancy will be assumed

```

END instruction - terminates each set of data instructions

```

cols
1- 3 end      punch instruction label END

```

FINISH instruction - terminates the run

```

cols
1- 6 finish   punch instruction label FINISH

```

Note

This instruction must come after the last end instruction. It initiates execution of the program.

3.6.2 List of allowed symbols

3.6.2.1 Wavelengths (variable symwl on CONDIT instruction)

The lines contain the symbols for $K\alpha_1$, $K\alpha_2$ and the weighted average of the $K\alpha$ radiation for chromium, iron, copper, molybdenum and silver. The wavelengths in Angstrom are given in parentheses

CRA1 (2.28970)	CRA2 (2.29361)	CR (2.2909)
FEA1 (1.93604)	FEA2 (1.93998)	FE (1.9373)
CUA1 (1.54056)	CUA2 (1.54439)	CU (1.5418)
MOA1 (0.70930)	MOA2 (0.71359)	MO (0.7107)
AGA1 (0.55941)	AGA2 (0.56380)	AG (0.5608)

3.6.2.2. ATOM identification (variable elemnt on ATOM instruction)

The atom symbols are used to retrieve the scattering factors for X-rays and the nuclear (but not the magnetic) scattering factors for neutrons. Do NOT include the asterisk preceeding an atom symbol. This asterisk indicates atom identifiers that are allowed for both X-ray and neutron diffraction. All other symbols are allowed for x-ray diffraction only. Atom symbols followed by a point have a special meaning (see below). In case of X-ray diffraction use neutral atoms unless you know exactly what you want to calculate.

*AC	*C	*DY	*HE	LU3+	NP4+	*PU	SI.	TM3+
AC3+	C.	DY3+	*HF	*MG	NP6+	PU3+	SI4+	*U
*AG	*CA		HF4+	MG2+		PU4+	*SM	U3+
AG1+	CA2+	*ER	*HG	*MN	*O	PU6+	SM3+	U4+
AG2+	*CD	ER3+	HG1+	MN2+	O1-		*SN	U6+
*AL	CD2+	*EU	HG2+	MN3+	O2-	RA	SN2+	
AL3+	*CE	EU2+	*HO	MN4+	*OS	RA2+	SN4+	*V
*AM	CE3+	EU3+	HO3+	*MO	OS4+	*RB	*SR	V2+
*AR	CE4+			MO3+		RB1+	SR2+	V3+
*AS	CF	*F	*I	MO5+	*P	*RE		V5+
AT	*CL	F1-	I1-	MO6+	*PA	*RH	*TA	
*AU	CL1-	*FE	*IN		*PB	RH3+	TA5+	*W
AU1+	*CM	FE2+	IN3+	*N	PB2+	RH4+	*TB	W6+
AU3+	*CO	FE3+	*IR	*NA	PB4+	RN	TB3+	
	CO2+	FR	IR3+	NA1+	*PD	*RU	*TC	*XE
*B	CO3+		IR4+	*NB	PD2+	RU3+	*TE	
*BA	*CR	*GA		NB3+	PD4+	RU4+	*TH	*Y
BA2+	CR2+	GA3+	*K	NB5+	PM		TH4+	Y3+
*BE	CR3+	*GD	K1+	*ND	PM3+	*S	*TI	*YB
BE2+	*CS	GD3+	*KR	ND3+	PO	*SB	TI2+	YB2+
*BI	CS1+	*GE		*NE	*PR	SB3+	TI3+	YB3+
BI3+	*CU	GE4+	*LA	*NI	PR3+	SB5+	TI4+	
BI5+	CU1+		LA3+	NI2+	PR4+	*SC	*TL	*ZN
BK	CU2+	*H	*LI	NI3+	*PT	SC3+	TL1+	ZN2+
*BR		H.	LI1+	*NP	PT2+	*SE	TL3+	*ZR
BR1-	D.	H1-	*LU	NP3+	PT4+	*SI	*TM	ZR4+

Symbols with special meaning

H. Hydrogen HF scattering factor
 C. Carbon HF scattering factor
 SI. Silicon HF scattering factor
 D. Deuterium (for neutron diffraction ONLY)
 O2-. taken from Acta Cryst. vol.19, p.486(1965).

3.6.2.3 SPACE GROUP symbols (variable ipos on SPCGRP instruction)

Do NOT include the star preceeding some of the symbols. The star indicates centro-symmetric space groups which have been described with several settings. The program generates only the setting with the centre of symmetry at the origin of the unit cell. WARNING - a symbol that does not figure in this list may yield wrong equipoints.

3.7 LAZY-PULVERIX - Data Menu

WinGX v1.71

TRICLINIC

P 1 P -1

MONOCLINIC

P 2	P 21	C 2	P M	P C
C M	C C	P 2/M	P 21/M	C 2/M
P 2/C	P 21/C	C 2/C		

THE POINT POSITIONS GENERATED FROM THESE SYMBOLS CORRESPOND TO THE MONOCLINIC SETTING WITH B AS UNIQUE AXIS (ALPHA=GAMMA=90.)

ORTHORHOMBIC

P 2 2 2	P 2 2 21	P 21 21 2	P 21 21 21	C 2 2 21
C 2 2 2	F 2 2 2	I 2 2 2	I 21 21 21	P M M 2
P M C 21	P C C 2	P M A 2	P C A 21	P N C 2
P M N 21	P B A 2	P N A 21	P N N 2	C M M 2
C M C 21	C C C 2	A M M 2	A B M 2	A M A 2
A B A 2	F M M 2	F D D 2	I M M 2	I B A 2
I M A 2	P M M M	*P N N N	P C C M	*P B A N
P M M A	P N N A	P M N A	P C C A	P B A M
P C C N	P B C M	P N N M	*P M M N	P B C N
P B C A	P N M A	C M C M	C M C A	C M M M
C C C M	C M M A	*C C C A	F M M M	*F D D D
I M M M	I B A M	I B C A	I M M A	

TETRAGONAL

P 4	P 41	P 42	P 43	I 4
I 41	P -4	I -4	P 4/M	P 42/M
*P 4/N	*P 42/N	I 4/M	*I 41/A	P 4 2 2
P 4 21 2	P 41 2 2	P 41 21 2	P 42 2 2	P 42 21 2
P 43 2 2	P 43 21 2	I 4 2 2	I 41 2 2	P 4 M M
P 4 B M	P 42 C M	P 42 N M	P 4 C C	P 4 N C
P 42 M C	P 42 B C	I 4 M M	I 4 C M	I 41 M D
I 41 C D	P -4 2 M	P -4 2 C	P -4 21 M	P -4 21 C
I -4 M 2	P -4 C 2	P -4 B 2	P -4 N 2	P -4 M 2
I -4 C 2	P -4 2 M	I -4 2 D	P 4/M M M	P 4/M C C
*P 4/N B M	*P 4/N N C	P 4/M B M	P 4/M N C	*P 4/N M M
*P 4/N C C	P 42/M M C	P 42/M C M	*P 42/N B C	*P 42/N N M
P 42/M B C	P 42/M N M	*P 42/N M C	*P 42/N C M	I 4/M M M
I 4/M C M	*I 41/A M D	*I 41/A C D		

TRIGONAL

P 3	P 31	P 32	R 3	P -3
R -3	P 3 1 2	P 3 2 1	P 31 1 2	P 31 2 1
P 32 1 2	P 32 2 1	R 3 2	P 3 M 1	P 3 1 M
P 3 C 1	P 3 1 C	R 3 M	R 3 C	P -3 1 M
P -3 1 C	P -3 M 1	P -3 C 1	R -3 M	R -3 C

ALL R-SPACE GROUPS REFER TO THE HEXAGONAL SETTING
HEXAGONAL

P 6	P 61	P 65	P 62	P 64
P 63	P -6	P 6/M	P 63/M	P 6 2 2
P 61 2 2	P 65 2 2	P 62 2 2	P 64 2 2	P 63 2 2
P 6 M M	P 6 C C	P 63 C M	P 63 M C	P -6 M 2
P -6 C 2	P -6 2 M	P -6 2 C	P 6/M M M	P 6/M C C
P 63/M C M	P 63/M M C			

CUBIC

P 2 3	F 2 3	I 2 3	P 21 3	I 21 3
-------	-------	-------	--------	--------

P M 3	*P N 3	F M 3	*F D 3	I M 3
P A 3	I A 3	P 4 3 2	P 42 3 2	F 4 3 2
F 41 3 2	I 4 3 2	P 43 3 2	P 41 3 2	I 41 3 2
P -4 3 M	F -4 3 M	I -4 3 M	P -4 3 N	F -4 3 C
I -4 3 D	P M 3 M	*P N 3 N	P M 3 N	*P N 3 M
F M 3 M	F M 3 C	*F D 3 M	*F D 3 C	I M 3 M
I A 3 D				

3.6.3 Formulae for the Lorentz-polarisation factors

3.6.3.1 Debye-Scherrer technique

$$l = 1.0/(\sin^2\theta \times \cos\theta)$$

$$p = (1.0 + \cos^2 2\theta)/2.$$

3.6.3.2 Guinier technique

$$l = 1./(\sin^2\theta \times \cos\theta \times \cos(2\theta - \beta))$$

β = angle between the normal to the specimen and the direction of the incident beam.

$$p = (1.0 + \cos^2 2\theta \times \cos^2 2\alpha)/(1 + \cos^2 2\alpha)$$

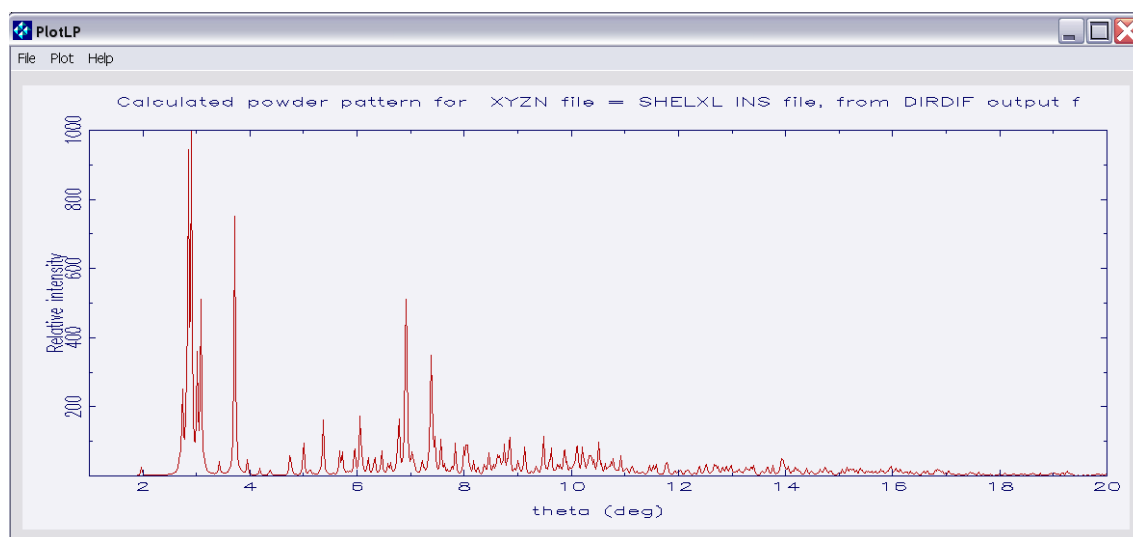
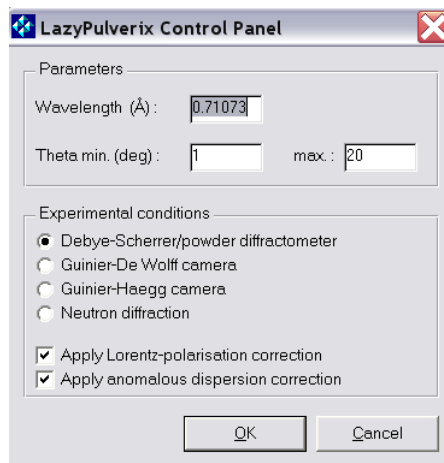
α = diffraction angle of monochromator.

α and β depend on the geometry of the Guinier camera and the d spacing of the reflecting planes of the monochromator crystal. For Guinier cameras other than Guinier-de Wolff or Guinier-Haegg or for monochromator crystals other than quartz changes in the program have to be made (see *remarks for adapting the program* in the source deck of PULVERIX).

3.6.4. LAZY-PULVERIX implementation in WinGX

The programs LAZY and PULVERIX have been compiled essentially unaltered, but the limits have been substantially increased. A maximum of 5000 reflections may be calculated, up to 1000 atoms of one type may be entered, and 20 different atomic types are allowed. A new program PLOTLP has been written which plots the calculated intensities listed in the output file of PULVERIX (called PULVERIX.LST). This program is run automatically after LAZY and PULVERIX. Lineshapes may be Lorentzian, Gaussian or a 50:50 mix and linewidths may be varied. Monochrome or colour PostScript files may also be written. The plot parameters may be selected from the Plot[Parameters] sub-menu.

The program instructions are written by the GUI shown below, and the above input description is for informational purposes only. The calculated spectrum is displayed as a plot, which may be saved in PostScript format.



3.6.5. Disclaimer

Although the program has been extensively tested by its authors, no warranty is made as to the accuracy and functioning. The values of wavelengths, scattering factor constants and the equi-points used by the program are part of the output (file *PULVERIX.LST*) and the user is advised to compare them with the values given in the International Tables for Crystallography.