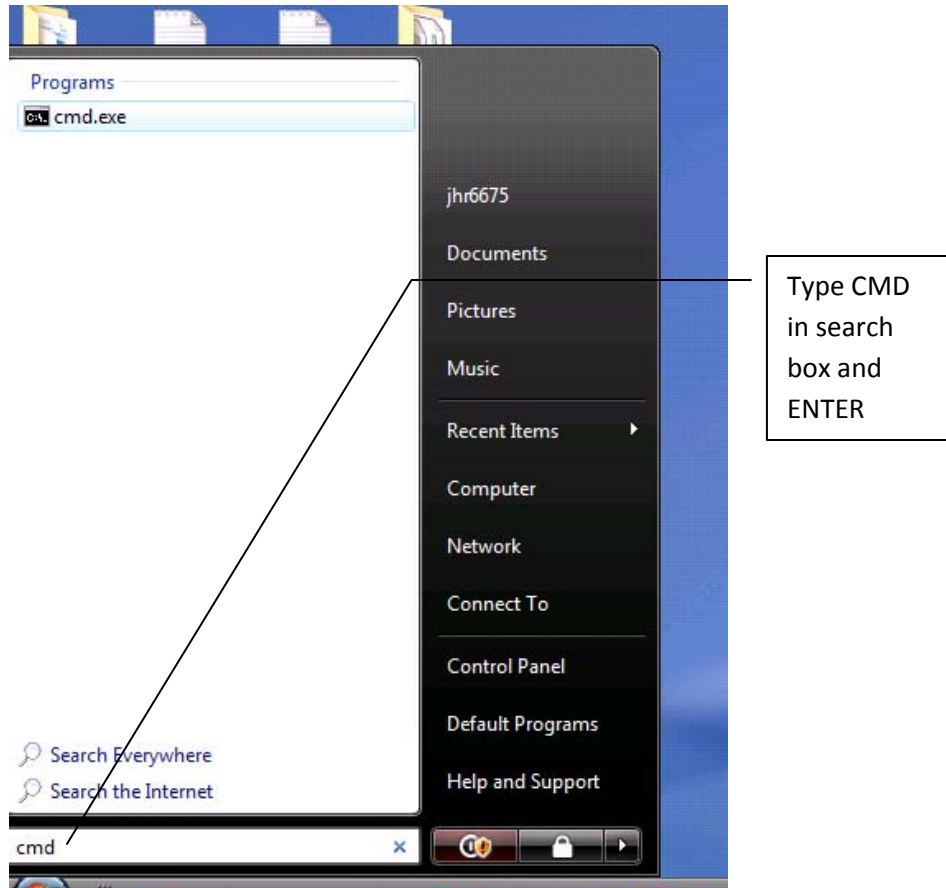
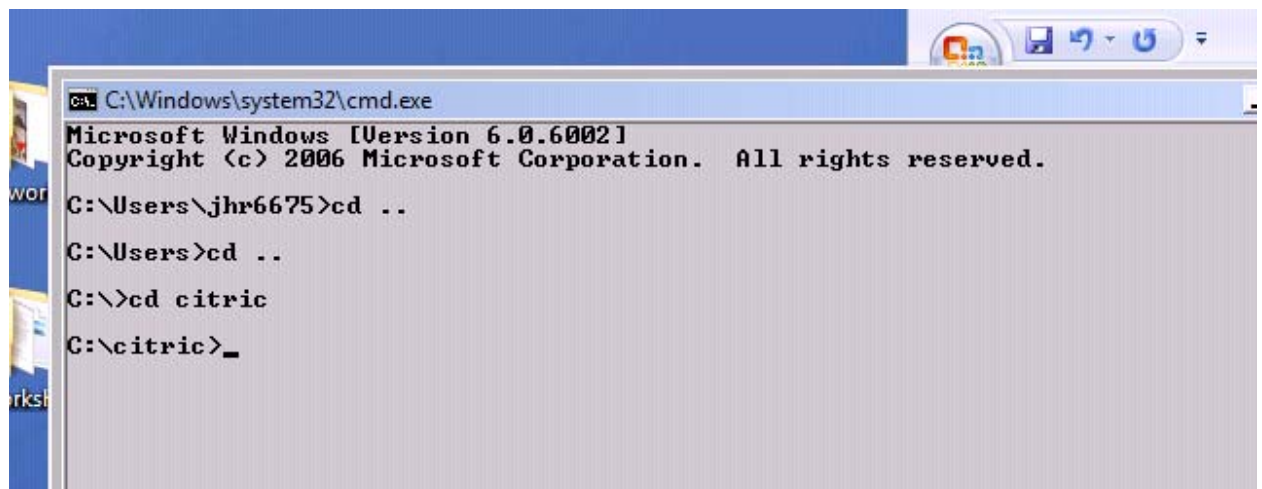


- Down load CRYSFIRE, PowderCell and Citric zip files.
- Unzip all folders.
- Move Citric, CRYSFIRE and PowderCell folders to the C: drive
- Call up a command window



- 
- Navigate (with the cd command) to the citric folder



-

- Type “..\CRYSFIRE\CRYSFIRE.BAT” at the command prompt

```

C:\Windows\system32\cmd.exe
Microsoft Windows [Version 6.0.6002]
Copyright (c) 2006 Microsoft Corporation. All rights reserved.

C:\Users\jhr6675>cd..

C:\Users>cd..

C:\>cd citric

C:\citric>..\CRYSFIRE\CRYSFIRE.BAT

```

- Follow the prompts
- Type IM at the prompt

```

** Crysfire 2004 **   An Interactive Powder Indexing Support System
                        Robin Shirley
                        41 Guildford Park Avenue
                        Guildford, Surrey GU2 7NL, England
                        v9.52.11 (Delphi), 24Nov04 (32-bit beta)
                        running under Windows NT-family: (NT, 2000, XP)

```

```

Default directory for datasets: start-up directory (Use CD command to c
Max no. of obs lines: 1000
Max no. of calc lines: 20000
Main screen output will repeat to <Dset>.CF0 file by default
** No observed pattern loaded yet - to input one, run L0, IM or OB **
Crysfire 2004 v9.52.11 - Next command or ? for Help: IM

```

- The AS at the next prompt

```

Please select one of the above import formats (or EX to exit)
Enter AS, EU, PX, UD, WF, XF or EX (default: EX): AS_

```

- When asked to proceed type Y and at the next prompt hit a RETURN (no name)

```

If your data file does not fit within the above limitati
either first edit it manually, or use some other file-coi
Are you ready to proceed (Y/N, default=Y)? y
3-char extension for the datafile name (default: TXT):

```

- At the next screen enter “citric”

Give / directory information followed by listing of textfiles  
in the currently selected directory, sorted into name order: \*.TXT

Directory/File specification is: .\\*.TXT

CITRIC.TXT

Name of ASCII datafile to be loaded (omitting .TXT, default = exit): citric\_

- 
- Use the defaults for the next 7 queries. After the How many fields query you will see

Reading data file...

Reading peaks list...

No.	2Theta	d	Q
1	11.479	7.702519	168.5522
2	14.905	5.938965	283.5166
3	15.185	5.830017	294.2120
4	16.973	5.219682	367.0387
5	17.928	4.943613	409.1769
6	18.180	4.875866	420.6264
7	19.016	4.663240	459.8587
8	19.720	4.498271	494.2068
9	19.877	4.463233	501.9968
10	20.538	4.320992	535.5908
11	22.267	3.989129	628.4110
12	22.320	3.979939	631.3167
13	23.075	3.851261	674.2083
14	23.721	3.747836	711.9325
15	24.295	3.660571	746.2810
16	24.424	3.641615	754.0705
17	24.995	3.559703	789.1735
18	27.117	3.285702	926.2829
19	28.316	3.149241	1008.2964

- 
- For the working name enter citric

13	23.075	3.851261	674.2083
14	23.721	3.747836	711.9325
15	24.295	3.660571	746.2810
16	24.424	3.641615	754.0705
17	24.995	3.559703	789.1735
18	27.117	3.285702	926.2829
19	28.316	3.149241	1008.2964

Type ENTER to continue or F (and ENTER) to Finish without further pauses:

20	28.785	3.099011	1041.2471
21	29.283	3.047393	1076.8198

What is the working name for this new dataset (1 to 8 chars): citric

- 
- Save the dataset to citric and continue

Save the current dataset in Crysfire (CDI) format

Dataset name (max 8 letters/digits, default: citric) or \*Q to quit: citric\_

- 
- Give it a title and hit return (twice)

- You should see the Crysfire 2004 v9.52.11 line and the command prompt
- Type index and return
- Now select the indexing programs. We will run several as an internal check for the unit cell.
- Type TP first.
- Accept all default values
- Start the program. When it is complete a text file will appear. Close it for now.
- Hit a return and a 2<sup>nd</sup> text file will appear. This is a summary of what TP found. The figure of merit is off the scale (that is good).
- Close this text file. (you may need to do this twice).
- Repeat the process for the next indexing program.
- At the Crysfire 2004 v9.52.11 line type Index
- Chose DV this time
- Here is the summary file

```

Running summary of all solutions logged so far for this dataset name,
sorted in descending order of I20 then Merit (as defined by each program)

I20  Merit    Volume  V/V1  BL  IndexProg    Date      Time      Pedig    a        b        c
20*****  904.933  1.00  P  TAUPv3.3a    23Feb10  15:59:40  Ort__1   6.2985   9.3265  15.4050
20 5500.6    904.933  1.00  P  DICVOL91/log 23Feb10  16:04:50  Ort__1  15.4050   9.3265   6.2985
20 4950.1    904.930  1.00  P  DICVOL91/log 23Feb10  16:04:50  Mon__7  19.8997   9.3265   6.2985
20 4623.2    904.935  1.00  P  DICVOL91/log 23Feb10  16:04:50  Mon__4  16.6429   9.3265   6.2985
20 4367.2    904.935  1.00  P  DICVOL91/log 23Feb10  16:04:50  Mon__2  15.4050   6.2985   9.3265
20 4155.1    904.935  1.00  P  DICVOL91/log 23Feb10  16:04:50  Mon__1   9.3265  15.4050   6.2985
20 3998.5    904.937  1.00  P  DICVOL91/log 23Feb10  16:04:50  Mon__6  18.0082   6.2985   9.3265
20 3693.3    904.930  1.00  P  DICVOL91/log 23Feb10  16:04:50  Mon__5  11.2541  15.4050   6.2985
20 3265.0    904.935  1.00  P  DICVOL91/log 23Feb10  16:04:50  Mon__3  15.4050   9.3265   6.2985

```

- -
- Notice that there are several DP outputs the one with the highest Merit is of interest.
- Try out other indexing programs.
- If you do you may see something like this

```

2017691.8    904.94    1.00  P  ITO12/log    23Feb10  16:06:29    461    9.3265  15.4051   6.2985
2014646.88   904.934   1.00  P  FJZNV6.22a   23Feb10  16:07:05    241    9.3265  15.4050   6.2985
20*****    904.933   1.00  P  TAUPv3.3a    23Feb10  15:59:40  Ort__1   6.2985   9.3265  15.4050
20 5500.6    904.933   1.00  P  DICVOL91/log 23Feb10  16:04:50  Ort__1  15.4050   9.3265   6.2985

```

- 
- ITO, FJZM, TAUP and DICVOL will find solutions. The highest are for the orthorhombic cell 6.2985 9.3265 and 15.4050 (a,b and c are interchangeable!)
- CRYSFIRE has found a solution.
- It is time to go to PowderCell.

- Powder Cell .. Navigate your way to the powder cell folder (in explorer) and double click the PCW icon
- Goto file/new and input the cell in the Structure Data box. For now allow the space group to be P1.

structure data

initial data

Citric

lattice constants

space-group No 1 setting 1 P 1 atoms in cell: 0.0 (0 pos)

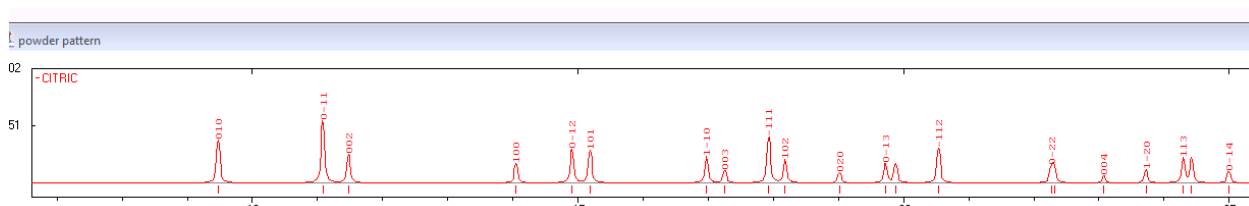
a	b	c	$\alpha$	$\beta$	$\gamma$
6.2985	9.3265	15.4050	90.0000	90.0000	90.0000

cell vol: 0.000 Å<sup>3</sup> density: 0.000 g/cm<sup>3</sup> rel. mass: 0.000 mass abs coef: 0.000 cm<sup>2</sup>/g

name	Z	ion	Wyck	x	y	z	SOF	B (temp)

+ atom - atom comment ? Help X Cancel OK

- 
- Point to Ok. The program will say no valid atom coordinates. Ignore this.
- An idealized pattern will appear at the bottom of the screen



- 
- That's good. Load the powder data.



- Notice that there are a few peaks in the observed Citric acid pattern that are missing from the calculated. This is an impurity, most likely the anhydrous form of citric acid.
- This concludes this tutorial.
- You may wish to continue by inputting the coordinates of citric acid in PowderCell and refining the pattern.