

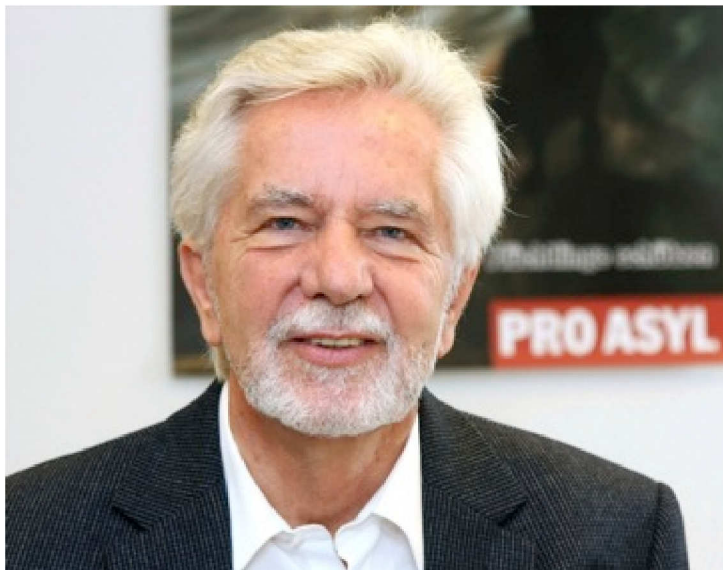
NSXTool - new Software for Data Reduction at Monochromatic Single Crystal Diffractometers

Tobias E. Schrader
Andreas Ostermann
Eric Pellegrini,
Jonathan Fisher
Laurent Chapon

MLZ is a cooperation between:

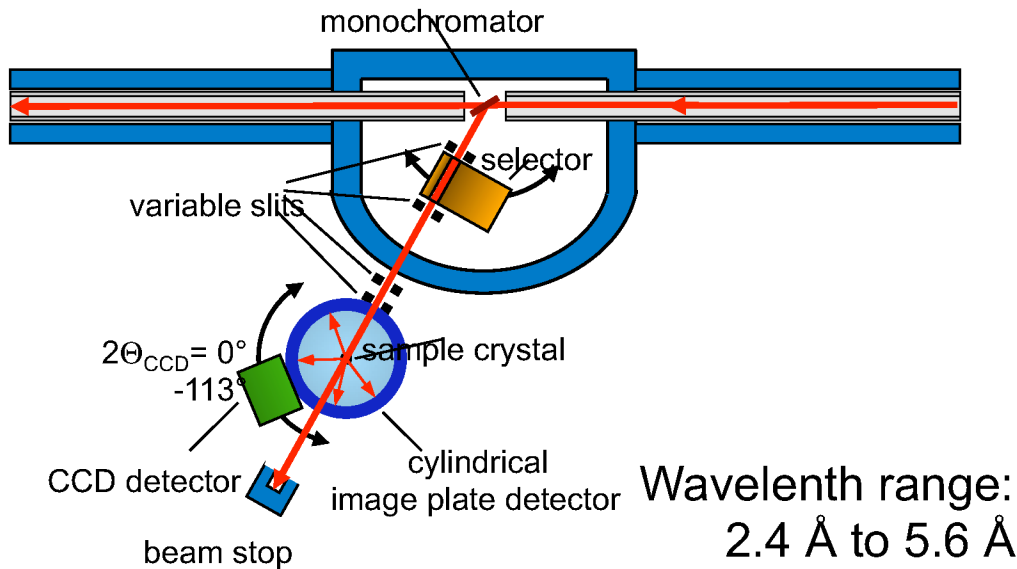
Jürgen Miksch, Founder of BISS

- 1984 bis 1993 als stellvertretender Direktor der Evangelischen Akademie Tutzing

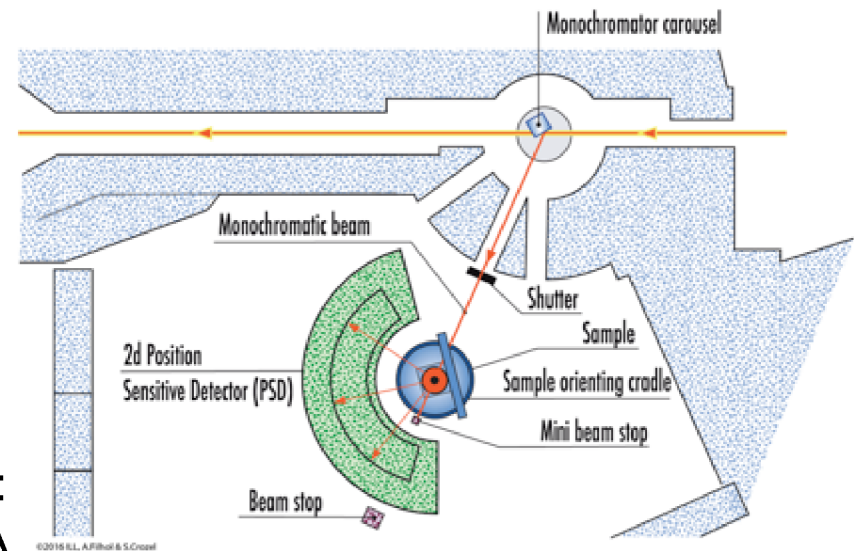


Instruments BIODIFF at MLZ and D19 at ILL

- Both monochromatic neutron diffractometers with cylindrical detector

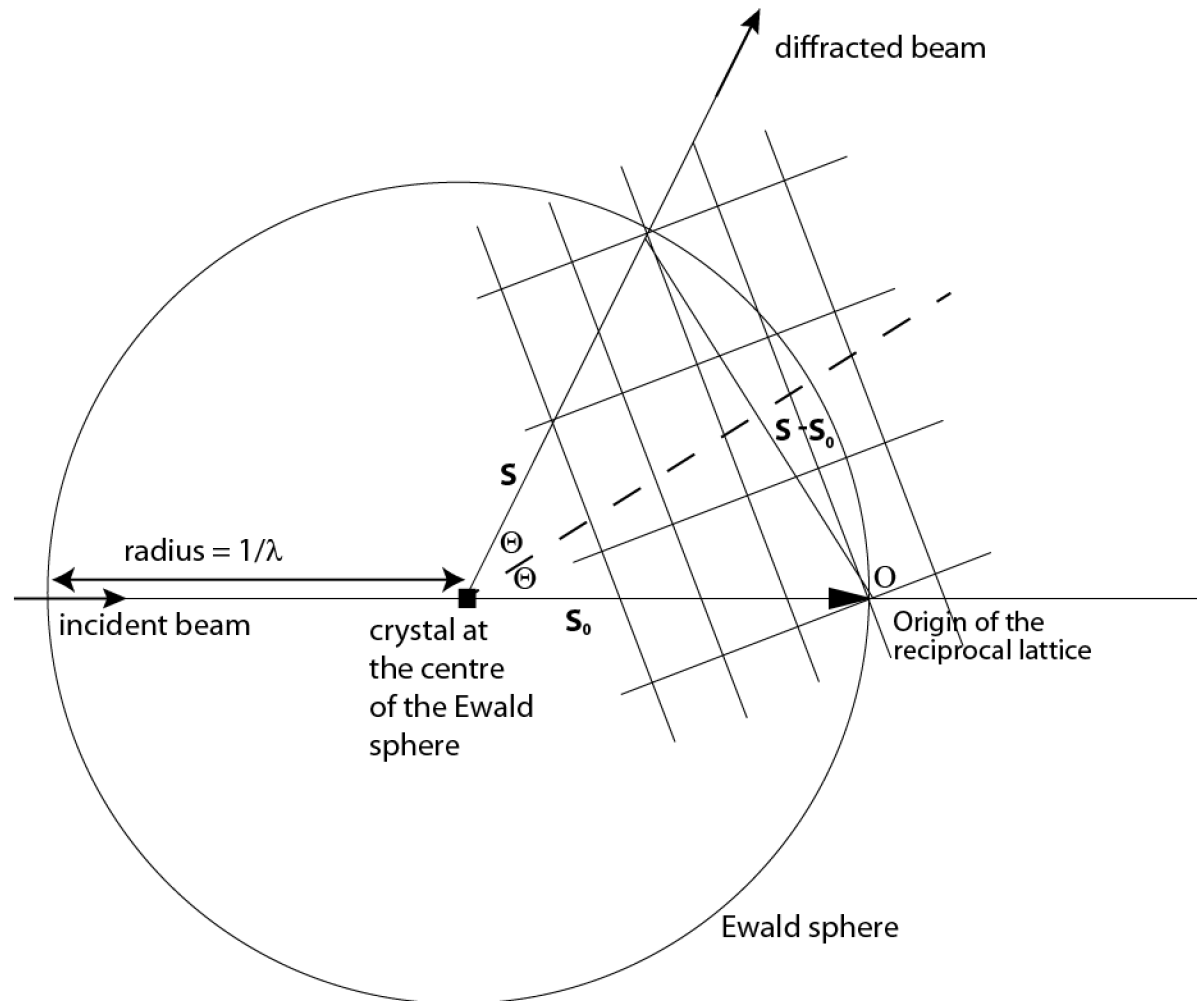


BIODIFF

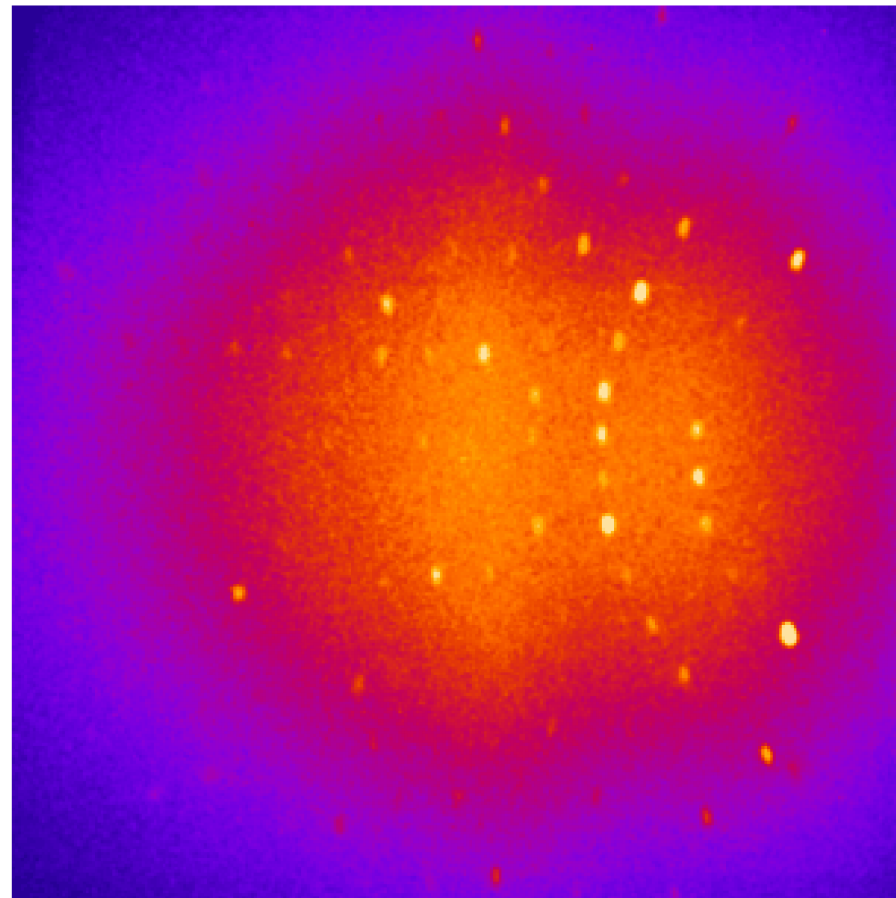
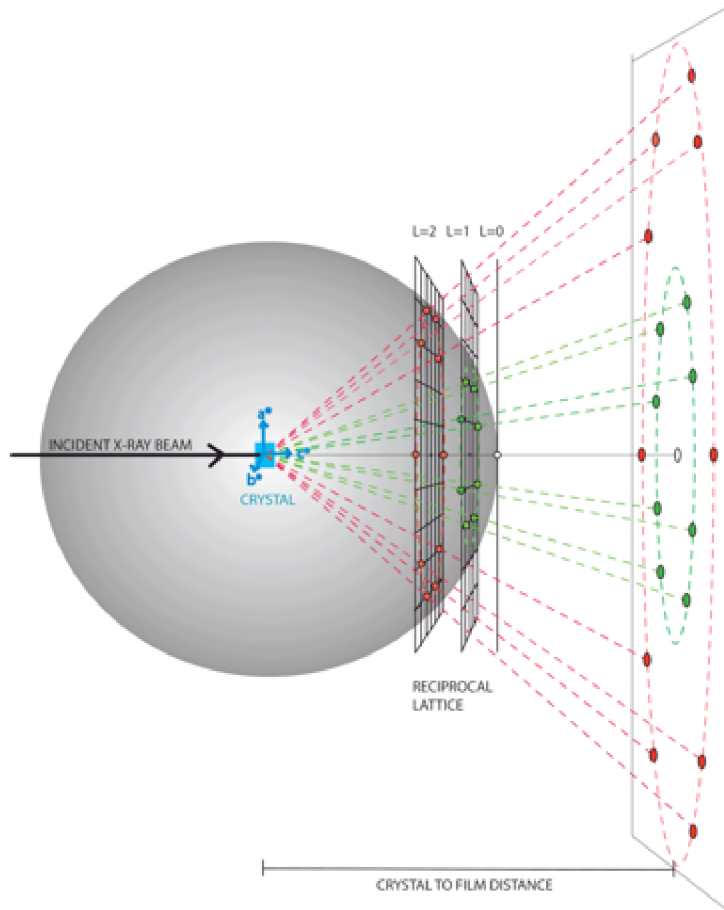


D19

Ewald construction and Bragg's Law



Myoglobin protein crystal (deuterated mother liquor) full data set recorded with CCD





 prim.
 beam

BioDiff: exposure time per frame: 20 minutes,
 sample: Myoglobin in deuterated mother liquor

Flow chart of data treatment and model building

Scans at varying crystal orientation
Scan := Series of detector images

Data reduction
 - determination of crystal orientation, unit cell dimensions etc.
 - Calculating integral of reflection intensities

hkl-list for each scan:
h k l Intensity Intensity error

Scaling of each hkl list to match each other

Unified hkl-list of measurement := complete data set

Calculation of a first map

-SCALA (CCP4-program package)

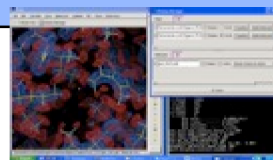
Additional information from the
solution of the phase problem

Structure refinement
 -Refinement of atom coordinates displacements
 - Calculation of scattering density maps (neutrons)
 or electron density maps (x-rays)

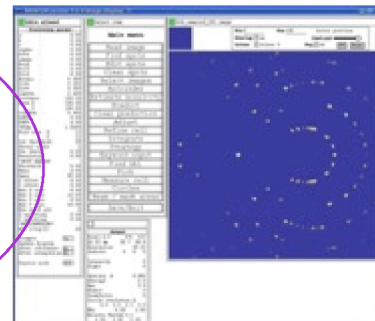
Map-plotting
 - inspection of model to fit the map)
 - real space changes and refinement to the model



-nCNS
-PHENIX



-XtalView
-Coot



-MOSFLM
-HKL-denzo
(commercial)

Peak search with hkl DENZO

The screenshot shows a desktop environment with the following components:

- hkl DENZO Main Window:** Displays a diffraction pattern with red circles around peaks. A sidebar on the right shows processing parameters: $I_{max}=1045720$, $I=1926$, $[213.8, .963, 4]$. Below this, it says "HKL Processing System" and "U. Minor, Z. Otwinowski". A vertical color scale bar is also visible.
- Terminal Window (jcms@phys:~/DENZO/denzo_1_96/real_data):** Shows a file listing:


```

-rw-rw-r-- 1 jcms jcms 5512500 Nov 5 18:03 309_01_001.raw
-rw-rw-r-- 1 jcms jcms 1013 Nov 5 18:04 auto_index_sim_spotb.dat
[jcms@phys real_data]$ ls -ltr
total 16148
-rwxr-xr-x 1 jcms jcms 4507524 Nov 2 18:51 Freimessen_291011_01_274_0.tif
-rw-rw-r-- 1 jcms jcms 5512500 Nov 2 19:01 Freimessen_291011_01_274_001.raw
-rw-rw-r-- 1 jcms jcms 496 Nov 2 19:03 refineone.dat-
-rw-rw-r-- 1 jcms jcms 474 Nov 2 19:03 refineall.dat
-rw-rw-r-- 1 jcms jcms 467 Nov 2 19:03 cr_info
-rwxr-xr-x 1 jcms jcms 626060 Nov 2 19:03 denzo
-rwxr-xr-x 1 jcms jcms 325804 Nov 2 19:03 xdisp
-rw-rw-r-- 1 jcms jcms 1049 Nov 2 19:05 auto_index_sim_spotb.dat-
-rw-rw-r-- 1 jcms jcms 1269 Nov 2 19:07 peaks.file
-rw-rw-r-- 1 jcms jcms 441 Nov 2 19:09 refineone.dat
-rw-rw-r-- 1 jcms jcms 14288 Nov 2 19:13 hklpredictions
-rw-rw-r-- 1 jcms jcms 1013 Nov 5 18:04 auto_index_sim_spotb.dat
-rw-rw-r-- 1 jcms jcms 5512500 Nov 5 18:11 309_01_001.raw
[jcms@phys real_data]$
      
```
- Web Browser Window (Reaktor-Info):** Displays the website <http://www.frm2.tum.de/intern/funktionen/reaktor-info/index.html>. It shows a power reading of "19.8 MW" and a button labeled "Shutterstellung NL-Anlage".

auto-index

Applications Places System Sat Nov 5, 18:24 JCNS

/home/jcns/DENZO/denzo_1_96/real_data/309_01_001.raw

zoom wind Write/Print A/D test Floor Up Floor Down reverse color Remove pred Full scale Go Show Overl Peak Sear Edit P.S. Help dsa bright close Frame

imax=1046720
I=1805
[298.6, 506.6]

HWL
Processing
System
U. Minor
Z. Otwinowski

7272
6383
5454
4545
3636
2727
1818
909
0

new date was send, updating picture wind

jcns@phys:~/DENZO/denzo_1_96/real_data

File Edit View Terminal Help

```

autoindex unit cell 35.44 31.09 64.92 90.00 105.53 90.00
crystal rotx, roty, rotz -112.379 87.484 0.804
Autoindex Xbeam, Ybeam 225.65 490.29
position 73 chi**2 x 11.35 y 8.84 pred. decrease: 0.000 * 73 = 0.0
partiality 73 chi**2 0.64 pred. decrease: 0.000 * 73 = 0.0
Angles equivalent by space group symmetry for:
vertical axis 1 0 0
spindle axis 0 0 1
crystal rotx 67.621 roty 92.516 rotz 0.804
rotz -112.379 roty 87.484 rotz -179.196
crystal rotx -112.379 roty 87.484 rotz 0.804
rotz 67.621 roty 92.516 rotz -179.196

```

jcns@phys:~/DENZO/denzo_1_96/real_data

File Edit View Terminal Help

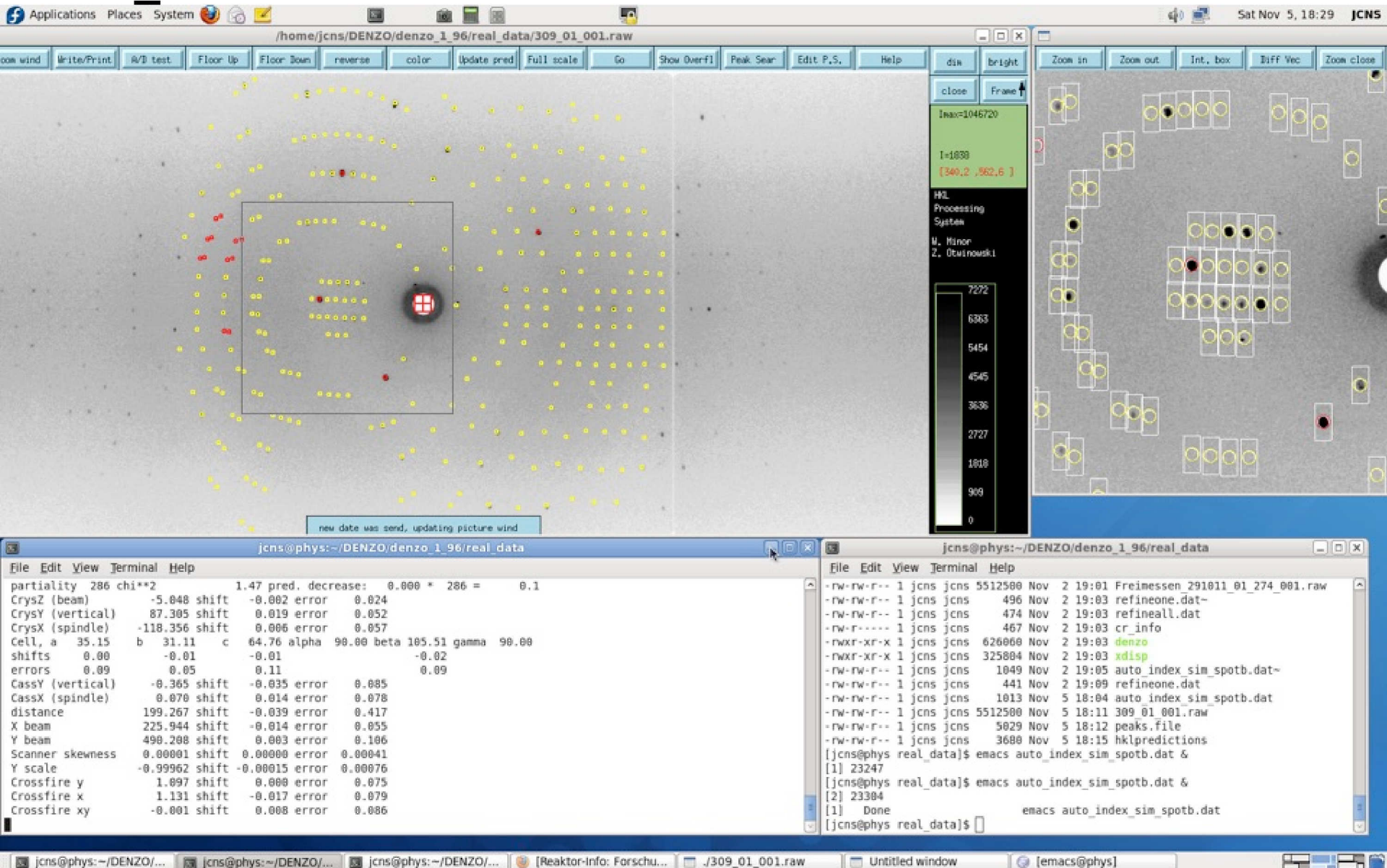
```

[jcns@phys real_data]$ ls -ltr
total 16140
-rwxr--r-- 1 jcns jcns 4507524 Nov 2 18:51 Freimessen_291011_01_274_0.tif
-rw-rw-r-- 1 jcns jcns 5512500 Nov 2 19:01 Freimessen_291011_01_274_001.raw
-rw-rw-r-- 1 jcns jcns 496 Nov 2 19:03 refineone.dat
-rw-rw-r-- 1 jcns jcns 474 Nov 2 19:03 refineall.dat
-rw-rw-r-- 1 jcns jcns 467 Nov 2 19:03 cr_info
-rwxr-xr-x 1 jcns jcns 626060 Nov 2 19:03 denzo
-rwxr-xr-x 1 jcns jcns 325804 Nov 2 19:03 xdisp
-rw-rw-r-- 1 jcns jcns 1049 Nov 2 19:05 auto_index_sim_spotb.dat
-rw-rw-r-- 1 jcns jcns 441 Nov 2 19:09 refineone.dat
-rw-rw-r-- 1 jcns jcns 1013 Nov 5 18:04 auto_index_sim_spotb.dat
-rw-rw-r-- 1 jcns jcns 5512500 Nov 5 18:11 309_01_001.raw
-rw-rw-r-- 1 jcns jcns 5029 Nov 5 18:12 peaks.file
-rw-rw-r-- 1 jcns jcns 3680 Nov 5 18:15 hklinpredictions
[jcns@phys real_data]$ emacs auto_index_sim_spotb.dat &
[1] 23247
[jcns@phys real_data]$

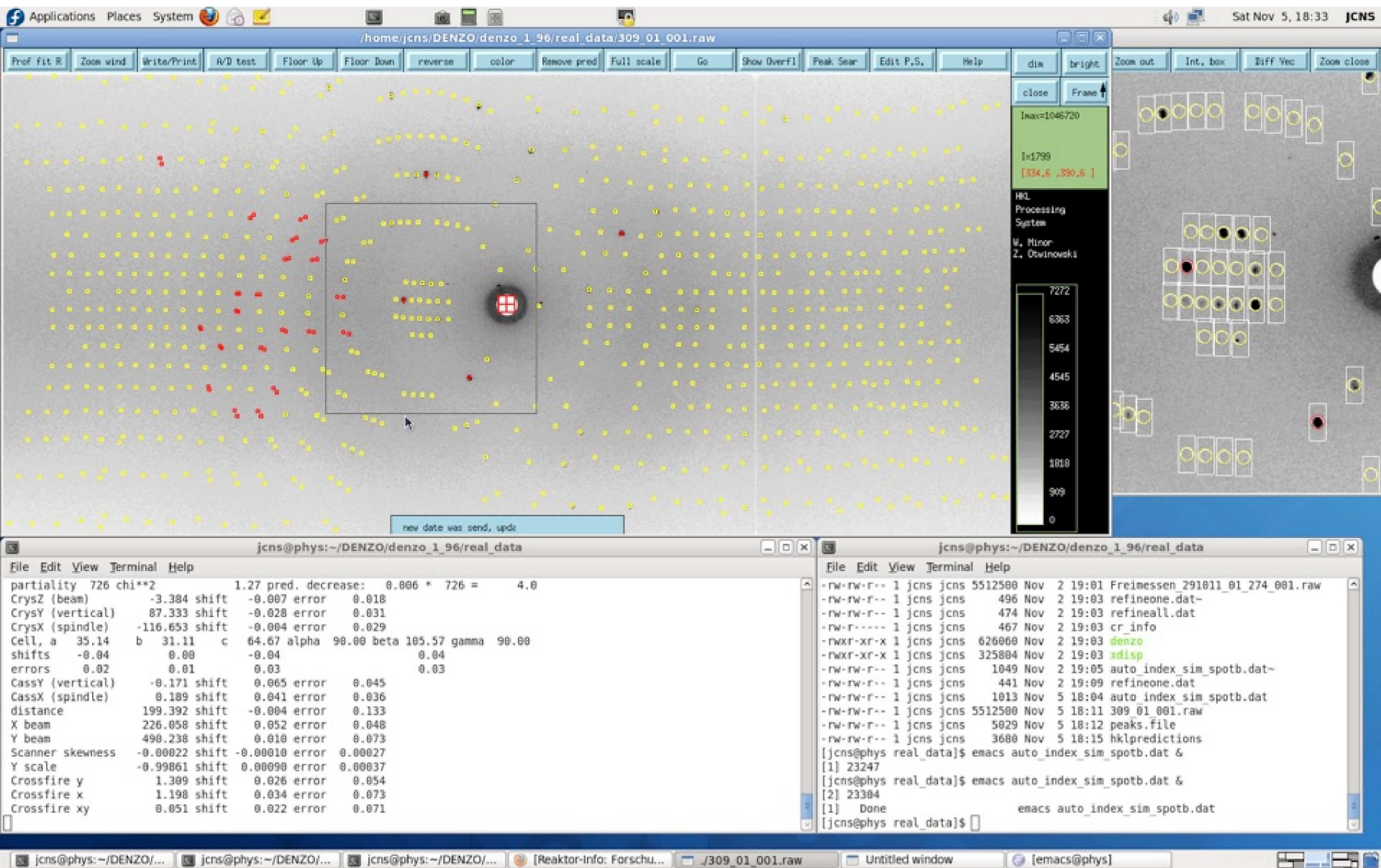
```

jcns@phys:~/DENZO/... jcns@phys:~/DENZO/... jcns@phys:~/DENZO/... [Reaktor-Info: Forschu... ./309_01_001.raw Untitled window

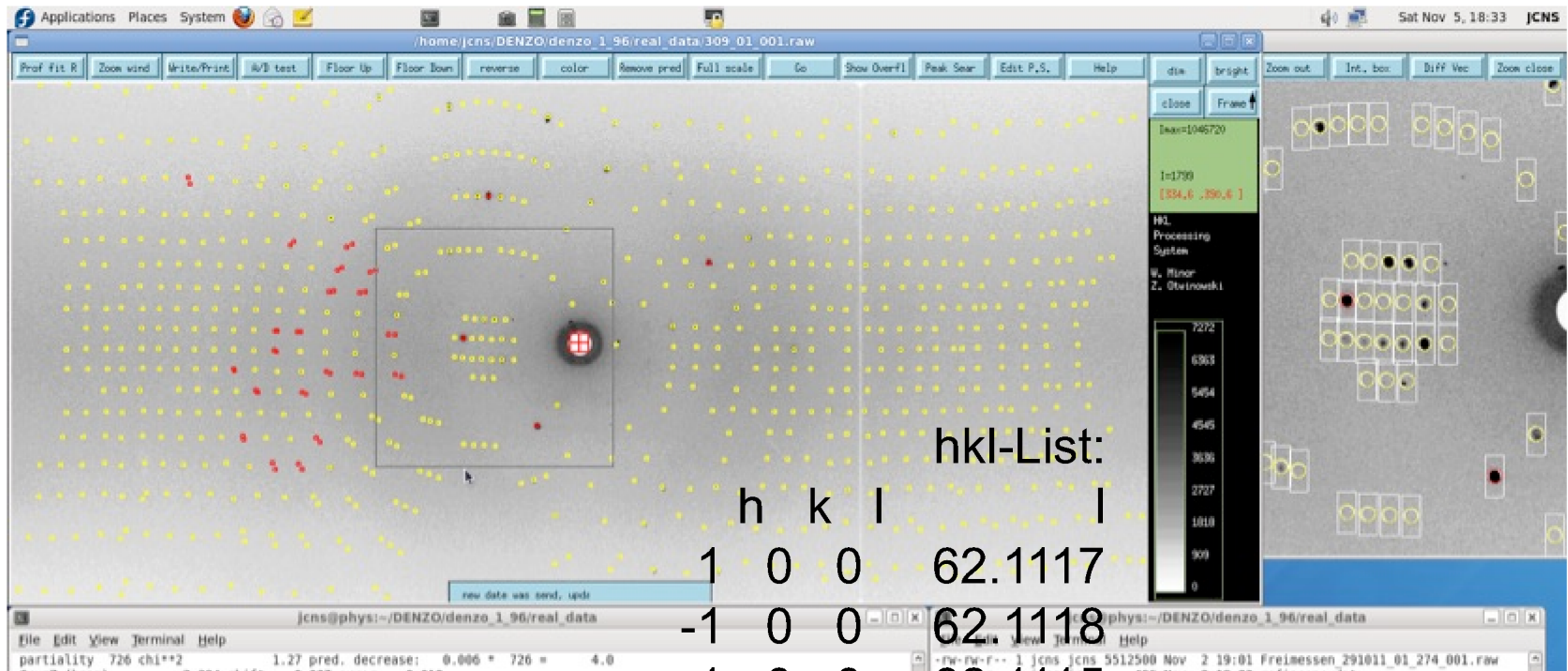
d_min=2.5 Å



d_min=1.5 Å



Integration of partial Bragg peaks with the commercial software hkl-denzo up to $d_{\min}=1.5 \text{ \AA}$



The „old“ HKL2000 software

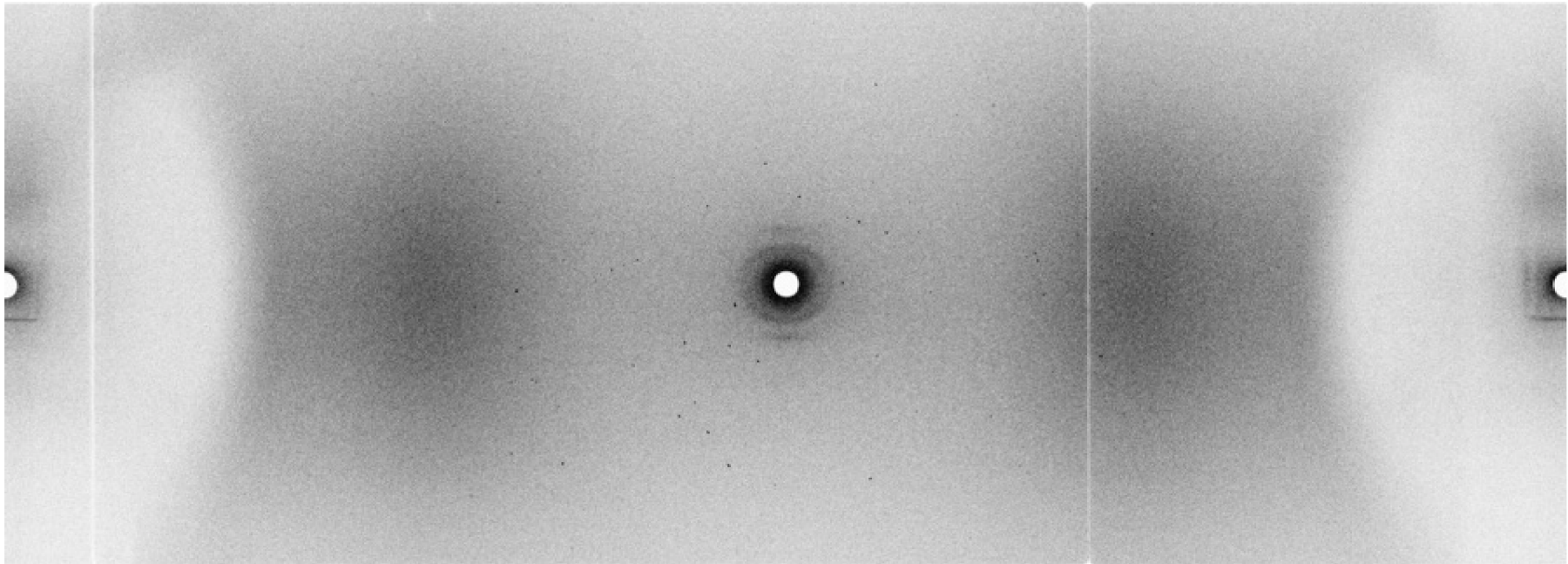
- Not supported,
- Running under academic licence
- Need to re-new every 6 months
- crashes frequently, if you press the buttons in the wrong order (also reboots the computer for you)
- No automatization, arbitrary choice of integration boxes...
- **Need something new!**

Other alternatives

- XDS: no circular detector geometry
- IP mosflm
- DIALS: We are in contact with them...
- Mantid
- Leighton Coates: Recent Publication on 3-D
- ESS: Plans from Esko Oksanen for NMX
- ...???

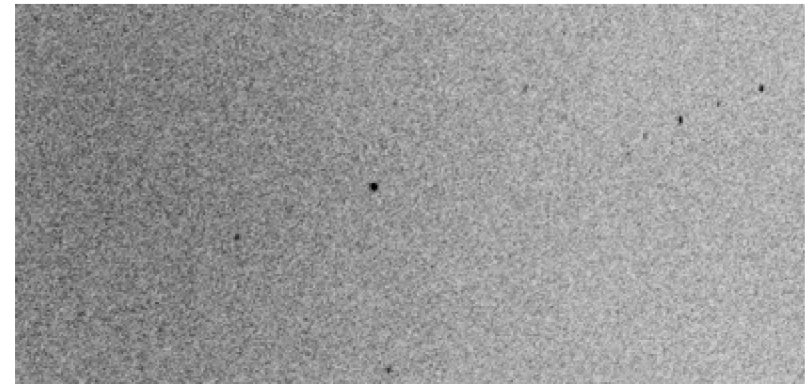
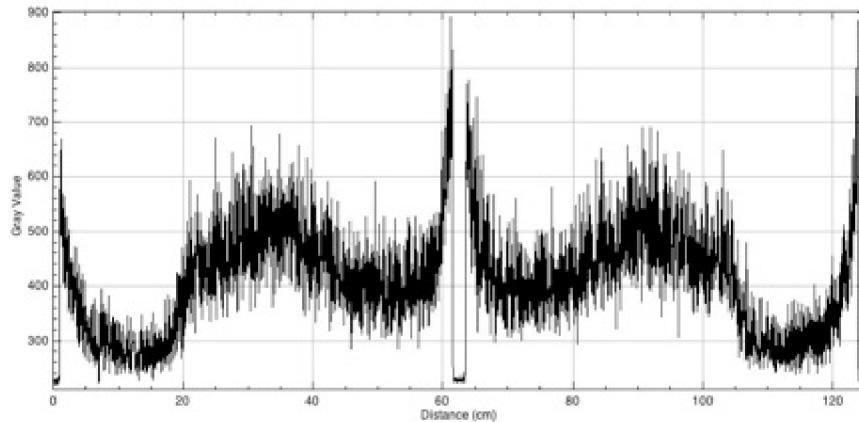
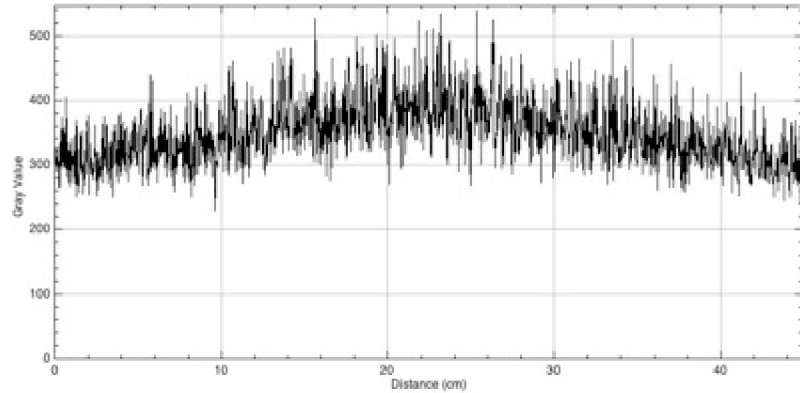
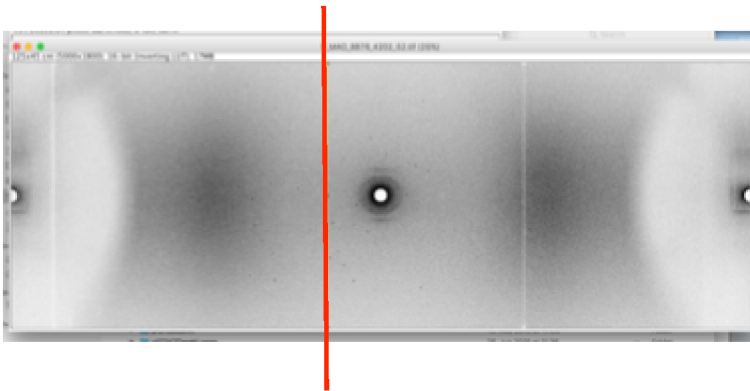
Outline of the problem

- We have to integrate the Bragg spots belonging to one Bragg reflection...



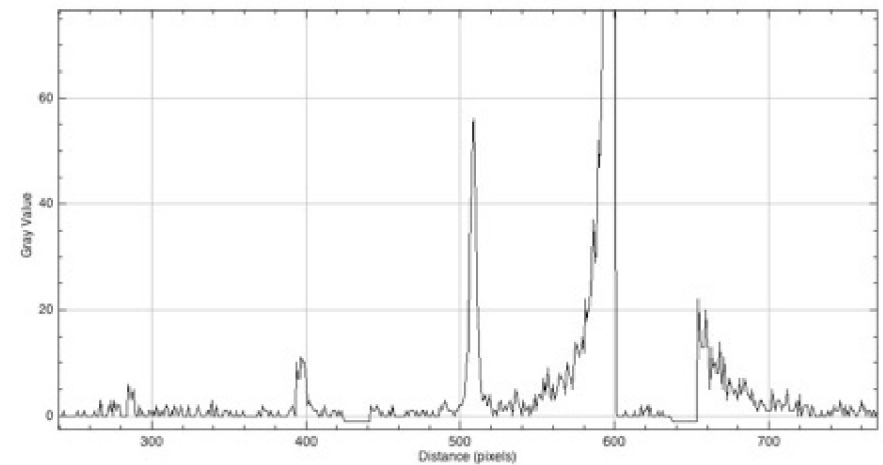
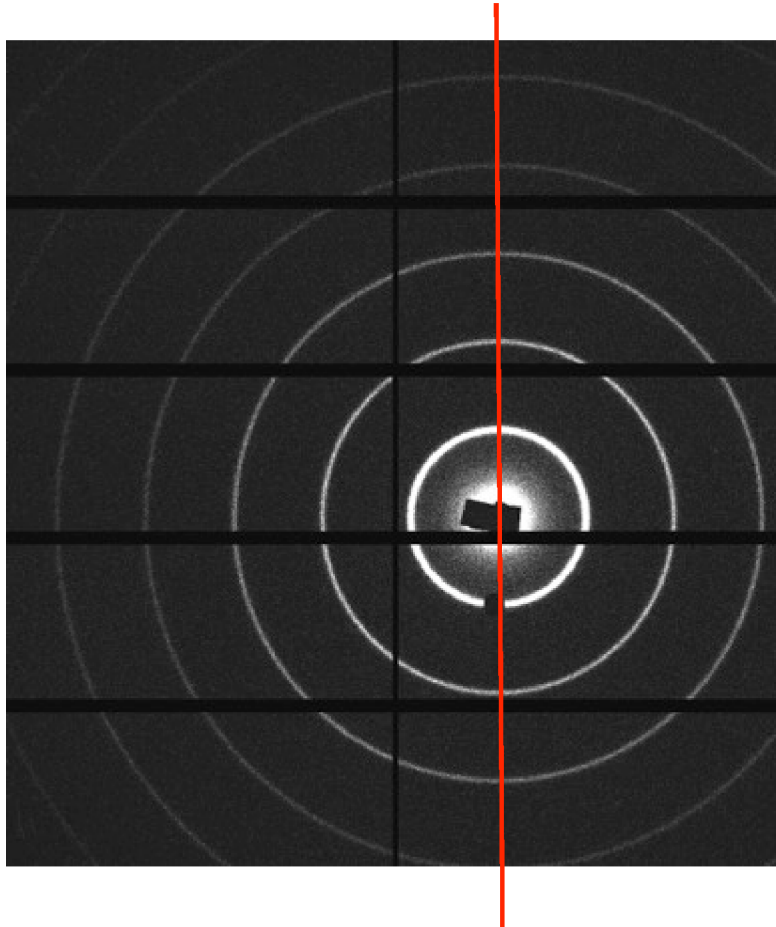
Special issues regarding neutron crystallography as compared to x-ray

- Background much more pronounced and not so homogenous

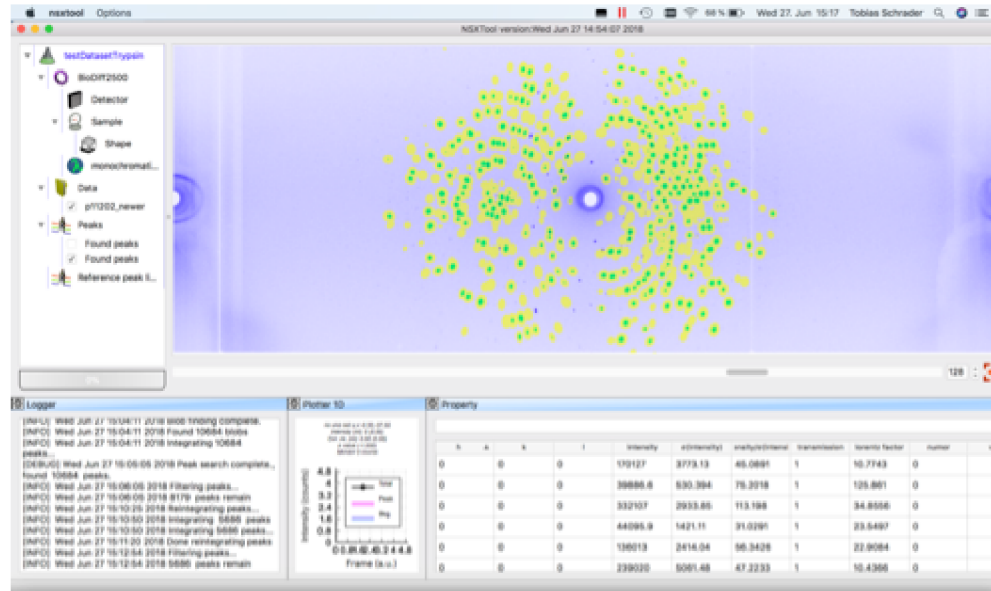


The x-ray case for comparison

- Horizontal cut showing the background



New Data Reduction Software: NXStool



Allows to incorporate

- Absorption correction of crystal with convex hull
- Systematic optimization of integration box size by python scripting
- Pushing resolution limits
- Monitoring detector parameters and detecting detector problems

To be completed...

- Incorporation of a strategy software
- open source, well documented

People involved

- Laurent Chapon



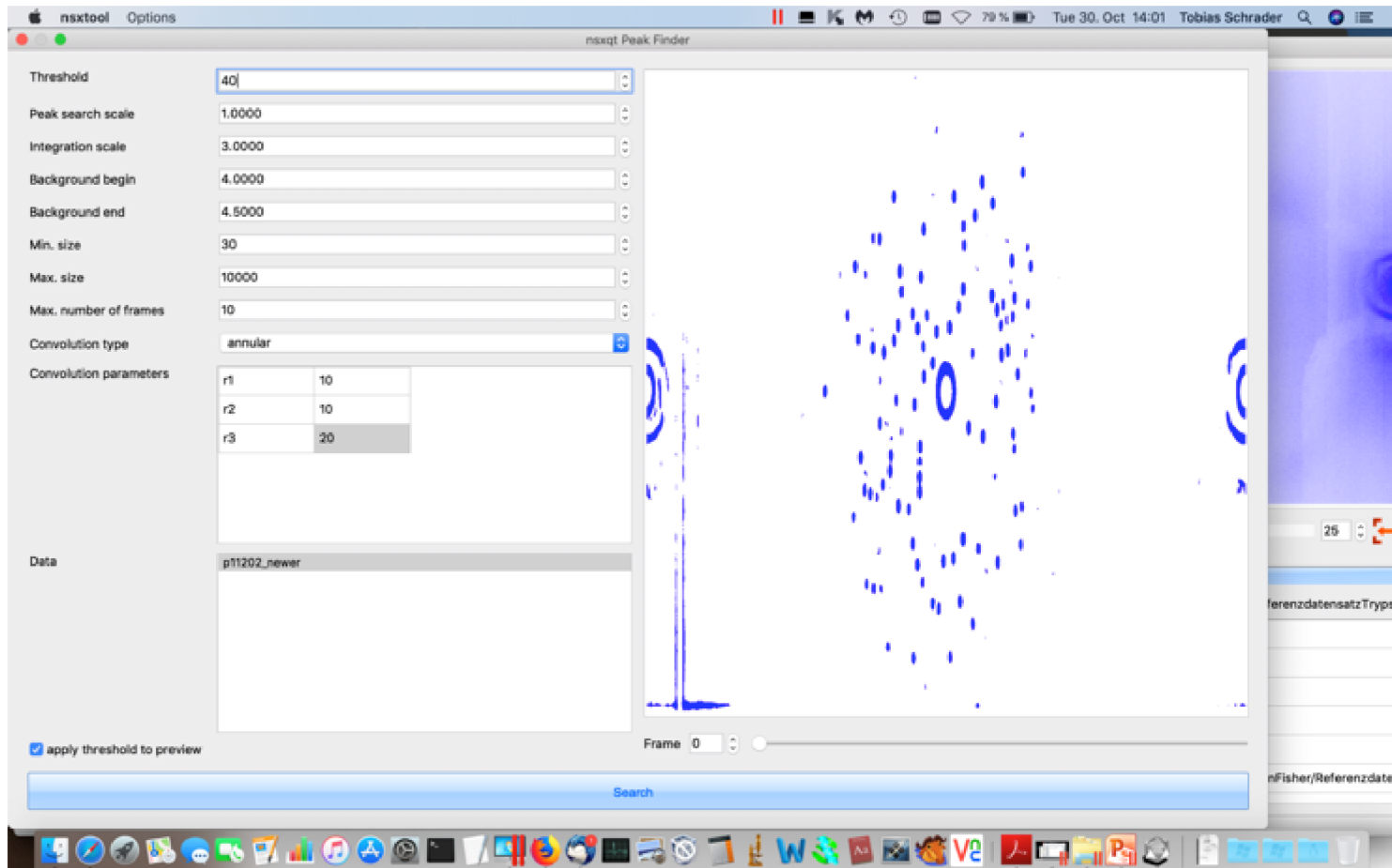
Eric Pellegrini

Jonathan Fisher



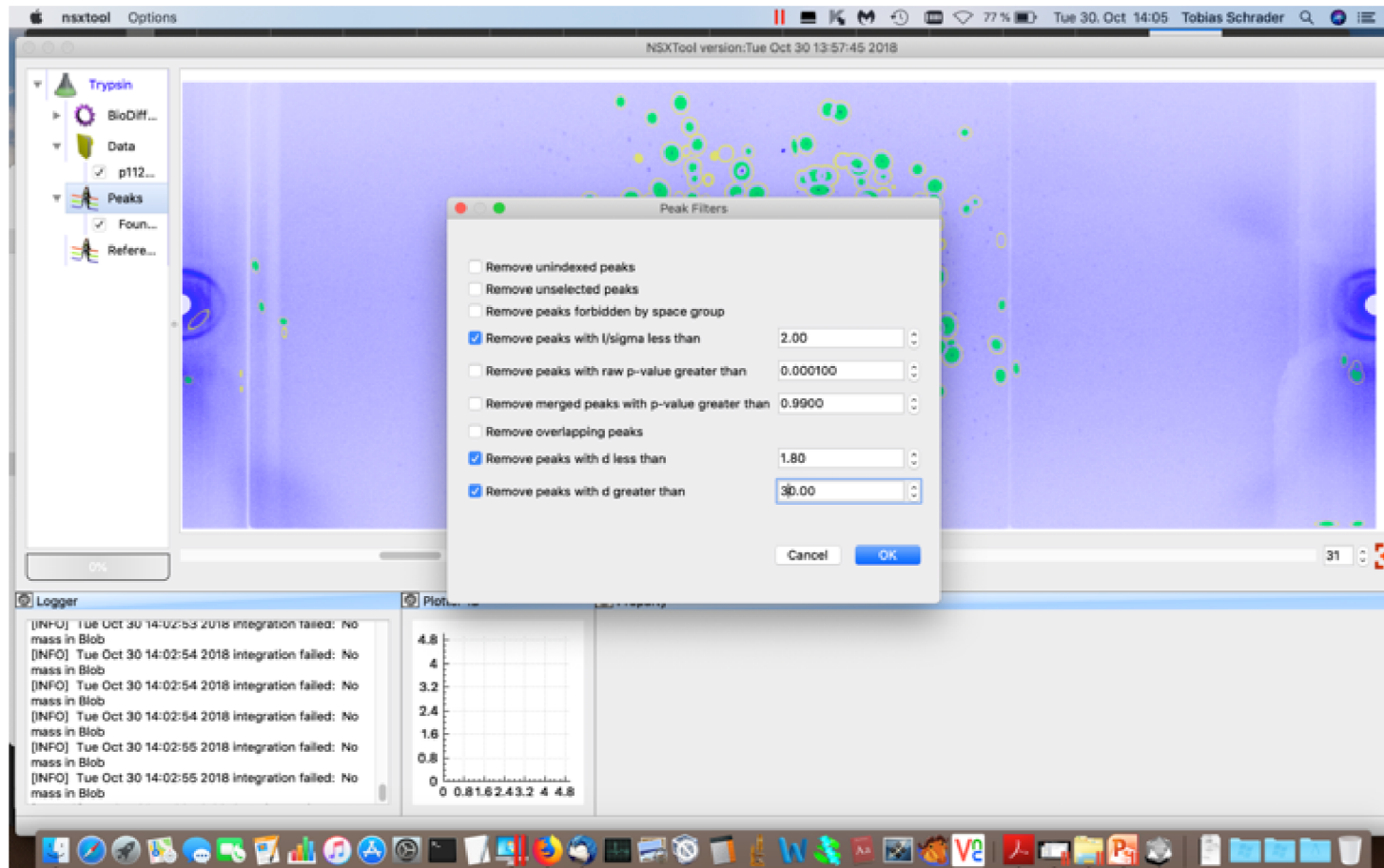
NXStool step by step

- Peak finding based on image analysis



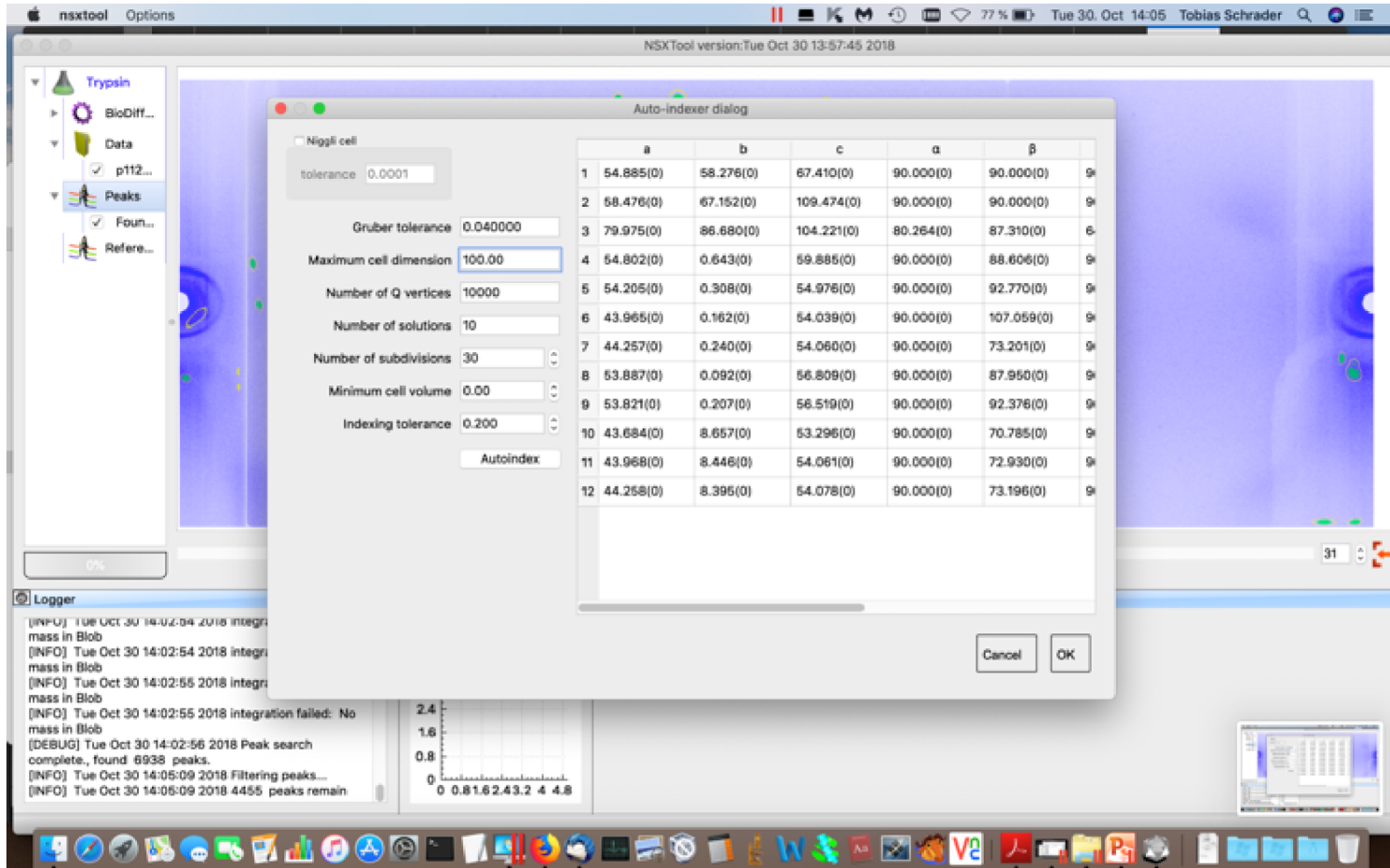
NXStool step by step

- Filtering of the peaks found by the image analysis



NXStool step by step

- Indexing of the Peaks, finding the right unit cell



Auto-indexer dialog

☐ Niggli cell

tolerance 0.0001

Gruber tolerance 0.040000

Maximum cell dimension 100.00

Number of Q vertices 10000

Number of solutions 10

Number of subdivisions 30

Minimum cell volume 0.00

Indexing tolerance 0.200

Autoindex

	a	b	c	α	β
1	54.885(0)	58.276(0)	67.410(0)	90.000(0)	90.000(0)
2	58.476(0)	67.152(0)	109.474(0)	90.000(0)	90.000(0)
3	79.975(0)	86.680(0)	104.221(0)	80.264(0)	87.310(0)
4	54.802(0)	0.643(0)	59.885(0)	90.000(0)	88.606(0)
5	54.205(0)	0.308(0)	54.976(0)	90.000(0)	92.770(0)
6	43.965(0)	0.162(0)	54.039(0)	90.000(0)	107.059(0)
7	44.257(0)	0.240(0)	54.060(0)	90.000(0)	73.201(0)
8	53.887(0)	0.092(0)	56.809(0)	90.000(0)	87.950(0)
9	53.821(0)	0.207(0)	56.519(0)	90.000(0)	92.376(0)
10	43.684(0)	8.657(0)	53.296(0)	90.000(0)	70.785(0)
11	43.968(0)	8.446(0)	54.061(0)	90.000(0)	72.930(0)
12	44.258(0)	8.395(0)	54.078(0)	90.000(0)	73.196(0)

Cancel OK

Logger

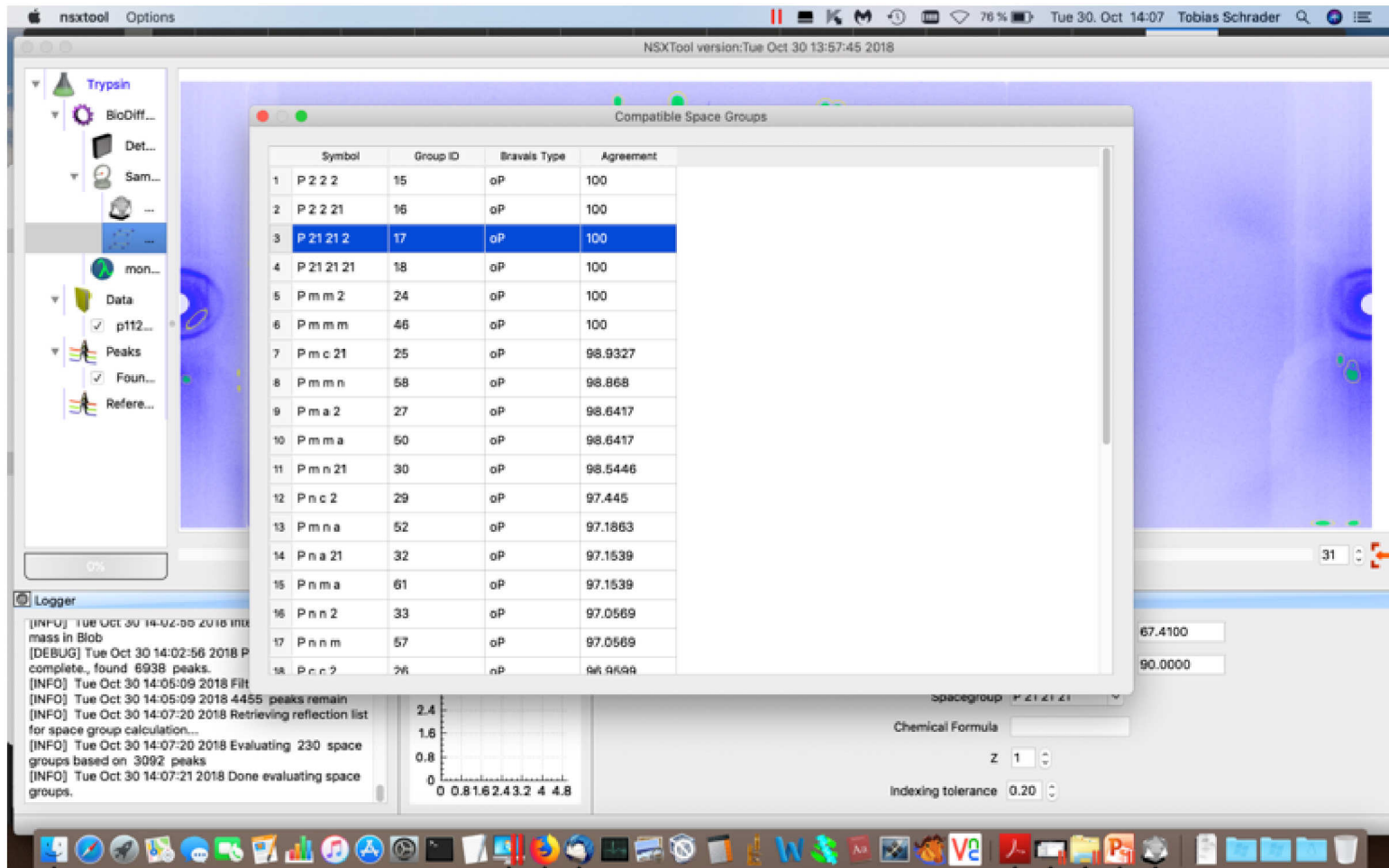
```

[INFO] Tue Oct 30 14:02:04 2018 integr
mass in Blob
[INFO] Tue Oct 30 14:02:54 2018 integr
mass in Blob
[INFO] Tue Oct 30 14:02:55 2018 integr
mass in Blob
[INFO] Tue Oct 30 14:02:55 2018 integrat
mass in Blob
[INFO] Tue Oct 30 14:02:56 2018 integration failed: No
mass in Blob
[DEBUG] Tue Oct 30 14:02:56 2018 Peak search
complete., found 6938 peaks.
[INFO] Tue Oct 30 14:05:09 2018 Filtering peaks...
[INFO] Tue Oct 30 14:05:09 2018 4455 peaks remain
  
```

0% 31

NXStool step by step

- Selecting the space group



NSXTool version: Tue Oct 30 13:57:45 2018

Compatible Space Groups

	Symbol	Group ID	Bravais Type	Agreement
1	P 2 2 2	15	oP	100
2	P 2 2 21	16	oP	100
3	P 21 21 2	17	oP	100
4	P 21 21 21	18	oP	100
5	P m m 2	24	oP	100
6	P m m m	46	oP	100
7	P m c 21	25	oP	98.9327
8	P m m n	58	oP	98.868
9	P m a 2	27	oP	98.6417
10	P m m a	50	oP	98.6417
11	P m n 21	30	oP	98.5446
12	P n c 2	29	oP	97.445
13	P m n a	52	oP	97.1863
14	P n a 21	32	oP	97.1539
15	P n m a	61	oP	97.1539
16	P n n 2	33	oP	97.0569
17	P n n m	57	oP	97.0569
18	P c c 2	26	oP	96.9699

Logger

[INFO] Tue Oct 30 14:02:30 2018 Indexing mass in Blob
 [DEBUG] Tue Oct 30 14:02:56 2018 P complete., found 6938 peaks.
 [INFO] Tue Oct 30 14:05:09 2018 Filtered 4455 peaks remain
 [INFO] Tue Oct 30 14:07:20 2018 Retrieving reflection list for space group calculation...
 [INFO] Tue Oct 30 14:07:20 2018 Evaluating 230 space groups based on 3092 peaks
 [INFO] Tue Oct 30 14:07:21 2018 Done evaluating space groups.

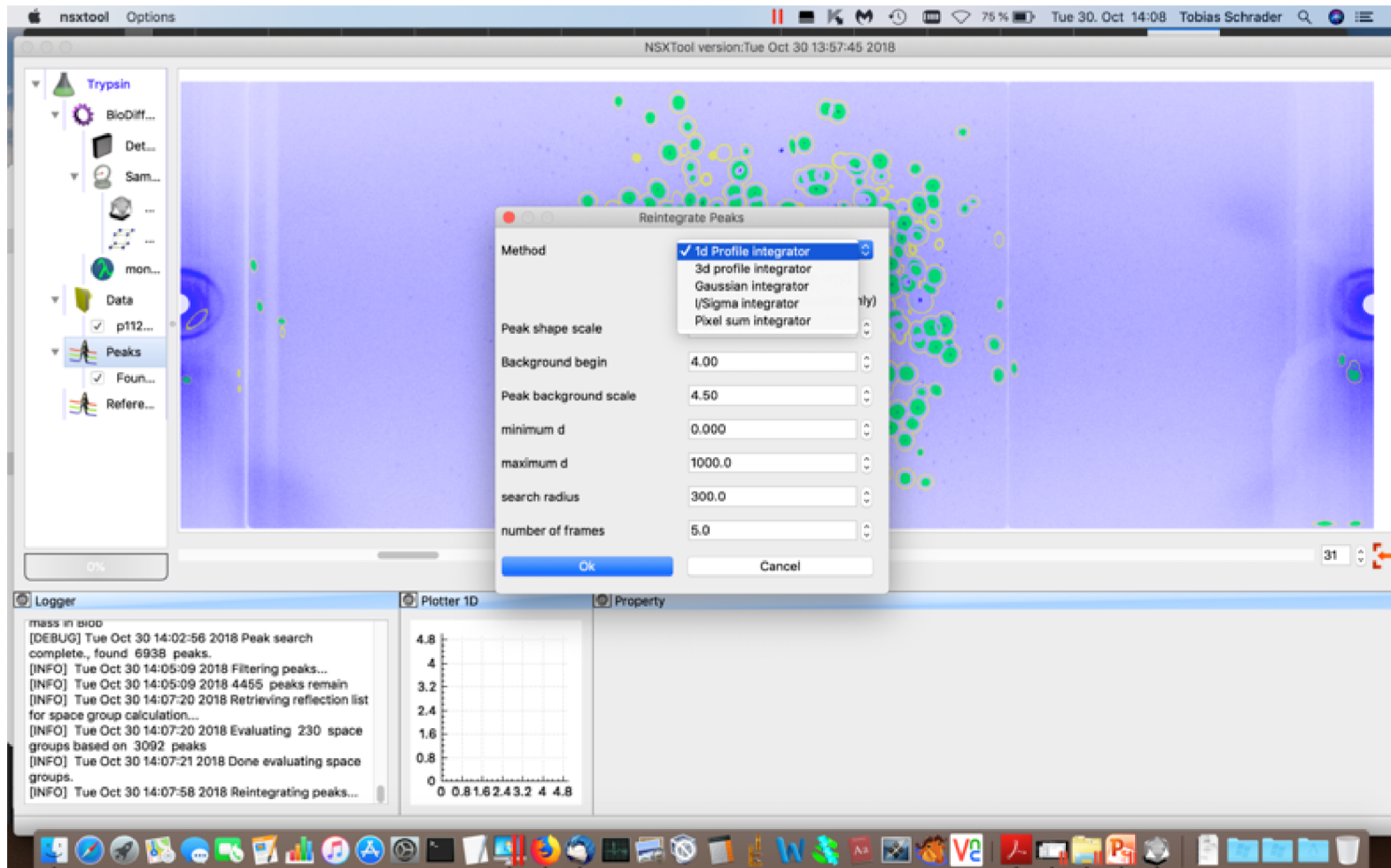
Chemical Formula

Z 1

Indexing tolerance 0.20

NXStool step by step

- Selecting the integration method

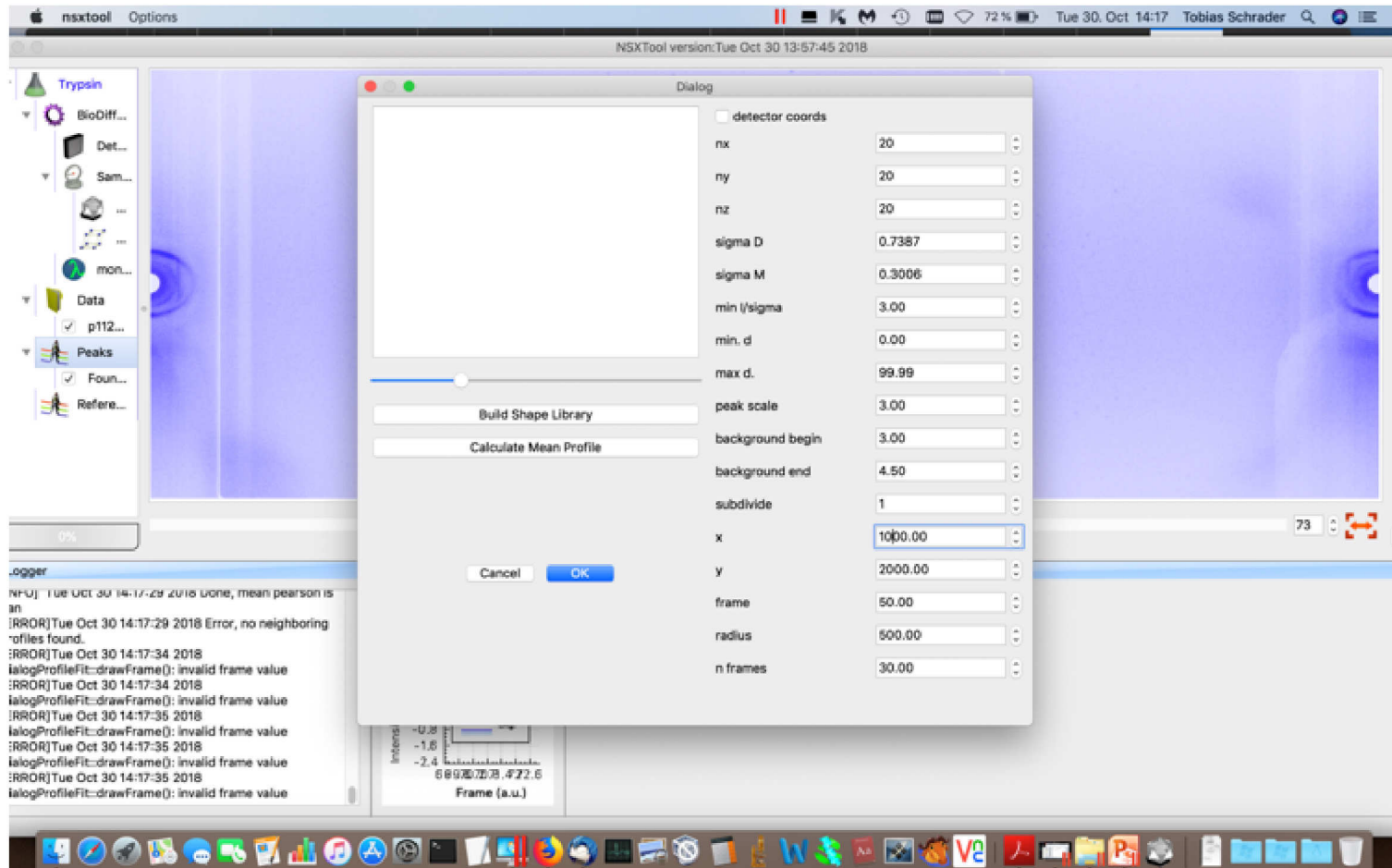


- Refinement of beam position and direction and other instrumental parameters



NXStool step by step

- Building a shape library for profile fitting



- Result of the integration

dmax	dmin	nobs	nmerged	redundancy	Rmeas	Rmeas(est.)	CChalf
19.78	4.19	1612	1153	1.398	0.166	0.016	0.853
4.19	3.33	1187	859	1.382	0.167	0.019	0.837
3.33	2.91	630	484	1.302	0.146	0.022	0.855
2.91	2.64	203	158	1.285	0.177	0.028	0.787
2.64	2.45	100	81	1.235	0.169	0.029	0.818
2.45	2.31	68	58	1.172	0.382	0.056	0.509
2.31	2.19	41	33	1.242	0.104	0.035	0.889
2.19	2.10	18	13	1.385	0.135	0.033	0.866
2.10	2.02	6	5	1.200	0.899	0.054	nan
2.02	1.95	5	4	1.250	0.230	0.040	nan
19.78	1.95	3871	2845	1.361	0.165	0.019	0.842

- Result of HKL2000, the old Software in use:

Shell limit	Lower Angstrom	Upper I	Average error	Average stat.	Norm. Chi**2	Linear R-fac	Square R-fac	Rmeas	Rpim	CC1/2	CC*	
50.00	3.23	7052.9	216.6	166.1	1.593	0.034	0.117	0.043	0.026	0.942	0.985	
	3.23	2.56	3943.3	243.3	226.3	1.403	0.061	0.058	0.077	0.046	0.991	0.998
	2.56	2.24	2919.6	321.9	315.4	1.505	0.099	0.101	0.128	0.080	0.968	0.992
	2.24	2.04	2634.7	384.7	380.1	1.418	0.123	0.123	0.161	0.103	0.950	0.987
	2.04	1.89	2324.2	416.1	412.6	1.160	0.141	0.132	0.188	0.123	0.946	0.986
	1.89	1.78	1894.2	422.0	419.7	1.211	0.173	0.167	0.235	0.157	0.905	0.975
	1.78	1.69	1596.5	407.5	405.8	1.154	0.194	0.185	0.261	0.173	0.894	0.972
	1.69	1.62	1284.7	376.6	375.4	1.211	0.222	0.211	0.294	0.191	0.871	0.965
	1.62	1.55	1003.5	335.2	334.4	1.187	0.263	0.253	0.348	0.225	0.814	0.947
	1.55	1.50	732.3	289.9	289.4	1.203	0.314	0.316	0.417	0.272	0.714	0.913
All reflections			2847.7	332.0	320.6	1.368	0.085	0.114	0.111	0.070		

Current status

- Programme runs on our Mac-Computers
- Support available from Eric Pellegrini at ILL
- Python scripting available but not rolled out to Andreas and me
- R-factors are still a bit worse than in the old denzo/HKL2000 case
- New programmer is needed to remove all remaining bugs and to do some more testing...
- **Close contact and many discussions with this programmer is needed**
- Ideas: To use McStas Calculations for improving the prediction of the centre of the predicted reflections
- Better absorption correction by using the convex hull of the crystal in correcting the Bragg intensities

Thanks

- Jonathan Fisher
- Andreas Ostermann
- Eric Pellegrini
- Laurent Chapon
- Joachim Wuttke

- and you for the attention to my talk!!