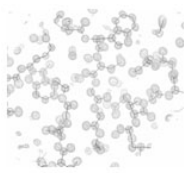
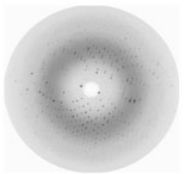
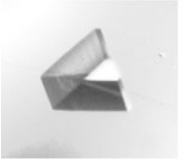


Data Processing

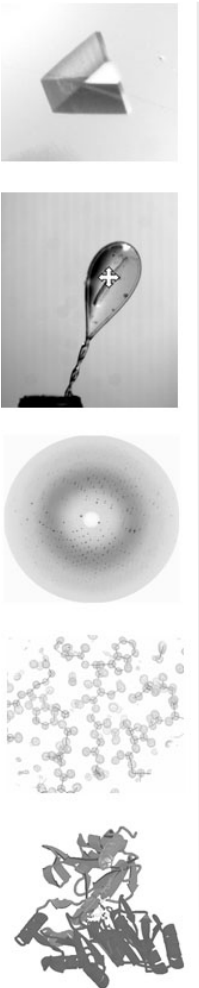
XDSAPP and future developments

Karine Röwer

*Helmholtz-Zentrum Berlin für Materialien und Energie
Macromolecular Crystallography (NP-GMX)
Albert-Einstein-Str. 15
D-12489 Berlin, Germany
karine.sparta@helmholtz-berlin.de*



XDSAPP layout



Summary | Integrate plots | CORRECT plots | XDSSTAT plots | Settings | Logfiles and pictures | Running output

XDSAPP 2.0
Based on python and Qt
xdsapp@helmholtz-berlin.de

Macromolecular Crystallography group
Helmholtz Zentrum Berlin
<http://www.helmholtz-berlin.de/bessy-mx>

Reference:
K. Sparta, M. Krug, U. Heinemann, U. Mueller, M. S. Weiss (2016).
J. Appl. Cryst. **49**, 1085-1092.

Did you know?
When loading a data set that has already been processed, this tab will show the last processing summary.

Control panel

Summary

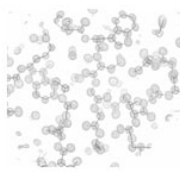
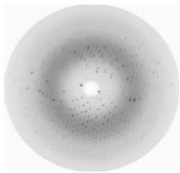
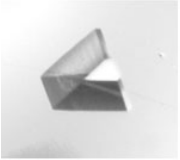
Detailed summary

Status line

No software, however sophisticated, can make up for badly planned experiments. -- Manfred S. Weiss

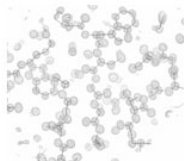
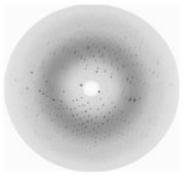
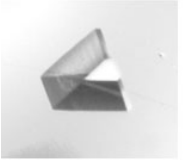
K. Sparta *et al.* (2016). *J. Appl. Cryst.* **49**, 1085-1092.

XDSAPP features



- Processing of diffraction data using XDS, CCP4 and Phenix
- Live graphical representation of data statistics
- Automated decision-making
 - Space group determination
 - Detection of anomalous signal
 - Resolution cutoff
- Conversion of intensities to various formats: mtz, hkl, cns
- Live mode for processing during data collection

XDSAPP workflow



Index

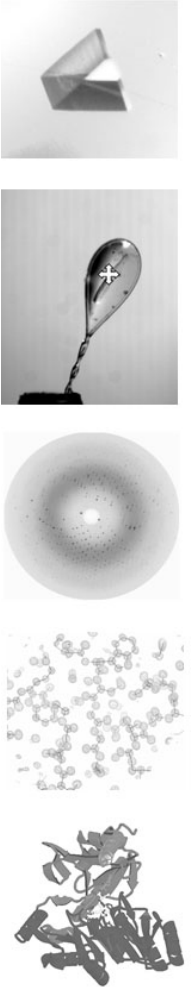
- XYCORR
INIT
COLSPOT
- IDXREF
- DEFPIX

Integrate + CORRECT

- INTEGRATE in P1
CORRECT
- Determination of
 - maximum resolution
 - anomalous signal
 - point group and Bravais lattice
- Smart reintegration cycles
- Check aliens, ice rings
- Space group determination
 - pointless
 - CORRECT

Analyze

- XDSSTAT
- Data conversion
 - MTZ
 - Shelx
 - CNS format
- Twinning analysis
 - SFCHECK
 - phenix.xtriage
- Enantiomorphic space groups
- Quality analysis



Select

Load

Index

Run DEFPPIX

Integrate + Correct

Rerun Correct

Analyse

Do all

In brief

[Prol_wt_Mg_LP_X9_3m15s_1](#)

Images:
1 to 1600

Wavelength:
0.918409 Å

Spots indexed:
5065 of 7410 (68.4%)

Finished



Summary Integrate plots CORRECT plots XDSSTAT plots Settings Logfiles and pictures Running output

Indexing results for data set Prol_wt_Mg_LP_X9_3m15s_1

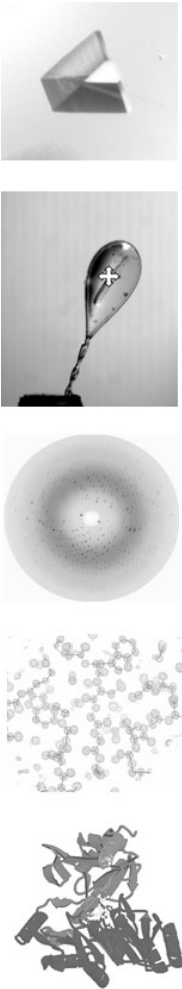
No. of subtrees: 15
No. of strong spots in subtree 1: 2982
No. of strong spots in subtree 2: 2
Quality value: 1.1
Delta: 0.1
DH, DK, DL: 0.01, 0.01, 0.05
Selected origin: 0 0 0
Computed (given) beam position on detector: 1240.4, 1250.2 (1239.0, 1248.0)
Std. of spot position (refined): 0.30 (0.35) px
Std. of spindle position (refined): 0.11 (0.14) deg.
Detector distance given (refined): 283.565 (282.39)
Spots indexed: 5065 out of 7410 (68.4 %)

Possible lattices are:

aP	0.0	74.4	74.4	216.9	90.0	90.0	88.2
aP	0.2	74.4	74.4	216.9	90.0	90.0	91.8
mP	1.3	74.4	216.9	74.4	90.0	91.8	90.0
mC	1.3	106.8	103.5	216.9	90.0	90.0	90.0
oC	2.0	103.5	106.8	216.9	90.0	90.0	90.0
mC	2.0	103.5	106.8	216.9	90.0	90.0	90.0
mP	16.0	74.4	74.4	216.9	90.0	90.0	91.8
mP	16.7	74.4	74.4	216.9	90.0	90.0	91.8
oP	16.9	74.4	74.4	216.9	90.0	90.0	91.8
tP	17.6	74.4	74.4	216.9	90.0	90.0	91.8

Idle since 15:51:44.

Reintegration cycles



Select

In brief

ProL_wt_Mg_LP_X9_3m15s_

Space group:
C222

Resolution:
1.15Å

Rmeas total:
9.5 %

Mosaicity:
0.082°

Completeness:
78.5 %

I/sigI:
7.87

I_{sa}:
55.57

Reprocessing



Summary

Integrate plots

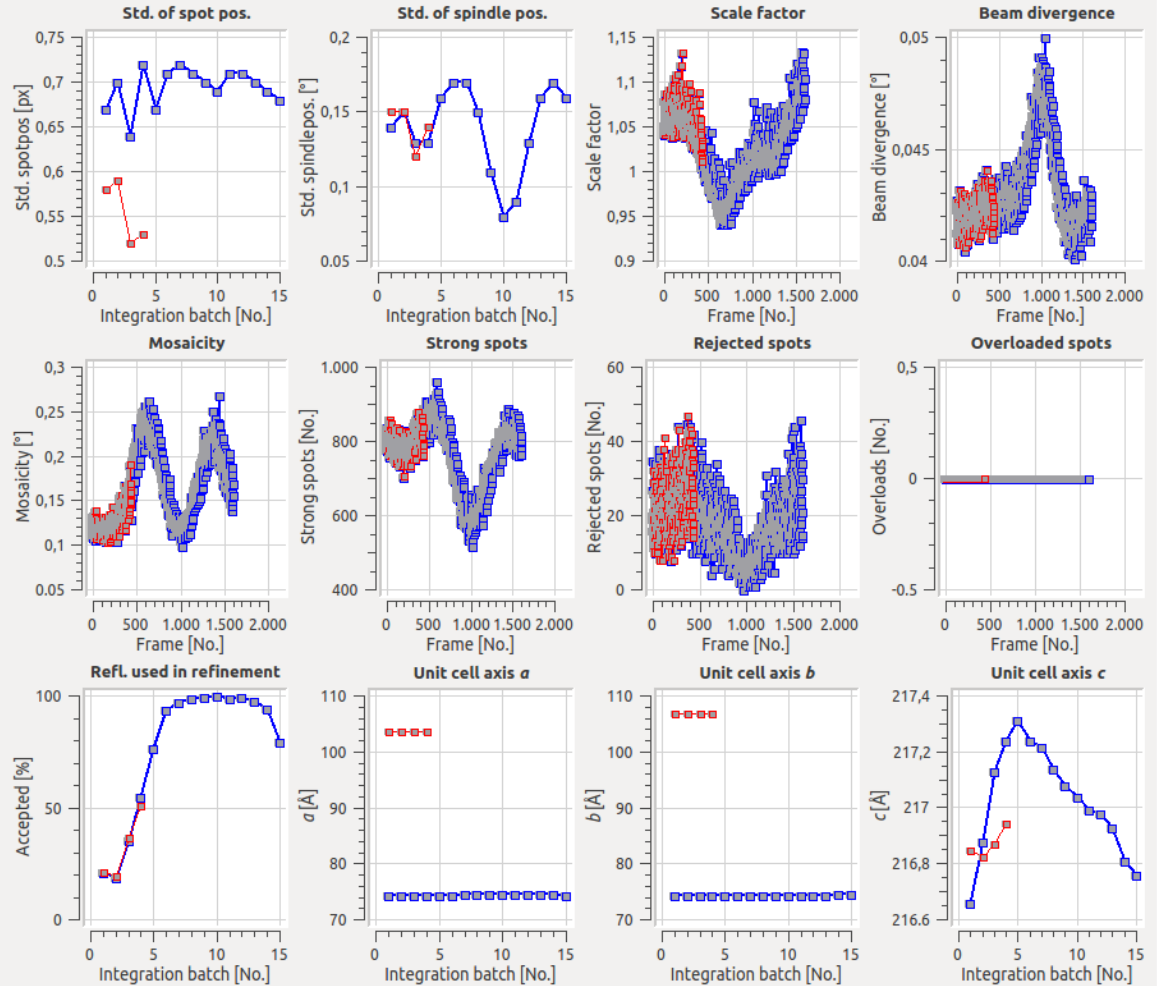
CORRECT plots

XDSSTAT plots

Settings

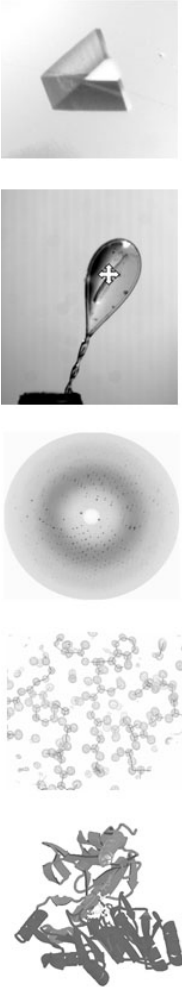
Logfiles and pictures

Running output



Processing of images 429 - 534, data set ProL_wt_Mg_LP_X9_3m15s_1, reintegration step 1 of max. 3

Data quality indicators (CORRECT)



Select

In brief

ProL_wt_Mg_LP_X9_3m15s_

Space group:
C222

Resolution:
1.15Å

Rmeas total:
9.5 %

Mosaicity:
0.082°

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78.5 %

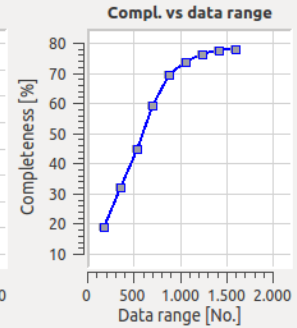
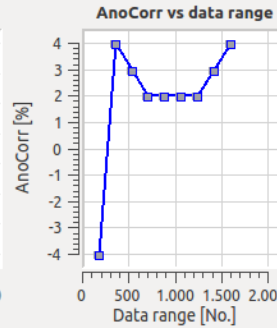
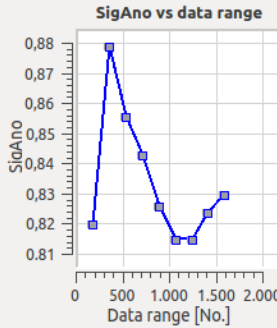
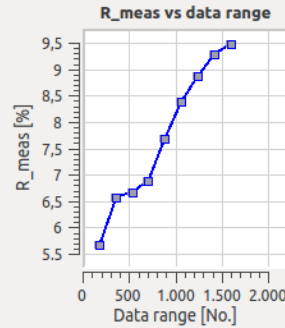
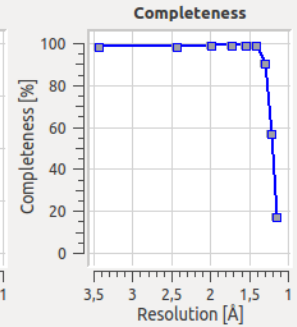
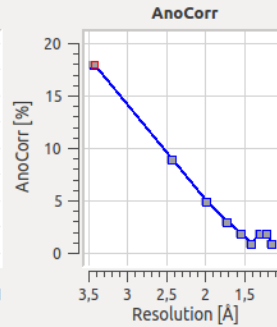
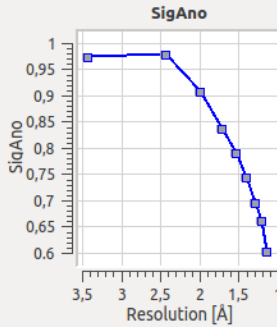
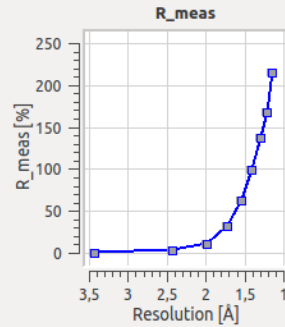
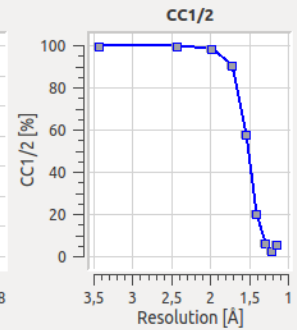
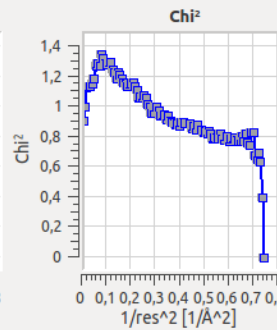
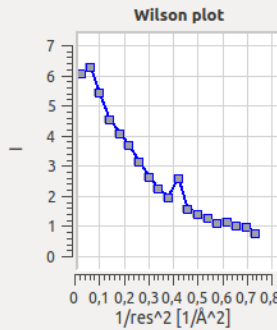
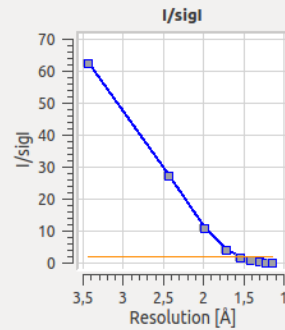
I/sigl:
7.87

Iσa:
55.57

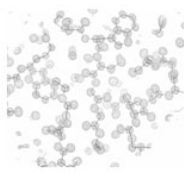
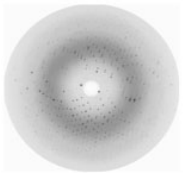
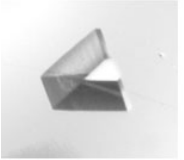
Reprocessing



Summary Integrate plots CORRECT plots XDSSTAT plots Settings Logfiles and pictures Running output

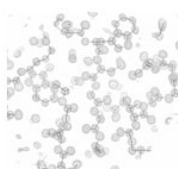
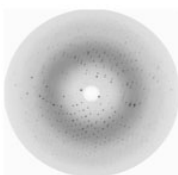
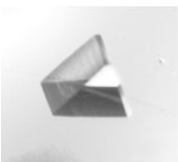


CORRECT Data set ProL_wt_Mg_LP_X9_3m15s_1



To be performed after the last CORRECT run

- XDSSTAT
- XDSCONV
- SFCHECK
- Quality analysis
 - Indexing quality
 - Twinning (POINTLESS, PHENIX.XTRIAGE)
 - Pseudo translation
 - Ice rings
- Things you may want to check



Select

In brief

[Prol_wt_Mg_LP_X9_3m15s_1](#)

Space group:
C222(1)

Resolution:
1.46Å

Rmeas total:
7.0 %

Mosaicity:
0.065°

Completeness:
99.8 %

I/sigI:
17.29

ISA:
50.99

Finished



Summary Integrate plots CORRECT plots XDSSTAT plots Settings Logfiles and pictures Running output

Processing summary for data set Prol_wt_Mg_LP_X9_3m15s_1

Results in: /home/karine/Prolidase/xdsapp/Prol_wt_Mg_LP_X9_3m15s_1

Things you may want to check:
Twinning

Table 1 like summary

Space group	C222(1) (20)
Unit cell parameters [Å]	103.61 106.84 217.00 90.0 90.0 90.0
Resolution limit [Å]	47.93-1.46 (1.55-1.46)
No. of reflections	1227068
No. of uniques	205595
Multiplicity	5.97
I/sigI	17.29 (2.01)
R_meas [%]	7.0 (92.9)
Completeness [%]	99.8 (99.3)
B(Wilson) [Å²]	24.3
Mosaicity [deg]	0.065
CC(1/2)	100.0 (52.8)
ISA	50.99

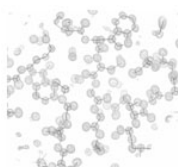
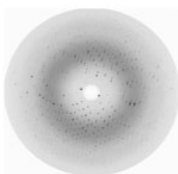
Result files generated for CCP4, CNS, SHELX

Prol_wt_Mg_LP_X9_3m15s_1_l.mtz
 Prol_wt_Mg_LP_X9_3m15s_1_f.mtz
 Prol_wt_Mg_LP_X9_3m15s_1_f_plus_f_minus.mtz
 Prol_wt_Mg_LP_X9_3m15s_1.hkl
 Prol_wt_Mg_LP_X9_3m15s_1.cv

Intensity checks

SFCHECK	
Perfect twinning test	2.0715
Pseudo translation	not detected

Idle since 16:34:33.



Select

Do all

In brief

[Prol_wt_Mg_LP_X9_3m15s...](#)

Space group:
C222(1)

Resolution:
1.46Å

Rmeas total:
7.0 %

Mosaicity:
0.065°

Completeness:
99.8 %

I/sigI:
17.29

Isa:
50.99

Finished



Summary

Intensity checks

SFCHECK	2.0715
Perfect twinning test	not detected
Pseudo translation	not detected
Phenix.xtriage	
<I ² >/<I>	2.033 (untwinned 2.000, perfect twin 1.500)
<F> ² /<>F ² >	0.790 (untwinned 0.785, perfect twin 0.885)
< E ² -1 >	0.740 (untwinned 0.736, perfect twin 0.541)
Pointless	
-test	0.442 (untwinned 0.500, perfect twin 0.375)
L ² -test	0.271 (untwinned 0.333, perfect twin 0.200)

Space group determination by Pointless

Selected space group	C 2 2 21 (20)
Total probability	0.961
Systematic absences probability	0.992
Next probable space group	none (-)
Total probability	-
Systematic absences probability	-

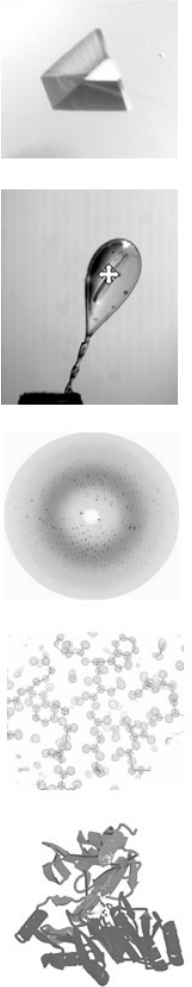
Indexing summary

No. of subtrees	15
No of strong spots in root subtree	2982
No. of strong spots in subtree 2	2
Quality value	1.1
Delta	0.1
DH, DK, DL	0.01, 0.01, 0.05
Selected origin	0 0 0
Computed (given) beam position	1240.4, 1250.2 (1239.0, 1248.0)
Std. of spot position (refined)	0.30 (0.35)
Std. of spindle position (refined)	0.11 (0.14)
Detector distance (refined)	283.565 (282.39)
No. of spots indexed (total)	5065 (7410) (=68.4 %)

Experimental settings

Idle since 16:34:33.

More statistics: XDSSTAT



Select

Load

Index

Run DEFPPIX

Integrate + Correct

Rerun Correct

Analyse

Do all

In brief

ProI_wt_Mg_LP_X9_3m15s_1

Space group:
C222(1)

Resolution:
1.46Å

Rmeas total:
7.0 %

Mosaicity:
0.065°

Completeness:
99.8 %

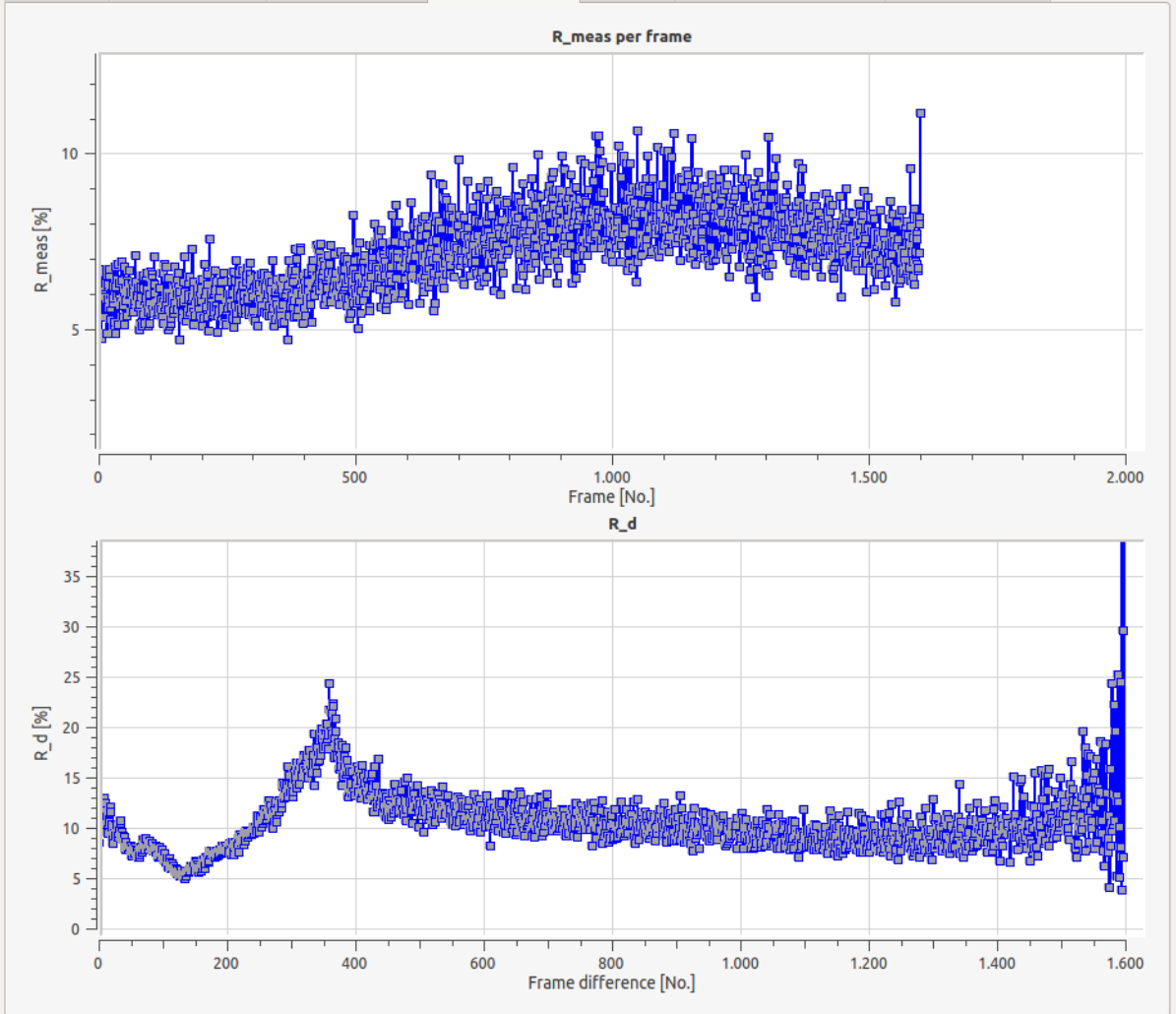
I/sigI:
17.29

I/si:
50.99

Analysing intensities



Summary Integrate plots CORRECT plots XDSSTAT plots Settings Logfiles and pictures Running output

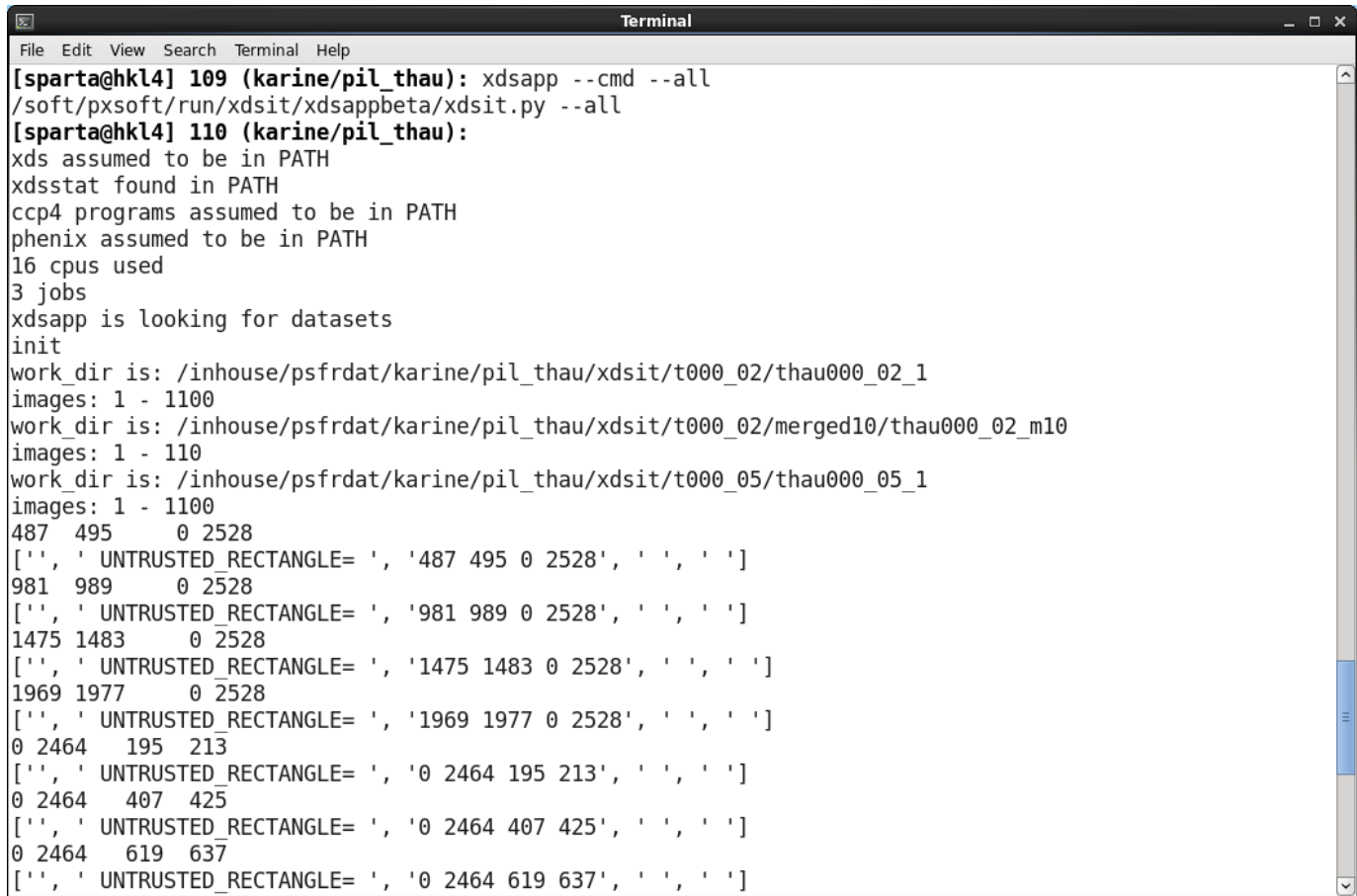


PHENIX.XTRIAE Data set ProI_wt_Mg_LP_X9_3m15s_1

XDSAPP in command line mode

Command line version of XDSAPP at the beamline, with full automatic server and database connection. Possibility to process all data sets at once.

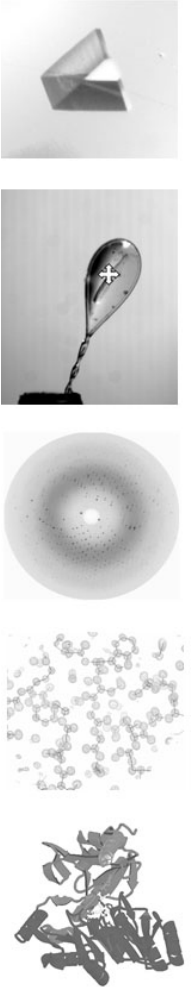
Particularly useful for fragment screening projects.



```

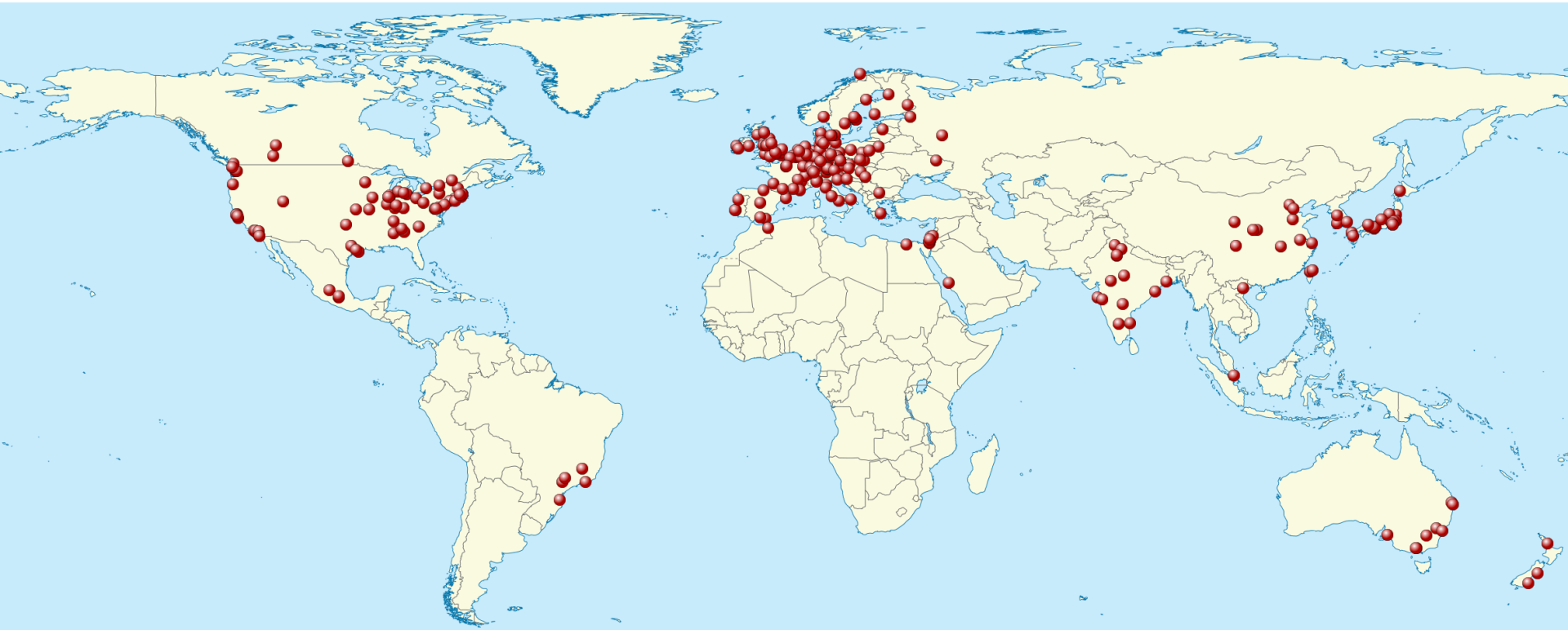
Terminal
File Edit View Search Terminal Help
[sparta@hkl4] 109 (karine/pil_thau): xdsapp --cmd --all
/soft/pxsoft/run/xdsit/xdsappbeta/xdsit.py --all
[sparta@hkl4] 110 (karine/pil_thau):
xds assumed to be in PATH
xdsstat found in PATH
ccp4 programs assumed to be in PATH
phenix assumed to be in PATH
16 cpus used
3 jobs
xdsapp is looking for datasets
init
work_dir is: /inhouse/psfrdat/karine/pil_thau/xdsit/t000_02/thau000_02_1
images: 1 - 1100
work_dir is: /inhouse/psfrdat/karine/pil_thau/xdsit/t000_02/merged10/thau000_02_m10
images: 1 - 110
work_dir is: /inhouse/psfrdat/karine/pil_thau/xdsit/t000_05/thau000_05_1
images: 1 - 1100
487 495 0 2528
['', ' UNTRUSTED_RECTANGLE= ', '487 495 0 2528', ' ', ' ', ' ']
981 989 0 2528
['', ' UNTRUSTED_RECTANGLE= ', '981 989 0 2528', ' ', ' ', ' ']
1475 1483 0 2528
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1969 1977 0 2528
['', ' UNTRUSTED_RECTANGLE= ', '1969 1977 0 2528', ' ', ' ', ' ']
0 2464 195 213
['', ' UNTRUSTED_RECTANGLE= ', '0 2464 195 213', ' ', ' ', ' ']
0 2464 407 425
['', ' UNTRUSTED_RECTANGLE= ', '0 2464 407 425', ' ', ' ', ' ']
0 2464 619 637
['', ' UNTRUSTED_RECTANGLE= ', '0 2464 619 637', ' ', ' ', ' ']

```

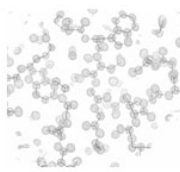
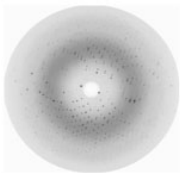
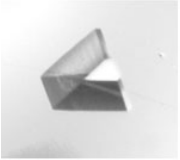


XDSAPP users community

821 downloaders, distributed over 488 institutes and 43 countries.



XDSAPP next features

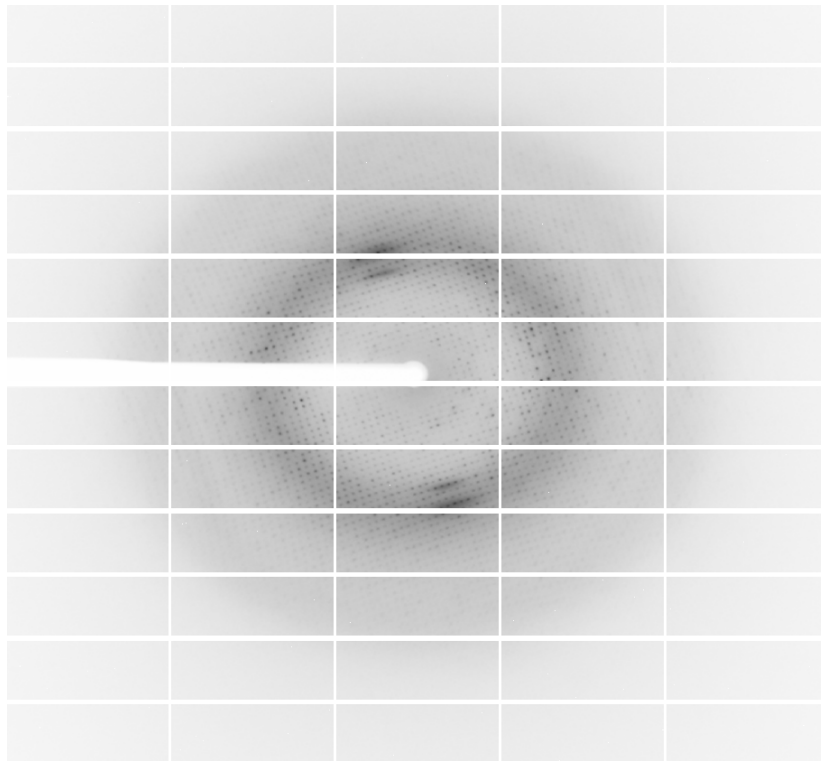
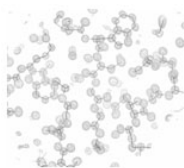
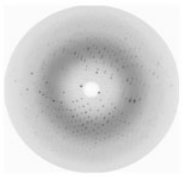
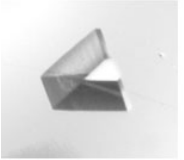


- Improved indexing procedure
- Resolution cutoff based on significant CC1/2
- Handling of Eiger data sets

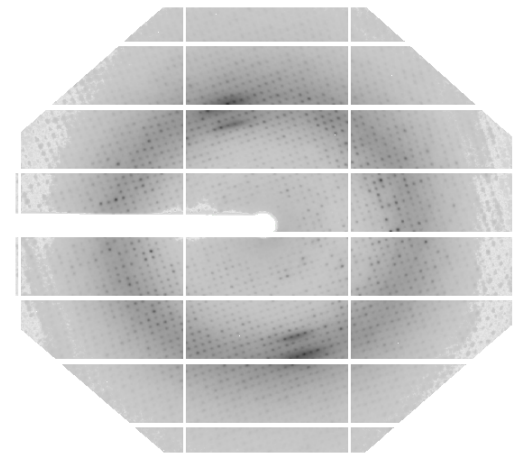
Improved performance

Regions of interest for COLSPOT and IDXREF

Limit resolution in BKGINIT.cbf



No resolution limit

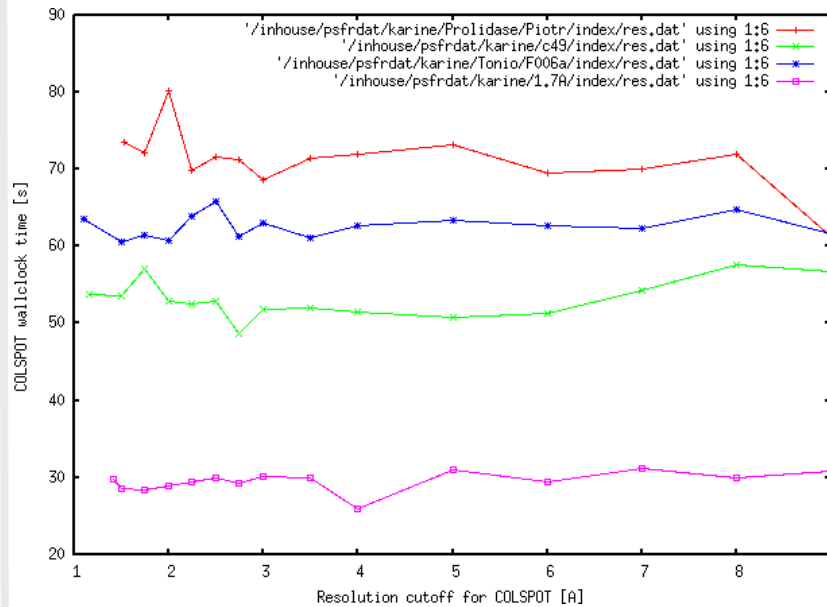


~ 2.5Å maximum resolution

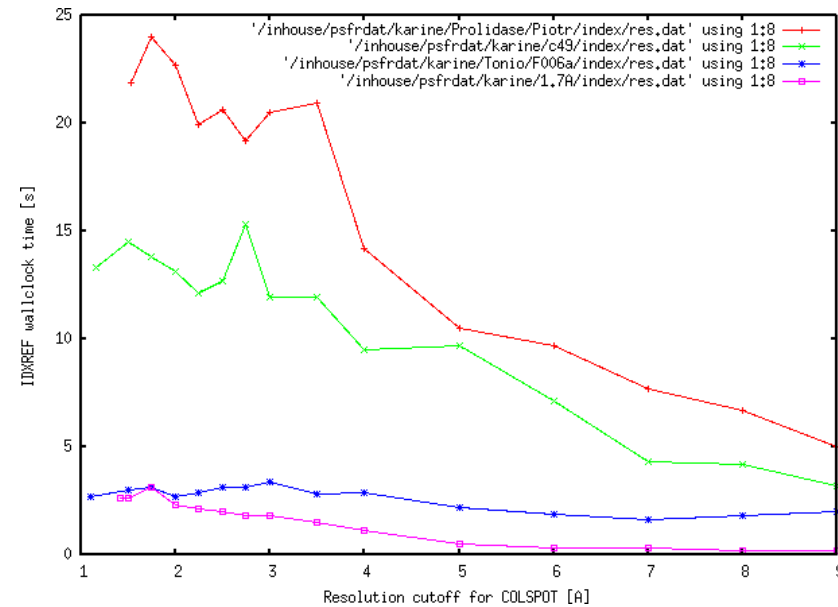
Improved performance

Regions of interest for COLSPOT and IDXREF

Time performance



COLSPOT



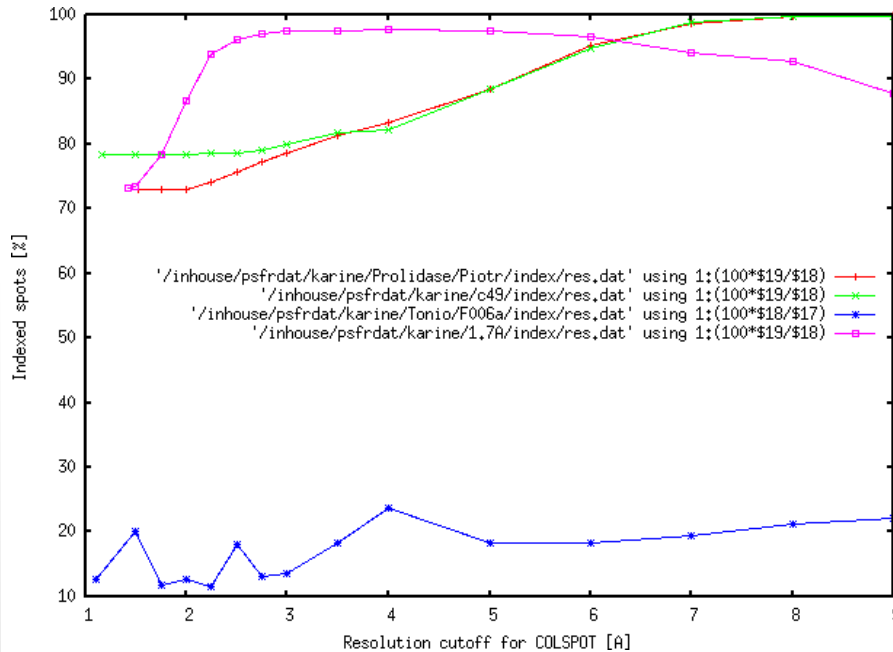
IDXREF

Time performance only improved for IDXREF: collect all spots

Improved performance

Regions of interest for IDXREF

Better indexing through resolution-based selection of spots for IDXREF



Useful for pathological crystals

- Mosaic spread
- Split crystals

Easier indexing at low resolution

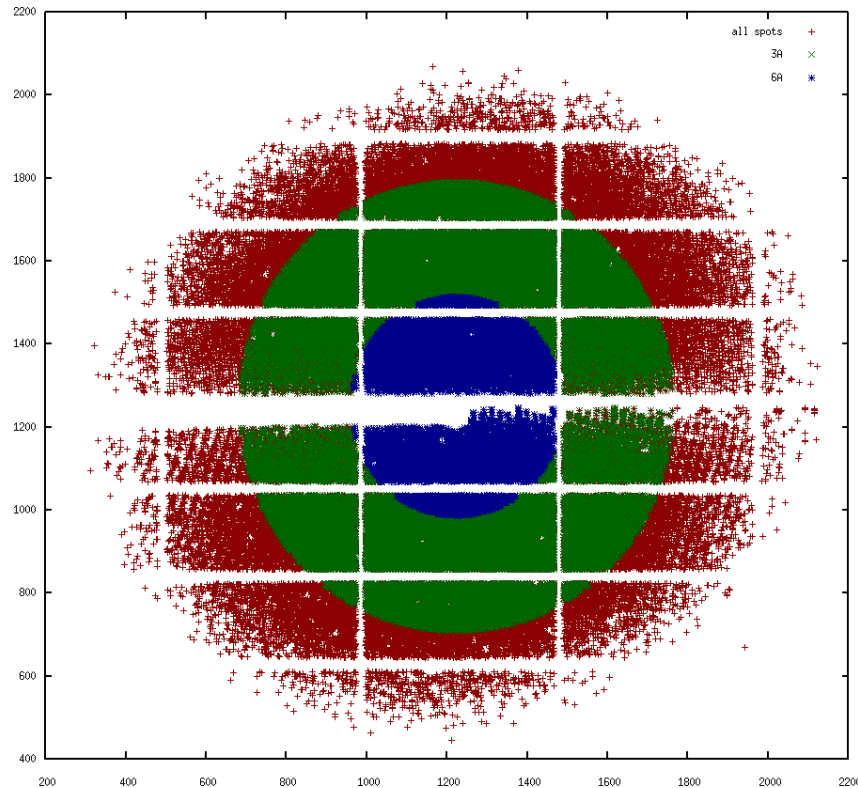
Successful indexing crucial for further processing steps

Indexed spots

Improved performance

Regions of interest for COLSPOT and IDXREF

Better indexing



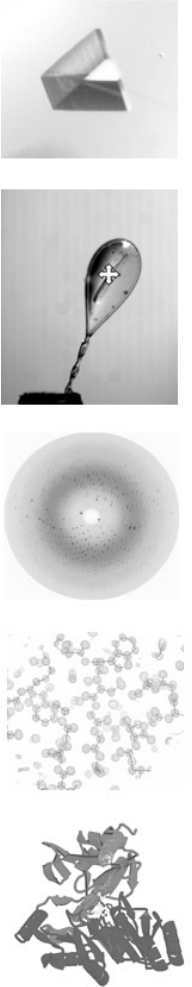
Spots used for IDXREF

Useful for pathological crystals

- Mosaic spread
- Split crystals

Easier indexing at low resolution

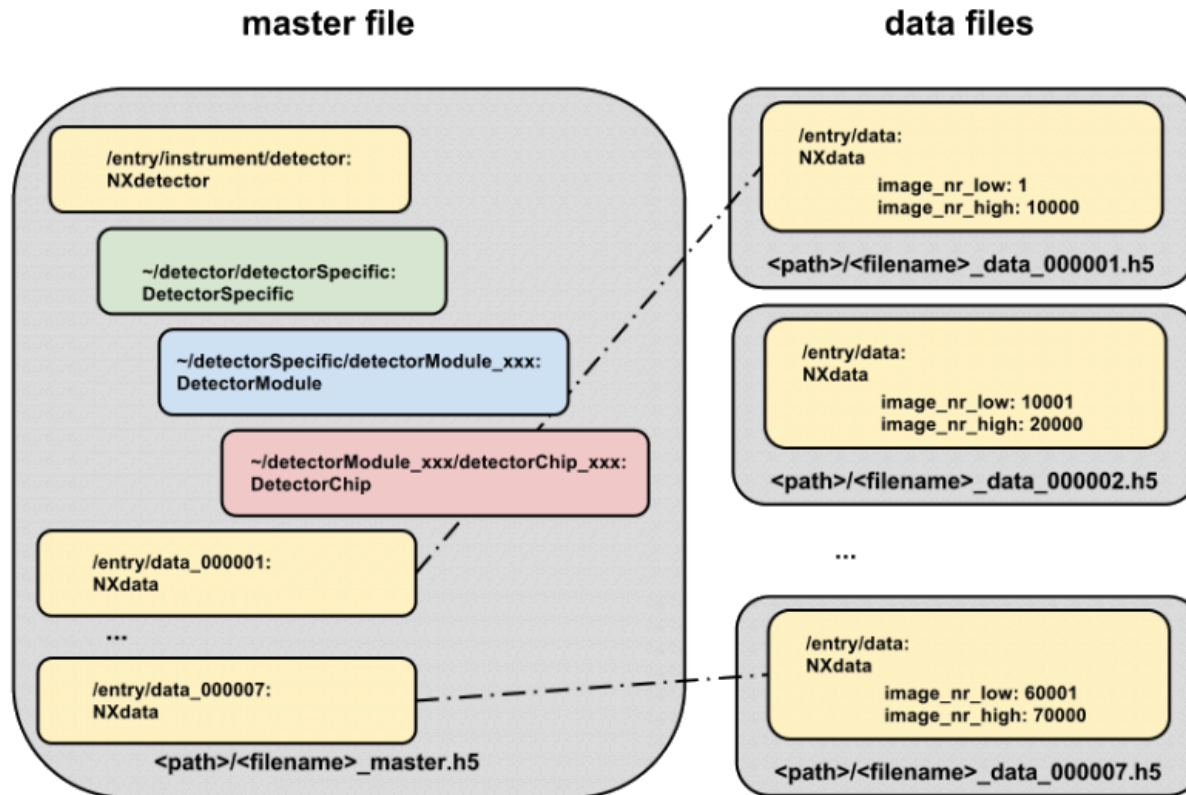
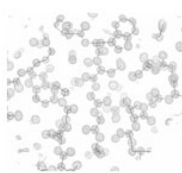
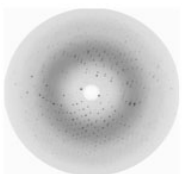
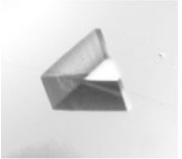
Successful indexing crucial for further processing steps



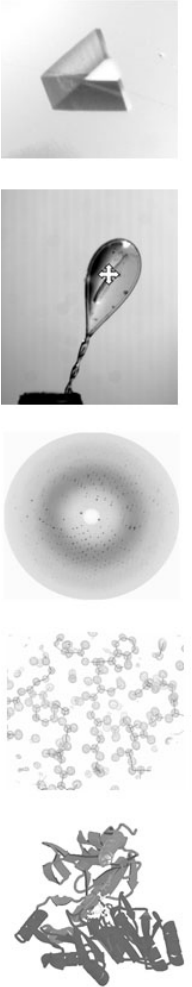
Eiger data sets

Even faster data collection
Finer ϕ -slicing
180 000 frames, 360GB / minute

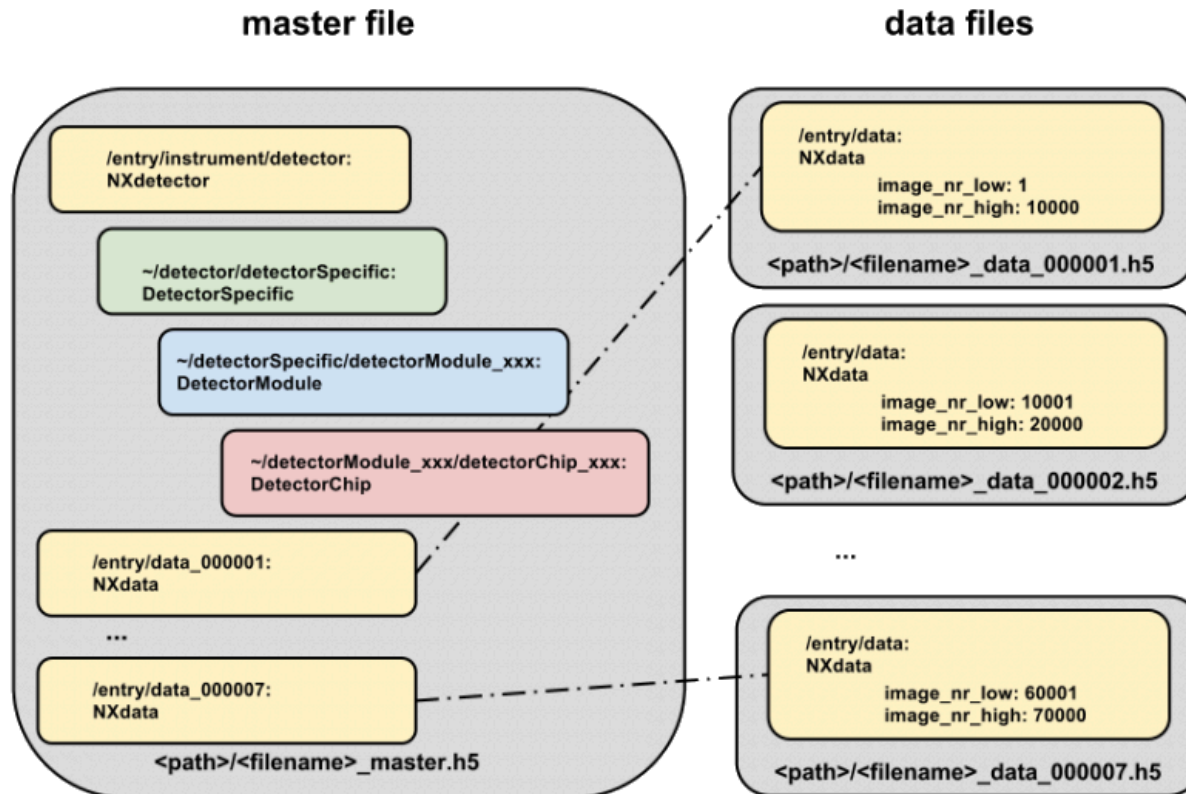
No native reading of .h5 files in XDS
Conversion to .cbf format



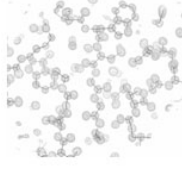
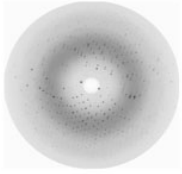
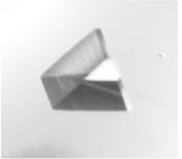
Eiger data sets



No native reading of .h5 files in XDS
Conversion to .cbf format



Eiger data sets



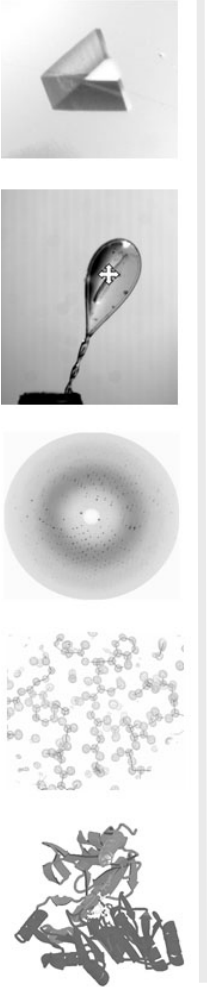
Solution compatible on all operating systems: eiger2cbf
Select either `_data_` or `_master` file

The screenshot shows a file manager window with the following table of files:

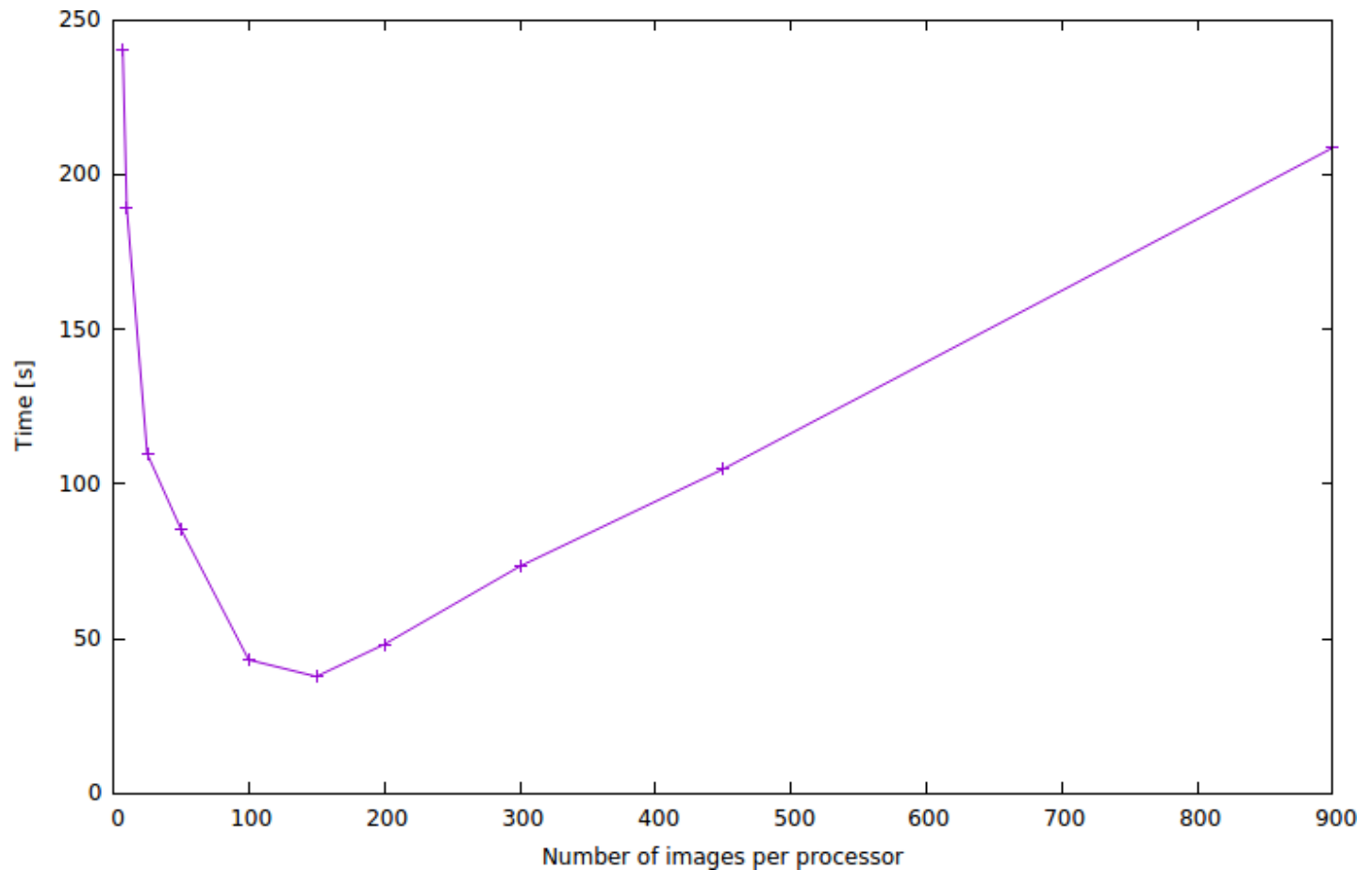
Name	Size	Modified
insu6_1_data_000006.h5	669,2 MB	08.02.2017
insu6_1_data_000007.h5	678,8 MB	08.02.2017
insu6_1_data_000008.h5	685,4 MB	08.02.2017
insu6_1_data_000009.h5	688,0 MB	08.02.2017
insu6_1_master.h5	366,2 MB	08.02.2017
tau1-tau_2_data_000001.h5	556,9 MB	08.02.2017
tau1-tau_2_data_000002.h5	547,9 MB	08.02.2017
tau1-tau_2_data_000003.h5	531,3 MB	08.02.2017
tau1-tau_2_data_000004.h5	519,8 MB	08.02.2017
tau1-tau_2_data_000005.h5	514,3 MB	08.02.2017
tau1-tau_2_data_000006.h5	504,6 MB	08.02.2017
tau1-tau_2_data_000007.h5	500,1 MB	08.02.2017
tau1-tau_2_data_000008.h5	494,7 MB	08.02.2017
tau1-tau_2_data_000009.h5	489,1 MB	08.02.2017
tau1-tau_2_master.h5	366,3 MB	08.02.2017

Buttons: Cancel, Open

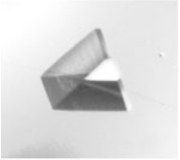
Eiger data sets



Solution compatible on all operating systems: eiger2cbf
Speed-up: 1 CPU per `_data_` file



Eiger data sets

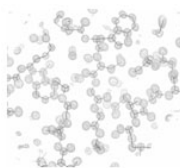
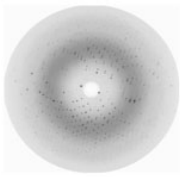


Possibility to load already converted .cbf files from eiger2cbf, thanks to modified generate_XDS.INP script by Olof Svensson (ESRF)



Tasks left

- Get rid of explicit data conversion in XDSAPP by using .h5 files directly in XDS, Linux only (H5ToXds)
- Delete converted .cbf files after processing
- Make the command line version of XDSAPP recognize Eiger data sets



Acknowledgment

Thank you for your attention

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