Processing data collected with Pilatus/Eiger detectors

James Parkhurst

IUCR Computing School, Bangalore, August 2017

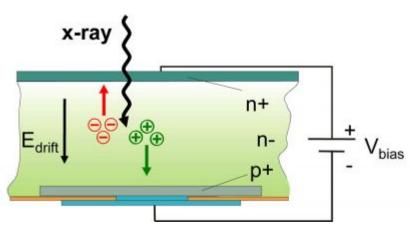
Introduction

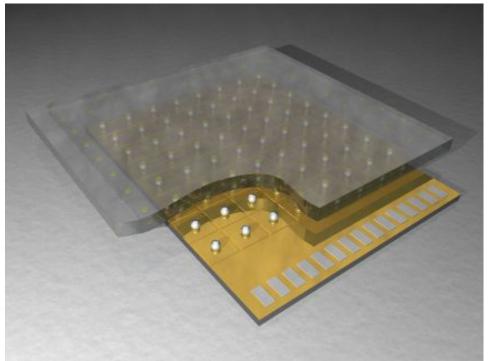
- Overview of Pilatus/Eiger detectors
- Overview of the DIALS integration program
- Data processing for Pilatus/Eiger detectors
 - Weak data
 - Spot finding
 - Background modelling
- Performance issues and parallelism

How does a Pilatus/Eiger detector work?

Sensor pixel: direct detection of X-ray photons -> one e-/hole pair per 3.6 eV.

Pixel electronics: counting of charge pulses.





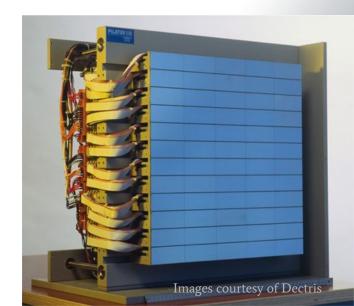
Modular detector

Pilatus/Eiger detectors are composed of modules: 8x2 array of CMOS ASICs

Each sensor module is a continuous 487 x 195 array of 94,965 pixels covering an active area of 83.8 mm x 33.5 mm

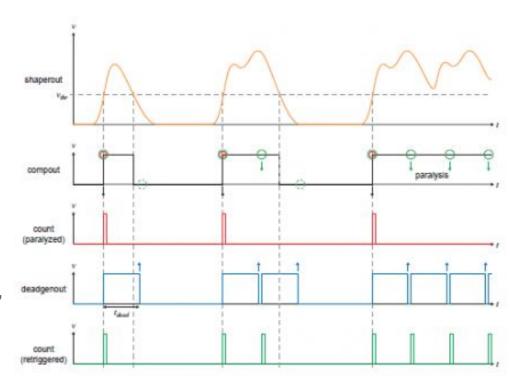
Modules are arranged to form larger detectors (Pilatus 6M contains 60 modules in a 5 x 12 grid)





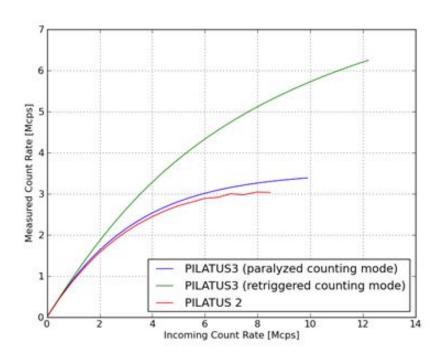
High flux - retriggering

- Incident X-rays converted to electric charge
- Once charge is greater than a threshold, a count is registered.
- In "paralyzed" mode (Pilatus 2/Eiger), another count is only registered after the charge decreases below and increases above the threshold again.
- In "retriggering" mode (Pilatus 3), if the charge stays above the threshold, another count is registered after a certain time.



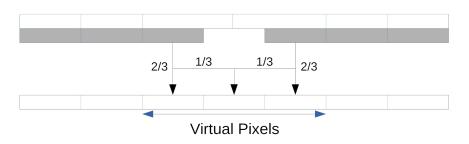
Count rate

- Due to the counting process there is a small dead time after each hit
- This becomes significant at high flux where some counts are lost
- The measured count rate is linear up to about 1Mcps



Virtual pixels

- Each module contains an 8x2 array of CMOS readout chips
- Chips have a small gap between them.
- This is spanned by 2 larger pixels (1.5x size of normal pixel)
- Counts are distributed into three "virtual pixels" after readout.
- The counts in the virtual pixels are therefore correlated.





Pixel array detectors

- Direct detection of X-rays
- Single-photon counting
- Good signal-to-noise ratio and high dynamic range (zero dark signal, zero noise)
- Low-energy X-ray suppression (energy resolution by single energy threshold)
- Short readout time and high frame rates
- Modular detectors enabling multi-module detectors with large active area

DIALS

Acknowledgements

research papers

Acta Crystallographica Section D Biological Crystallography ISSN 0907-4449

Wolfgang Kabsch

Max-Planck-Institut für Medizinische Forschung. Abteilung Biophysik, Jahnstrasse 29. 69120 Heidelberg, Germany

wolfgang.kabsch@mpimf-heidelberg.mpg.de

XDS

The usage and contro

package XDS for

described in the con

include automatic de

range and recognition

Moreover, the limita

number of correction

pixel contents have b

been restructured fo

and completeness of

research papers

Acta Crystallographica Section D Biological Crystallography ISSN 0907-4449

I. W. Pflugrath

Molecular Structure Corporation, 9009 New Trails Drive, The Woodlands, TX 77381, USA

Correspondence e-mail: jwp@msc.com

sensitive detectors can be cha depending on whether the rotatic is greater than or less than the c The expectations and consequen and thin images in terms of spa-X-ray background and $I/\sigma(I)$ a software suite for processing of

The finer things in

X-ray diffraction images from

introduced, and results from d

those from another popular pacl

1. Introduction

Two-dimensional position-sensiti for many years in X-ray diffract cular, data from crystals of maci oligonucleotides and their cor acquired with an area detector obsolete), a multi-wire system recently commercialized char coupled to a phosphor-coated fit detectors, the crystal, centered in oscillated around a single axis th ~2.0°, while counts from diffrac for a specified time. At the en detector is read out and the cou two-dimensional array with each to a distinct position on the c

research papers

Acta Crystallographica Section D Biological Crystallography ISSN 0907-4449

Andrew G. W. Leslie

MRC Laboratory of Molecular Biology, Hills Road, Cambridge CB2 2QH, England

Correspondence e-mail: andrew@mrc-lmb.cam.ac.uk

The integration

The objective of any

produce from a set of

with their associated uncertainties), togeth crystal unit-cell param reliable, but should i intervention. The pro three stages. The first parameters and the o parameters may indica The second step is to r parameters and also t known as post-refiner images, which consists reflections on each in

intensity of each reflect

out while simultaneou

parameters. Basic fea

each of these three so

with reference to the 1. Introduction

The collection of mac gone dramatic advan advent of two-dimensi and CCDs, crystal cry monochromatic and

Centre National de la Recherche Scientifique Université Paris-Sud

Laboratoire pour l'Utilisation du Rayonnement Electromagnétique

Proceedings

of the EEC Cooperative Workshop

on Position-Sensitive Detector Software

(Phases I & II)

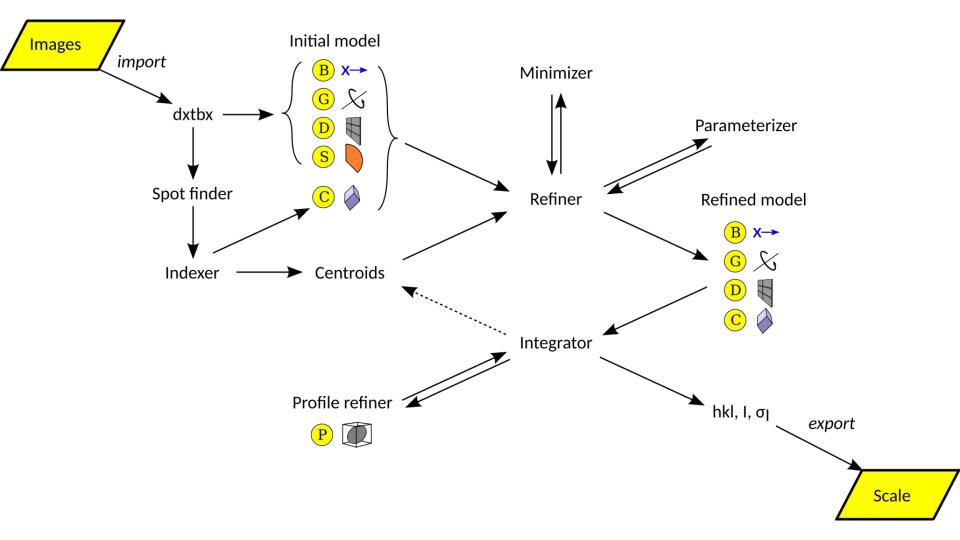
held at L.U.R.E. from May 26 to June 7, 1986.

1. Functional speci

measurement.

The program package developed for the re recorded on a plana monochromatic X-ra

XDS accepts a rotation images from and multiwire area metrics and produces of the reflections occur way. The program as positive amount of c incident beam and cr imposes no limitati directions of the rot oscillation range cov



Transitions

CCD PAD: Pilatus PAD: Eiger

PLATUS 3X

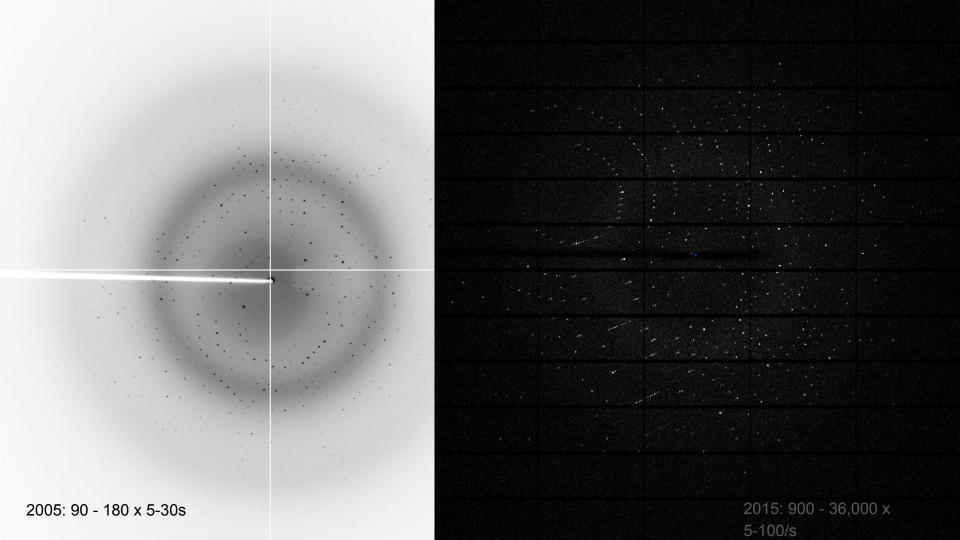
PILATUS3 S and X product pages ...

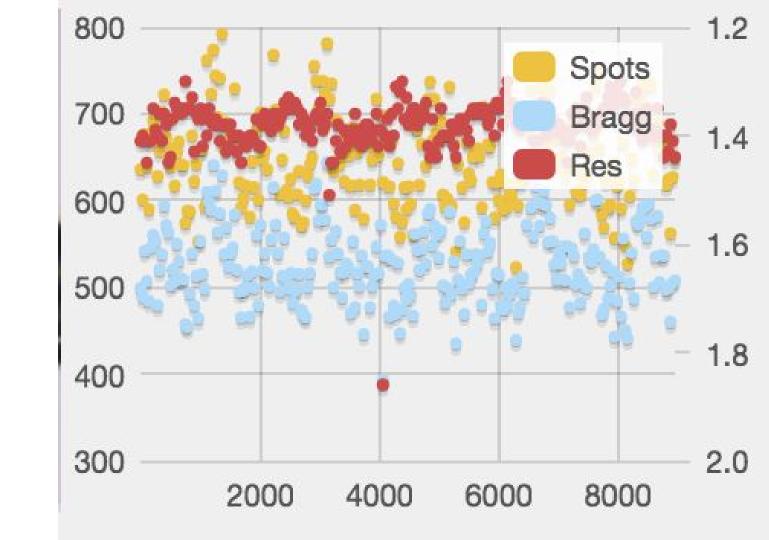


EIGER X product pages...

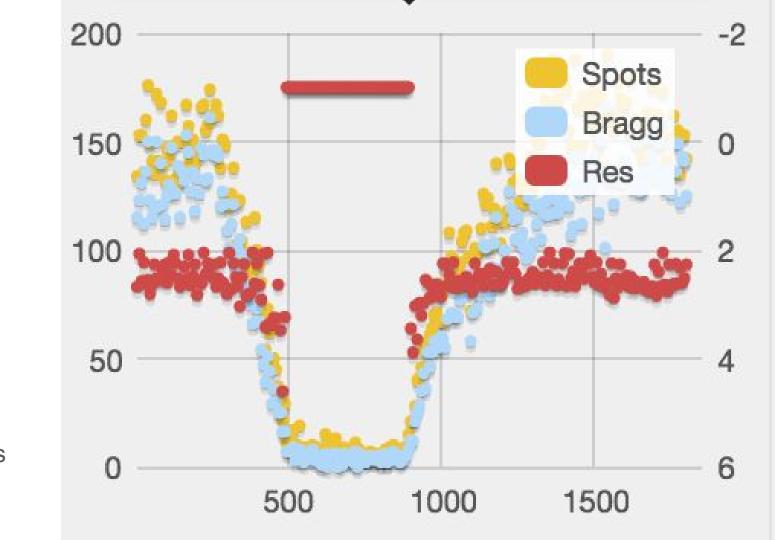
New Algorithms

New infrastructure



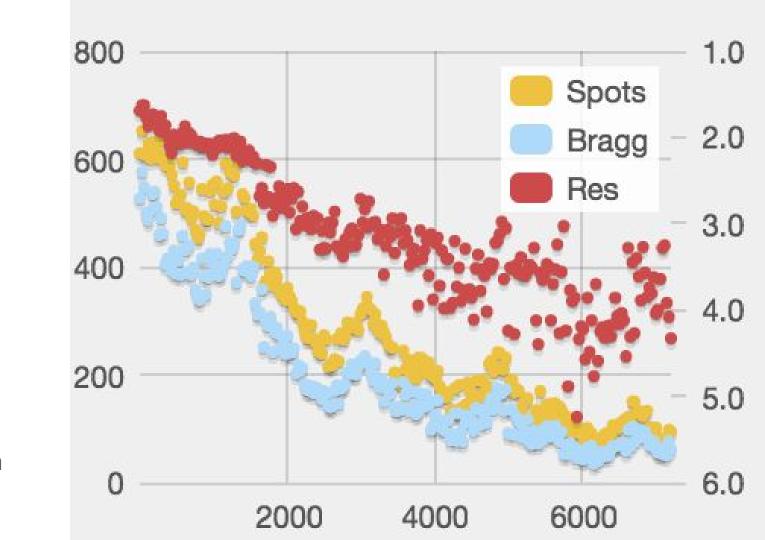


Good - lots of indexed spots



Bad - crystal leaving beam

Could treat as two sweeps



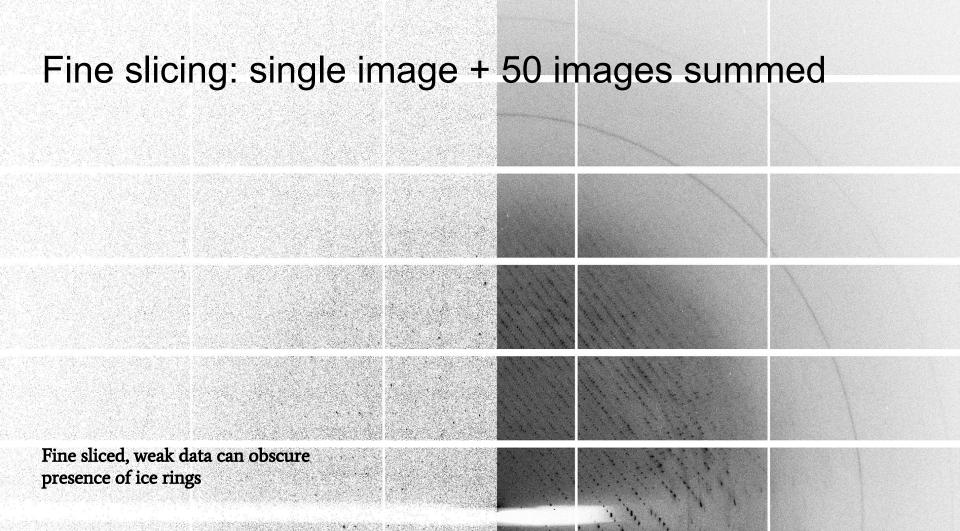
Bad - radiation damage

To what resolution do my spots extend?



Spot finder view of the image

Fine slicing: single image



Complex detectors: DLS BL-I23

Traditionally, integration programs supported collection from a single flat panel detector.

Modern integration programs need to support multi-panel detectors with complex configurations such as the Pilatus DLS 12M @ Diamond beamline I23

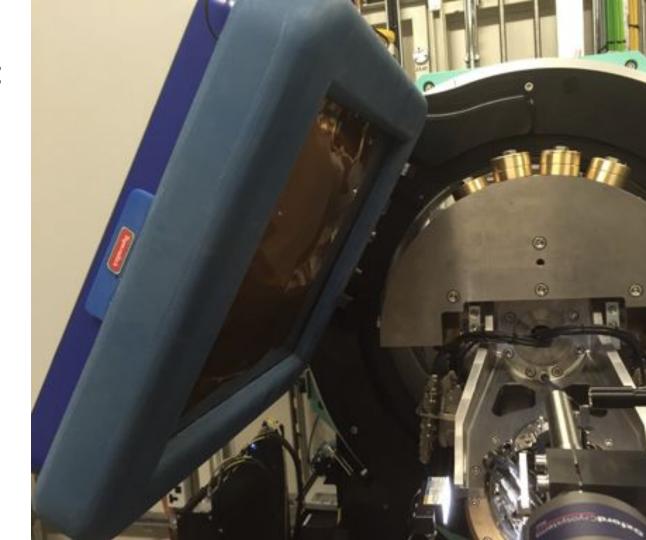


Image courtesy of Dectris

Complex detectors: DLS BL-I19

Detector is mounted on a goniometer so it can be positioned around the sample - including vertically above the sample (i.e. 90 degree to the incident beam).

The familiar concept of the "beam centre" is not really appropriate for this scenario.



CCD -> PAD: weak data

Data collection with CCD:

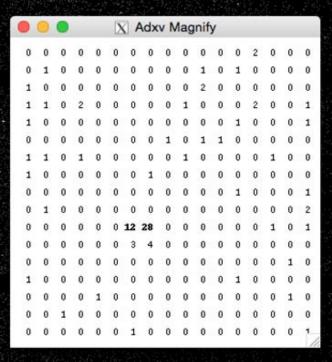
- Need to balance signal to readout noise, dark current, etc.
- Therefore need to collect strong data to get good I / Sig(I).

Data collection with PAD (Pilatus/Eiger):

- Very low readout noise in detection process means no compromise necessary
- Therefore dose / radiation damage can be spread around reciprocal space more uniformly.
- We can collect weak data with a very low background.

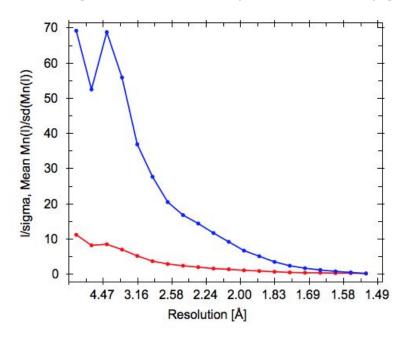


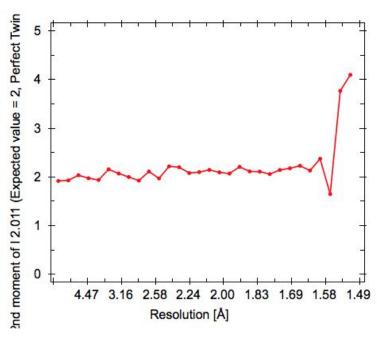
Example: Thaumatin recorded on 103, Pilatus 26M with 0.05% transmission. Most pixels are zero.



Data processing

- I/sig(I) well behaved, tend to 0 at high resolution
- 4th moment of E well behaved to high resolution
- Rmerge in outer shell crazy, but data broadly good, Rpim overall < 1%



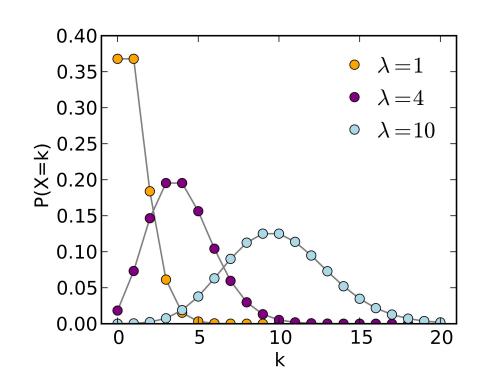


Poisson Distribution

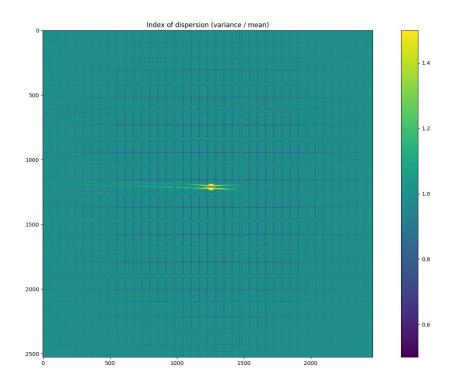
- Approximately normal for large lambda
- Not the case for small lambda

Useful properties

- variance = mean
- D = variance / mean = 1
- D is chi-squared distributed with n (n-1) degrees of freedom
- variance of D = 2 / (N-1)



Pixel array detectors: statistics



Analysed 9000 blank images and computed the index of dispersion (D = variance / mean) at each pixel.

For a Poisson distribution variance = mean, so we expect D = 1

Background data is Poisson distributed

Virtual pixels show under-dispersion due to correlations with neighbouring pixels

~7.2% of pixels are affected

Spot finding

dials.find_spots

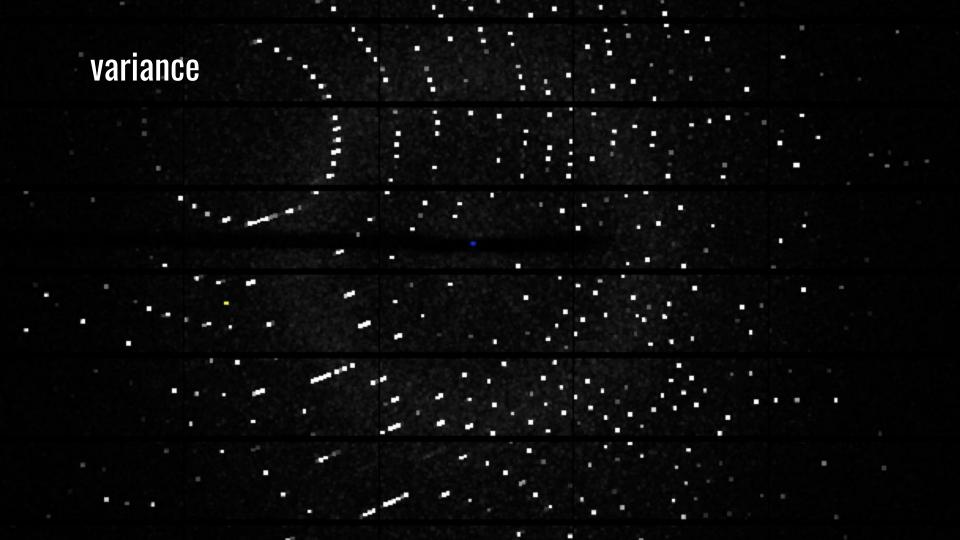
- Sequence of per-image filters to find strong pixels
- 3D analysis of strong pixels to identify strong spots
- Filter spots by
 - number of pixels
 - peak-centroid distance
 - resolution
 - ice rings
 - untrusted regions

\$ dials.find_spots datablock.json nproc=8

Saving 116082 reflections to strong.pickle Saved 116082 reflections to strong.pickle Time Taken: 31.768495

raw data

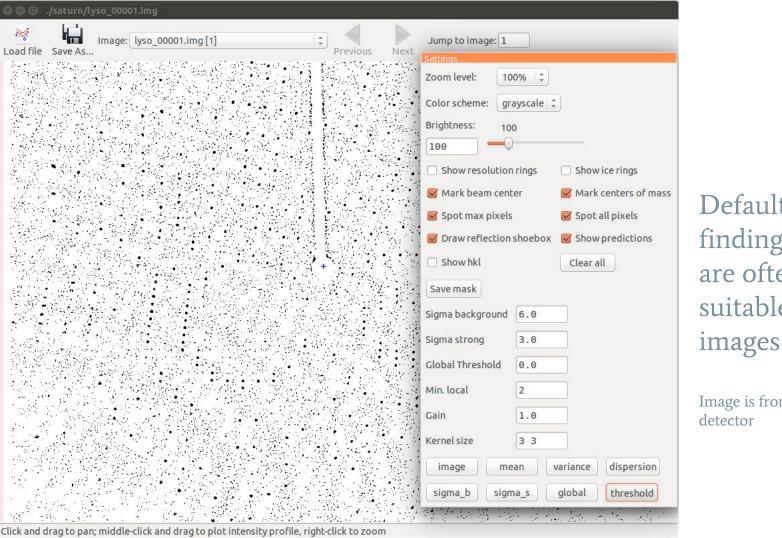




dispersion = variance / mean

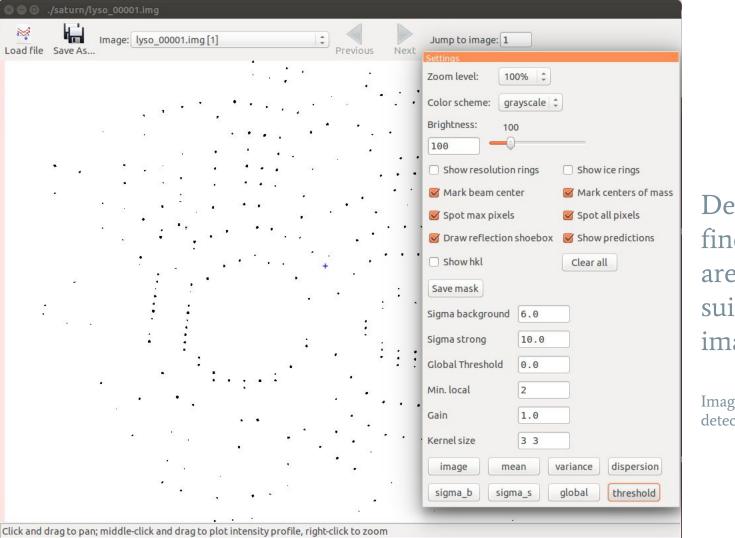
dispersion > $1 + sigma_s * sqrt(2/(n-1))$

raw data > mean + sigma_b * sqrt(variance)



Default spot finding parameters are often not suitable for CCD

Image is from Rigaku Saturn 92 detector



Default spot finding parameters are often not suitable for CCD images

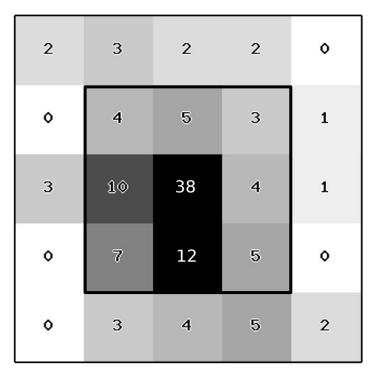
Image is from Rigaku Saturn 92 detector

Summary

- Pilatus and Eiger detectors are statistically well-behaved
- Pixels obey Poisson statistics
- Counts in virtual pixels are under-dispersed relative to a Poisson distribution
- Gain is equal to 1 across the detector, unlike CCDs which can have different per pixel gain values
- Spot finding works very well for Pilatus detectors, even when "strong" spots are "weak".

Background modelling

Integration



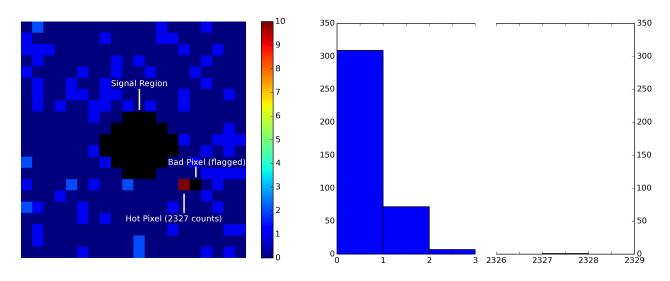
Summation integration: estimate the reflection intensity by summing the counts contributing to the reflection and subtracting the background

I = SUM(Counts - Background)

Profile fitting: fit a known profile shape to the reflection to estimate the intensity

Need to estimate background under reflection peak since it can't be measured directly

Background outlier pixels



	With Hot Pixel	Without Hot Pixel		
Mean	6.20	0.22		
Variance/Mean	2237.90	0.926		

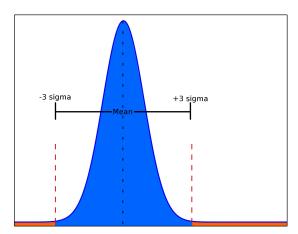
~1 for Poisson distribution

Outlier handling methods: simple

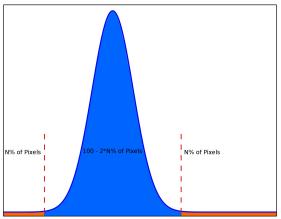
outlier.algorithm=nsigma

outlier.algorithm=truncated

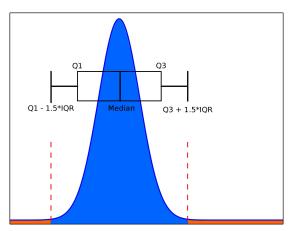
outlier.algorithm=tukey



Reject pixels N sigma from the mean



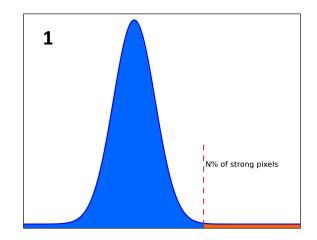
Reject N% of the highest and lowest valued pixels



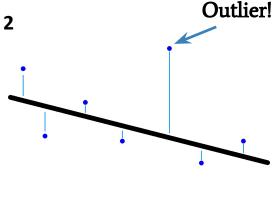
Reject pixels based on the interquartile range

Outlier handling methods: mosflm algorithm

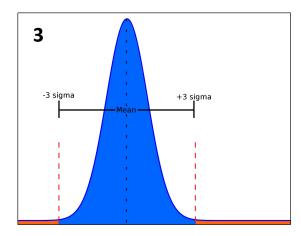




Remove N% of strongest pixels and compute the background plane



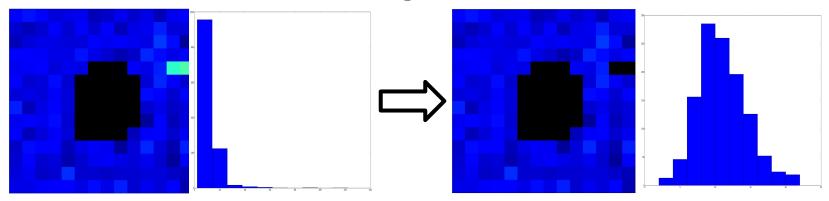
Compute the residuals of all background pixels to the plane



Remove pixels whose residuals are greater than N sigma from the plane

Outlier handling methods: xds algorithm*

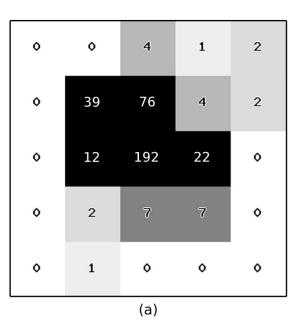
outlier.algorithm=normal

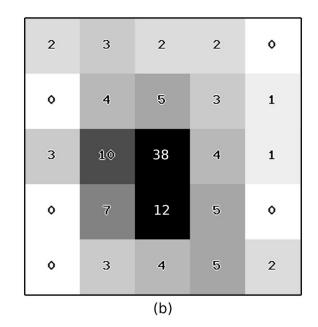


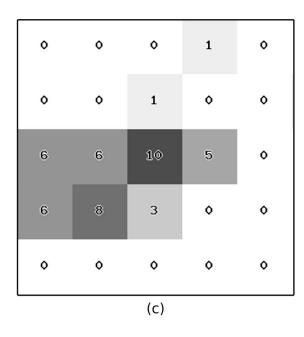
Iteratively remove high valued pixels until the distribution of pixel counts resembles a normal distribution

^{*} As described in Kabsch (2010) 'Integration, scaling, space-group assignment and post-refinement', Acta Cryst. D. 66(2), 133–44.

Pixel array detectors: low background





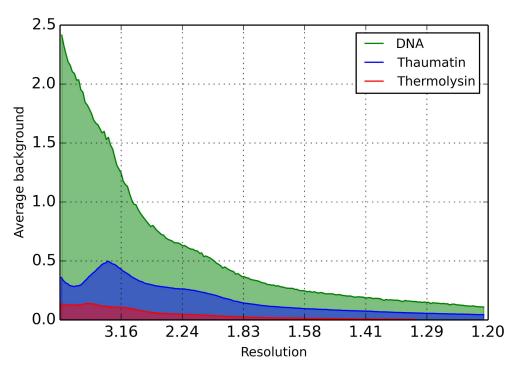


Thaumatin

DNA

Thermolysin

Pixel array detectors: low background



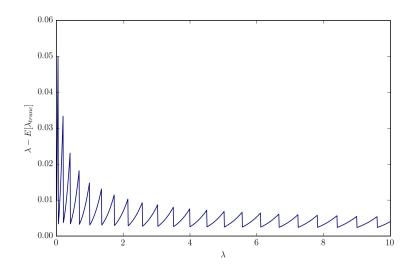
Each dataset has low background over entire resolution range.

Thaumatin and Thermolysin datasets have background less than 1 count per pixel over the whole resolution range

DNA dataset has background less than 1 count per pixel at high resolution

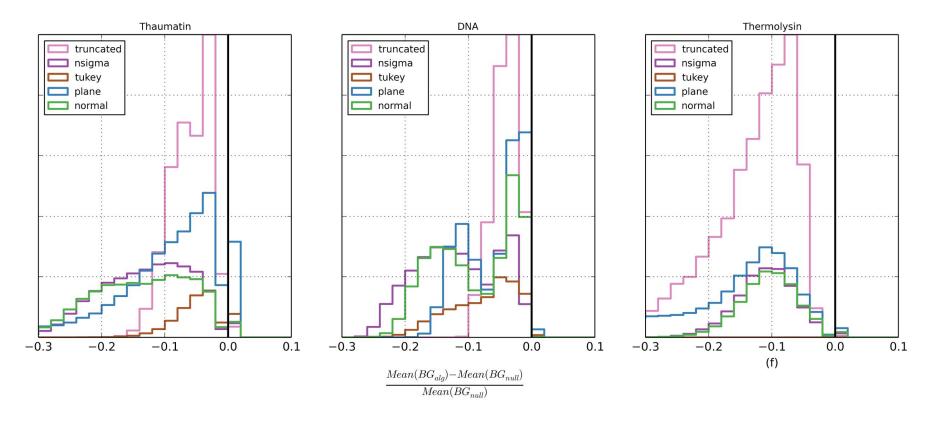
Bias in background determination

- Poisson distribution is asymmetric
- Truncation of the data results in bias in the background determination

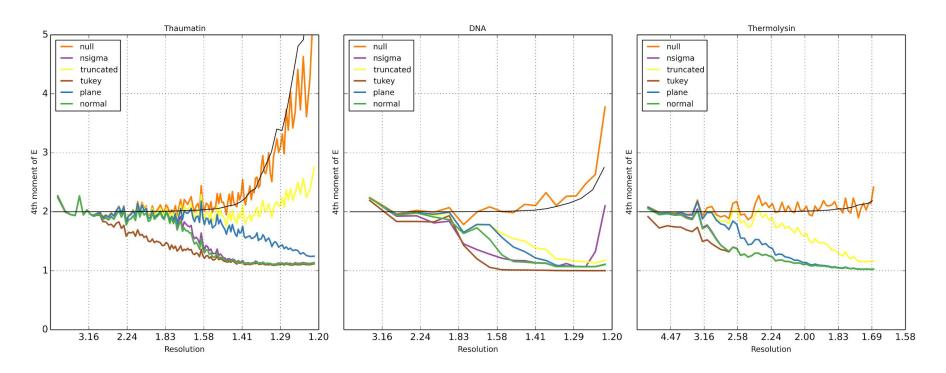


Q is the regularized gamma function

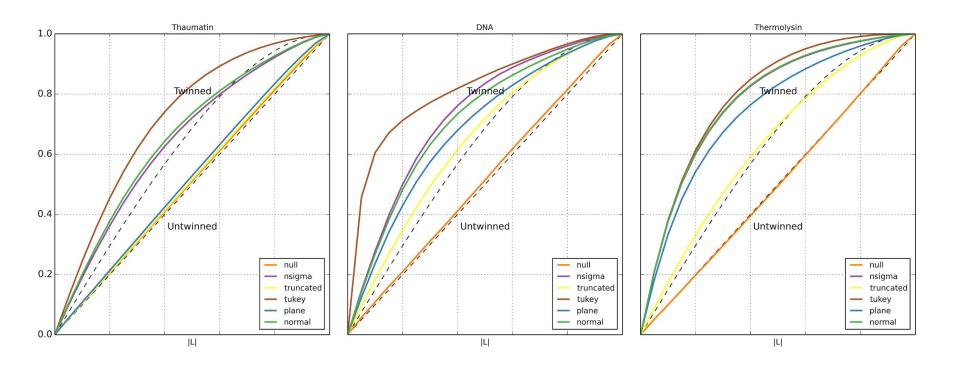
Bias in background determination



Bias in intensity statistics



Bias in intensity statistics



GLM background modelling

Eva Cantoni and Elvezio Ronchetti (2001), "*Robust Inference for Generalized Linear Models*", Journal of the American Statistical Association, Vol. 96, No. 455

Solve

Pearson residuals

Variance function

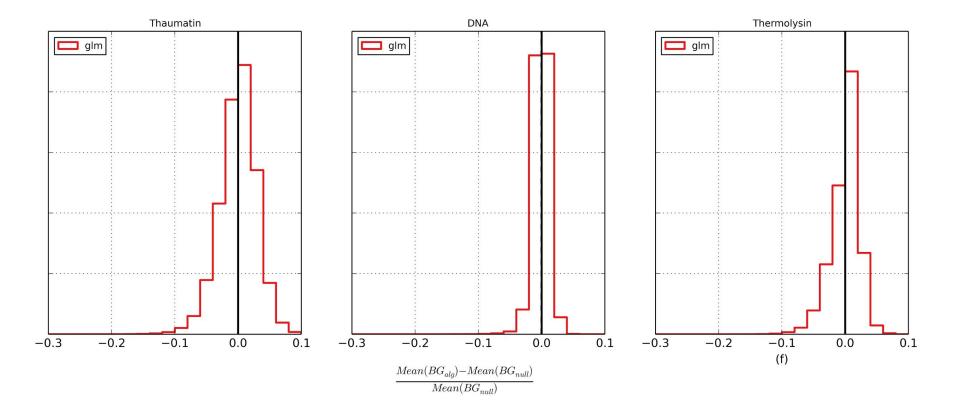
Weights for explanatory variables

Weights for dependant variables

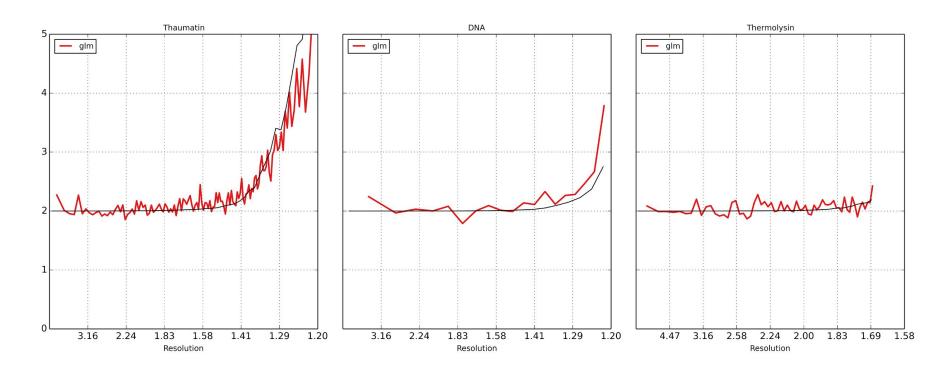
Tuning constant

Consistency correction

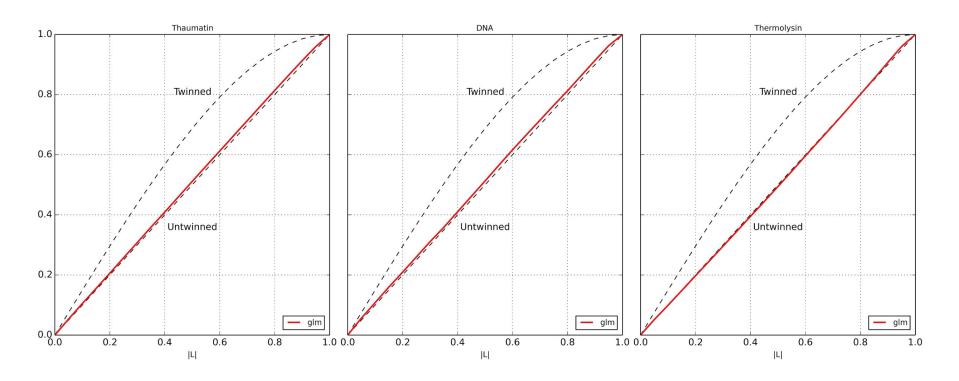
GLM method is unbiased



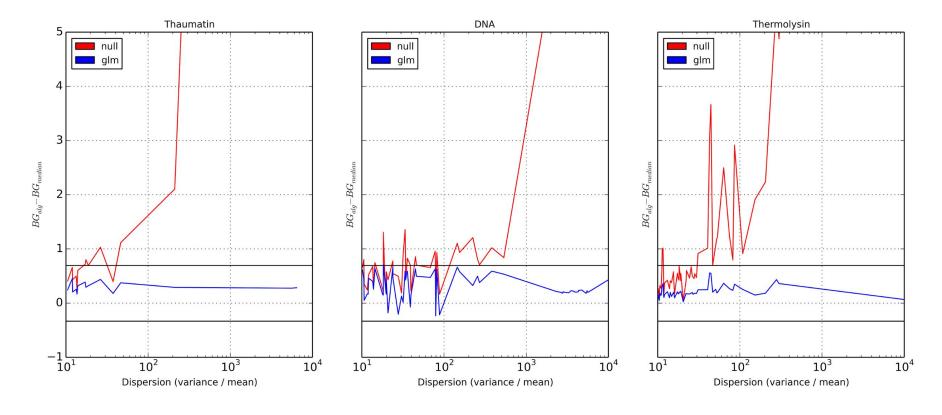
GLM method is unbiased



GLM method is unbiased



GLM method: handling pixel outliers



Twin test results

	Thaumatin		DNA		Thermolysin	
	L test	4th moment	L test	4th moment	L test	4th moment
truncated	0.04	0.00	0.50	0.28	0.50	0.23
nsigma	0.50	0.27	0.50	0.50	0.50	0.50
tukey	0.50	0.50	0.50	0.50	0.50	0.50
plane	0.06	0.01	0.50	0.42	0.50	0.50
normal	0.50	0.30	0.50	0.50	0.50	0.50
glm	0.03	0.00	0.04	0.00	0.03	0.00
null	0.03	0.00	0.05	0.00	0.03	0.00

Summary

- Traditional methods for handling pixel outliers systematically underestimate the background level
- Consequently they overestimate the reflection intensities even in the absence of any pixel outliers in the raw data.
- This can cause statistical tests to give the false impression that a crystal is twinned.
- The GLM method is robust against such effects.
 - When no outliers are present, the estimates given by the GLM algorithm are, on average, the same as those with no outlier handling;
 - When outliers are present, the method gives values within the expected bounds of the median.

Performance

Transitions

CCD PAD: Pilatus PAD: Eiger

PILATUS3 S and X product pages ...

EIGER X product pages...

New Algorithms

New infrastructure

Pilatus -> Eiger: algorithms and data

For DIALS:

- Detector behaviour is the same in both cases identical mathematical problem which is well supported
- One file per image (Pilatus CBF), now one file per scan (Eiger HDF5) easily handled via dxtbx
- Metadata stored in binary arrays in HDF5 easily handled via dxtbx
- HDF5 external references just work
- Fine slicing works fine with 3D profile fitting use the same algorithms.

HDF5 and Nexus

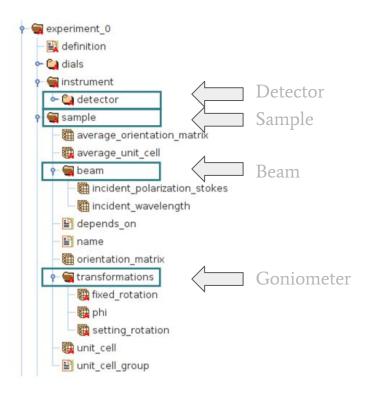
In the past, detectors typically wrote a file for each image. This is ok if the data rate is low and the number of files output is small.

This becomes difficult for the file system to handle when writing out huge numbers of files at a high rate.

The EIGER writes out 1 HDF5 file containing all the images from a single data collection.

EIGER HDF5 files use the Nexus data specification and can be read natively by DIALS.

Nexus HDF5 files



HDF5 is the file container. Image and metadata is stored in a hierarchical format.

Nexus provides the definition that allows programs to understand the HDF5 file.

Full NXmx specification available from:

http://download.nexusformat.org/sphinx/classes/applications/NXmx.html

CCD -> Pilatus -> Eiger: detector performance

- CCD detector processing data during collection feasible
- Pilatus @ 10 Hz data set around 3 minutes, fast processing OK, xia2 already "too slow" for interactive feedback
- Pilatus6M @ 10 Hz may as well wait for data to be finished before processing
- Pilatus6M @ 100 Hz data set in 18s time to give up on processing in real time, fast processing now too slow for real-time
- Eiger9M @ 200 Hz real-time effectively impossible

The problem

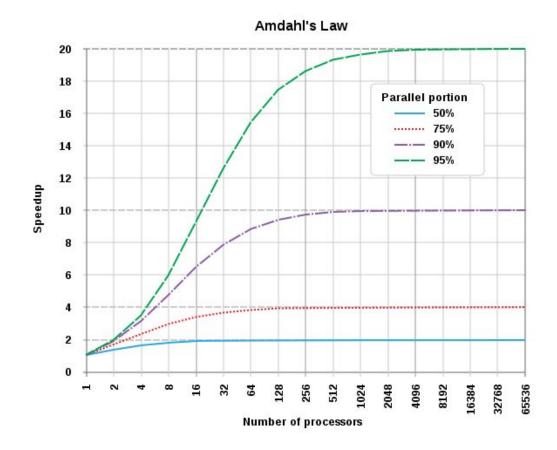
- Current detectors (DECTRIS Pilatus) run at up to 100 frames / s
- Next generation Eiger detectors run up to 750 frames / s (for 4M)
- This rate will probably not be routinely used for data collection
- This rate will be used for raster scanning i.e. to allow a large loop to be sampled with a fine beam in a short time (e.g. X-ray centering)
- For raster scanning the experiment has to wait for the results so this is time critical
- Therefore in first instance principle benchmarking problem is spot finding
- Need to make use of parallel processing

Amdahl's Law

Expected performance improvement from increased number of processors

p = parallel percentage

s = speed up (i.e. number of cores)

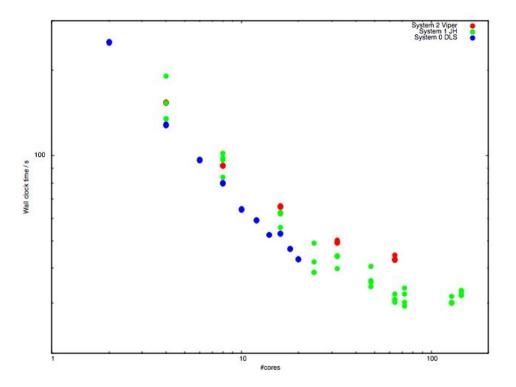


Benchmark

- Performed with dials 1.3.1 linux binaries (same binary set for all systems)
- dials.find_spots datablock.json nproc=\${nproc} shoebox=false
- Principle consideration wall clock time i.e. from starting process to results becoming available
- Here nproc=4...# in system
- Data come from RAMDISK => file system performance not a consideration

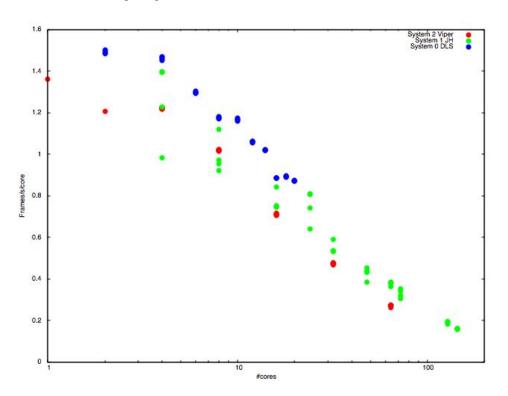
Wall clock time vs #cores

- Wall clock time decreases with increasing number of cores
- Decrease in wall clock time tails off at around 100 cores.



Frames/second/core ("efficiency") vs #cores

"Efficiency" of spot finding drops with increasing number of cores.



Summary

- "Efficiency" drops off rather quickly with increasing #cores [1]
- Wall clock time flattens off around 40 s for system 0 using 20 cores; ~ 30 s for system 1 using 144 cores
- For small #frames start up time (~ 4s) dominates
- For large #frames wall clock time ~ linear 0.08 s / frame (10 cores)
- We maybe need to put some effort into optimizing DIALS for many core architectures (e.g. system 1 above; Xeon phi; ...)
- Using small #cores but analysing each row of a grid scan on a separate node in a round-robin manner may be optimum for responsiveness

Acknowledgements

DIALS East

Gwyndaf Evans, Graeme Winter, David Waterman, James Parkhurst, Richard Gildea, Luis Fuentes-Montero, Markus Gerstel, Melanie Vollmar

DIALS West

Nick Sauter, Aaron Brewster, Iris Young

Lots of other people

Garib Murshudov, Andrew Leslie, Phil Evans, Harry Powell, Takanori Nakane, Andrea Thorn









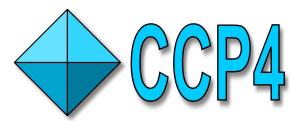












Thanks for listening!

https://dials.diamond.ac.uk