



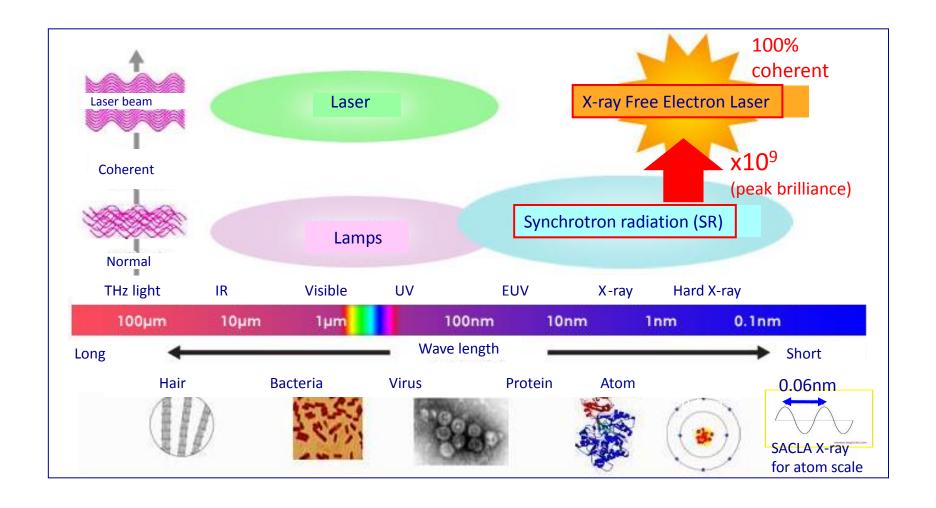
Data-analysis scheme and infrastructure at the X-ray free electron laser facility, SACLA

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X-ray Free Electron Laser (XFEL)



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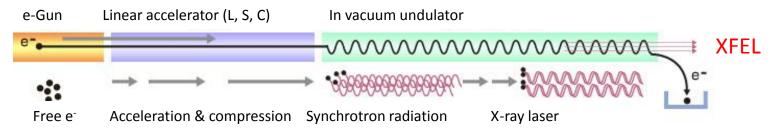
SPring-8 and SACLA

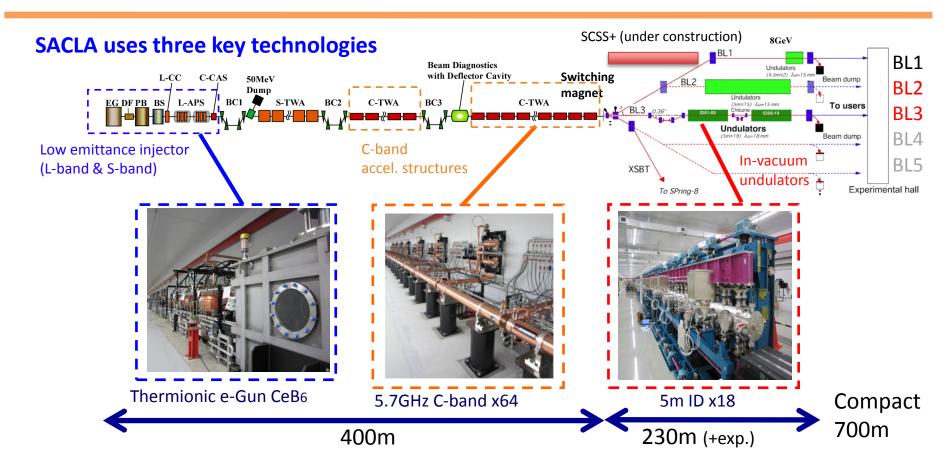




XFEL lasing scheme at SACLA

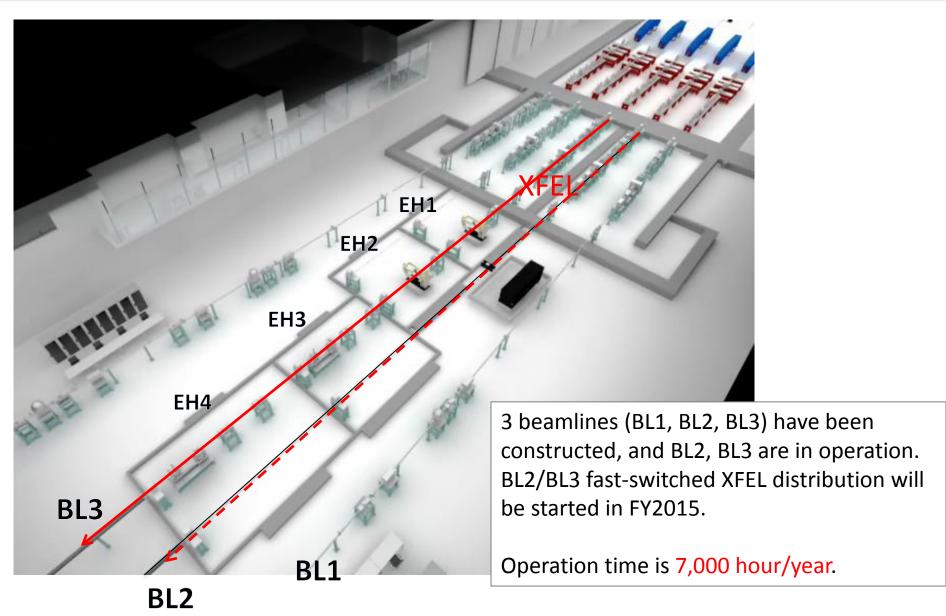
Lasing scheme





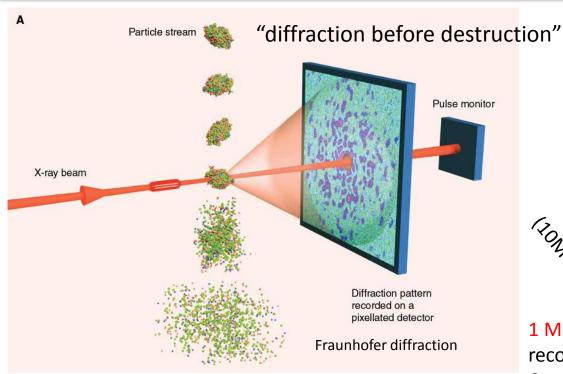


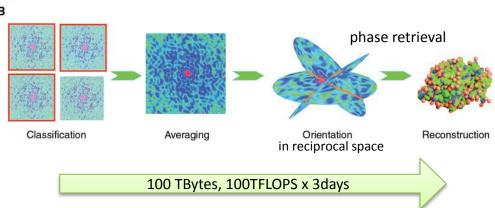
Beamlines



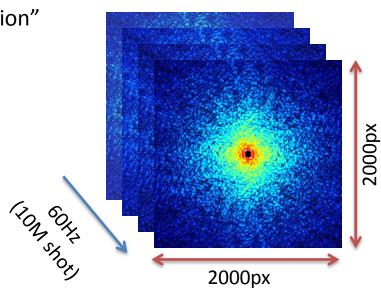


Coherent X-ray Diffraction Imaging (CXDI)





K. J. Gaffney and H. N. Chapman, Science 316, 1444 (2007)



1 M shots of diffraction images are necessary to reconstruct 3-D structure.

Considering XFEL hit rate on sample, 10 M shots data are required.

-> more than 100 TB storage per sample

To reconstruct 3-D image in real space, we must analyze relationship between 1 M images.

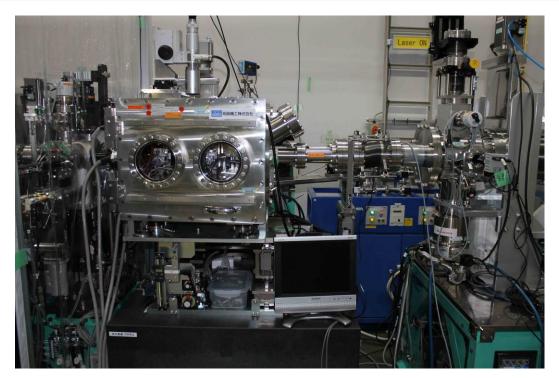
XFEL experiments requires heavy computing

resources.

-> 100 TFLOPS x 3 days (per sample)

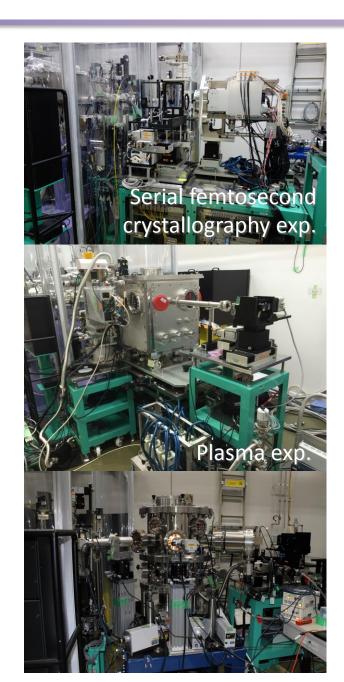


Experimental setups



CXDI Experiment

Experiment changes every few days. We must setup detectors and DAQs as fast as possible.



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Requirement for the DAQ and the analysis systems

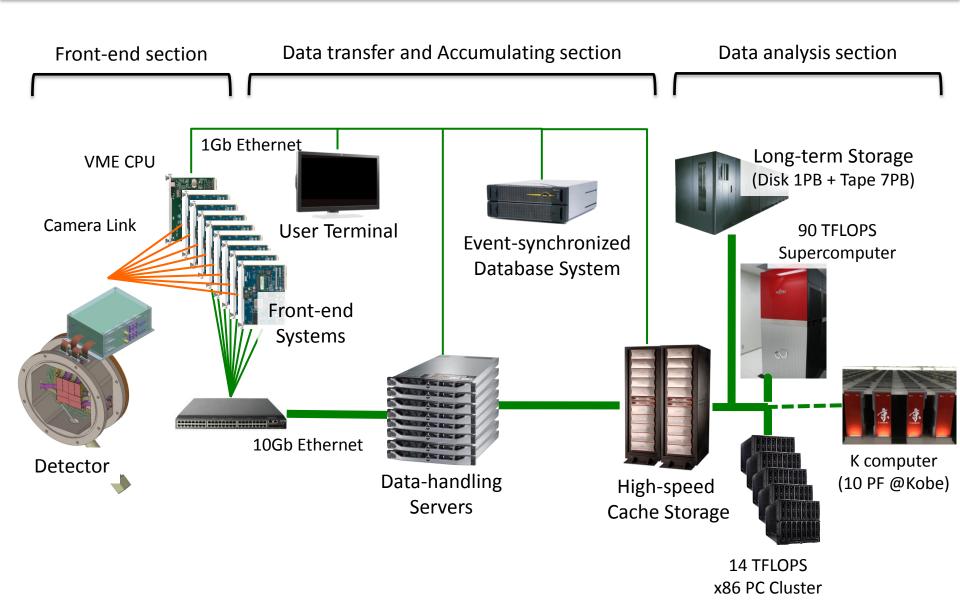
- Images are accumulated at 6 Gbps data rate
 - 12 MBytes (12 sensors) at 60 Hz repetition
- One experimental sample requires 120 TBytes storage
 - 12 MBytes x 10 M shots
- Short experimental period
 - Sample and/or detector setup change every few days
 - Users want to use the SACLA as a commercial microscope. (System tuning / DAQ is not their science)
- Analysis should be faster than the DAQ data rate
 - 10 TFLOPS for run-by-run analysis and pre processing (10M -> 1M)
 - 100 TFLOPS for 3-D structure reconstruction within a few days.



DAQ and Analysis System at SACLA

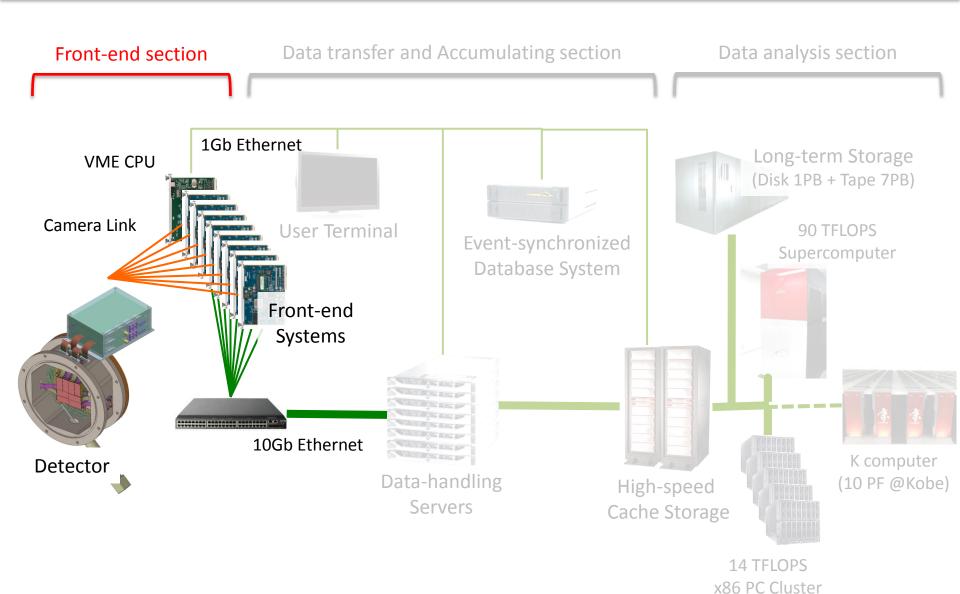


SACLA DAQ and Analysis System





Detector Front-end System





MPCCD sensor



MPCCD sensor (512 x 1024 px, 16bit depth)

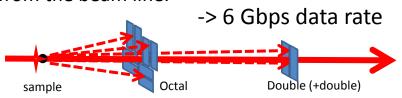
Trigger = accelerator repetition (60 Hz)

-> data rate is 0.5 Gbps/sensor



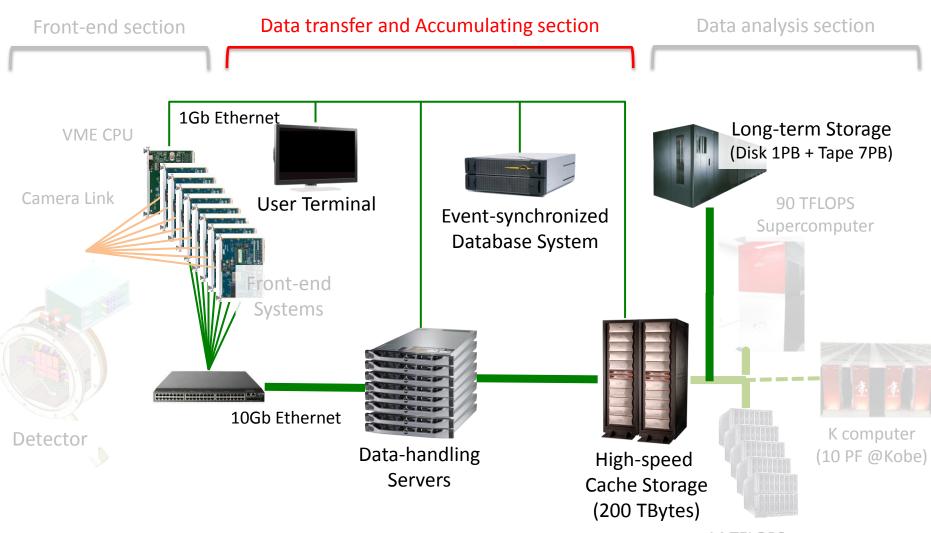
T. Kameshima, et al., Rev. Sci. Instrum. 85, 033110 (2014).

To avoid radiation damage from direct XFEL beam, 8+2+2 sensors are installed offset from the beam line.



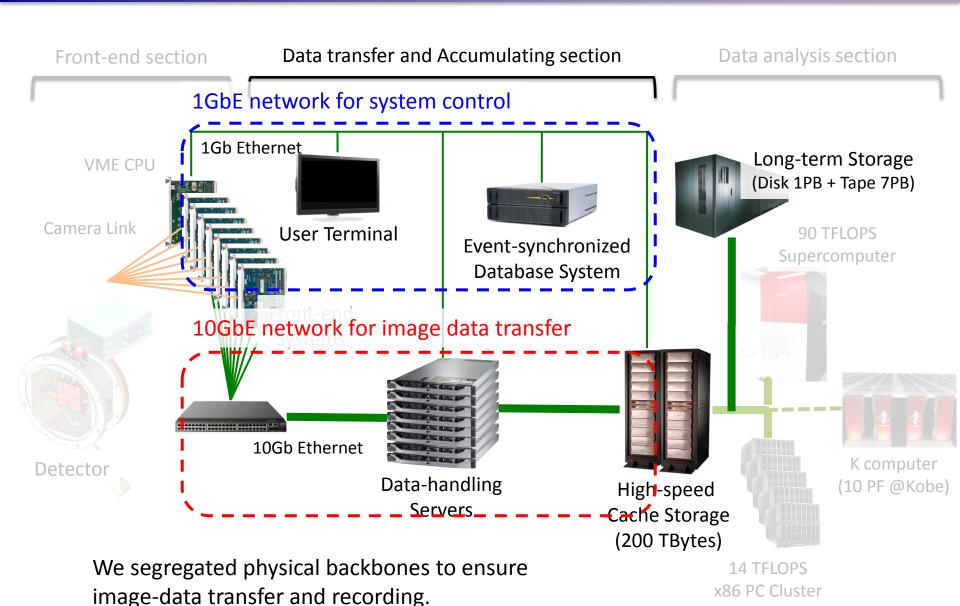
Note: All images (including junk) are recorded. Because it is difficult to determine reduction condition within few-day experimental period.





14 TFLOPS x86 PC Cluster

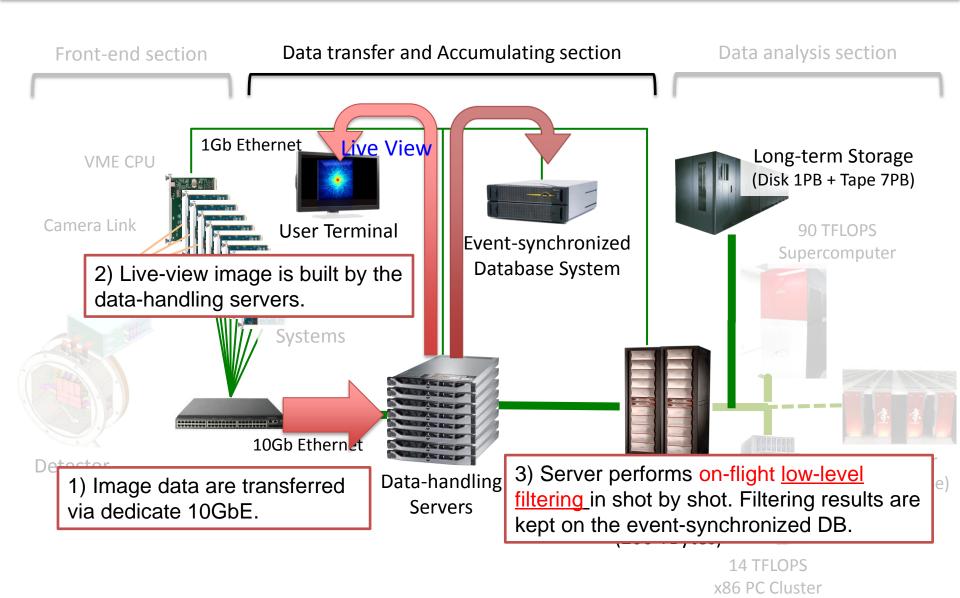




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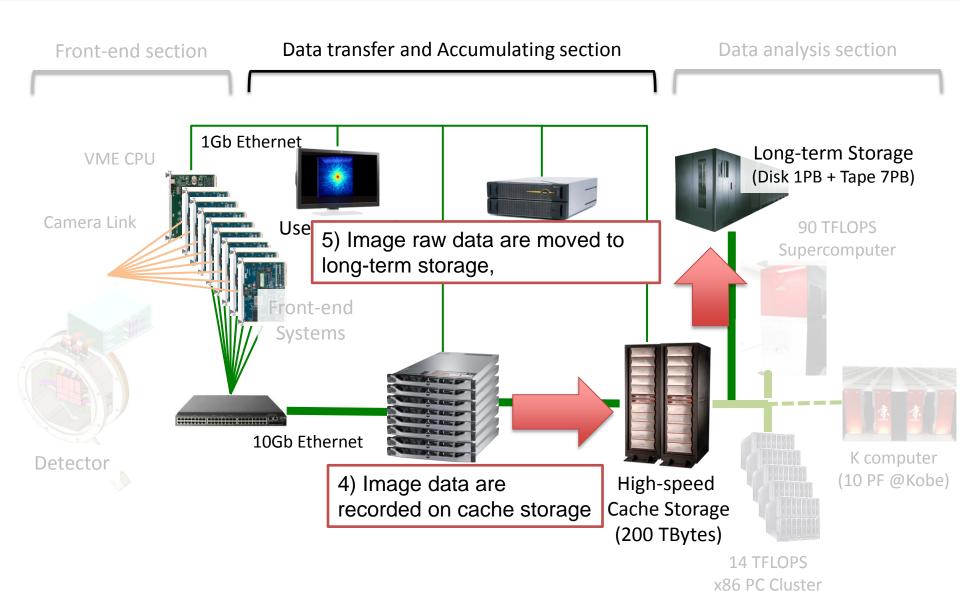
DAQ Data flow



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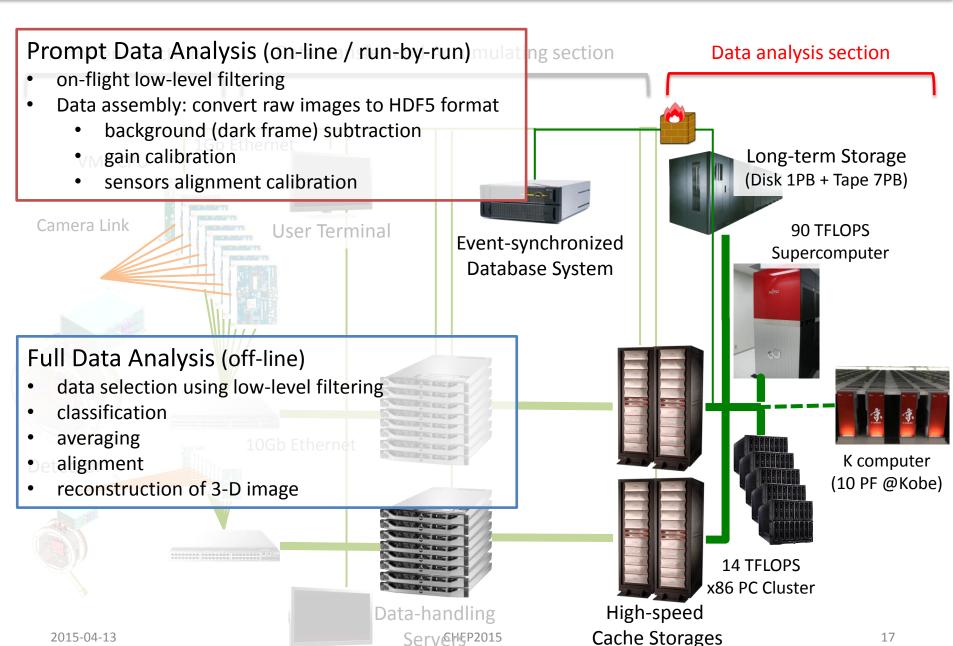
DAQ Data flow



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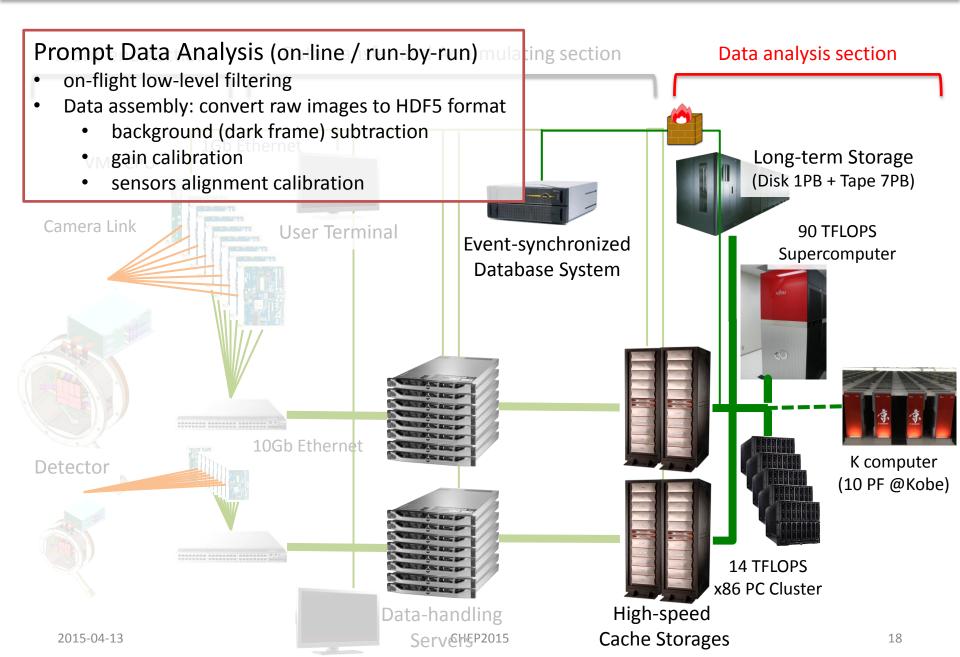


Data analysis section



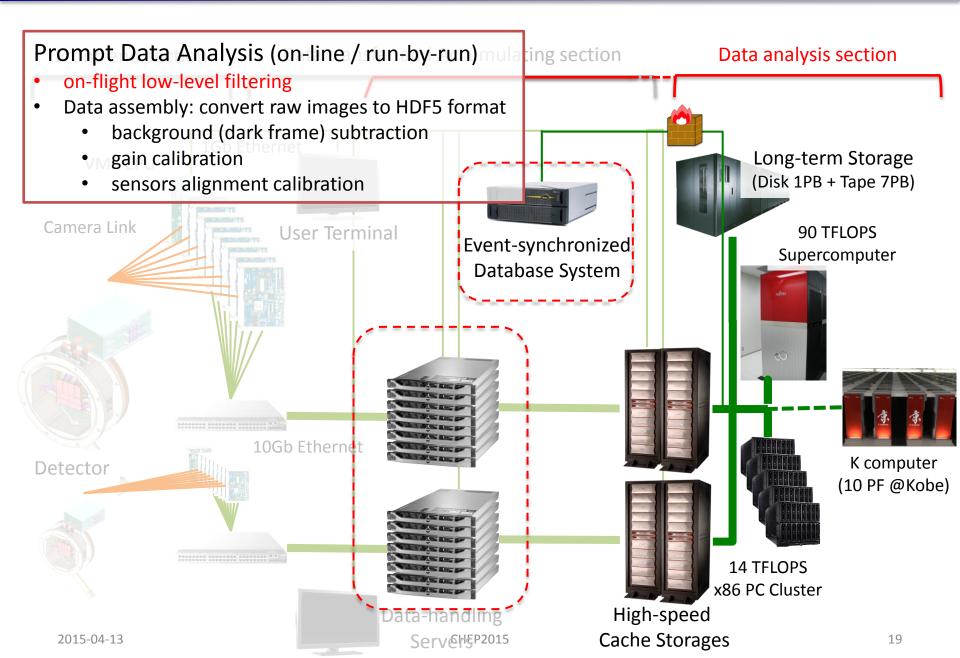


Prompt data analysis





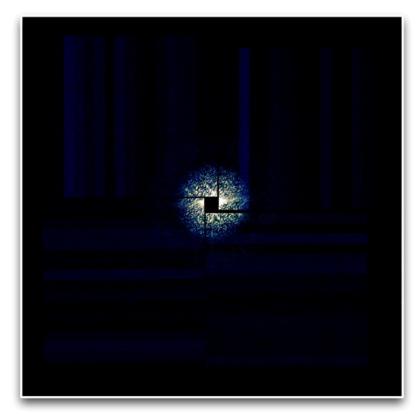
Prompt data analysis





Prompt data analysis: Low-level filtering

To reduce latter computing load, we want to identify candidates of GOOD event.



Exapmle of GOOD event (diffraction pattern)



Example of BAD event (scattered by water)

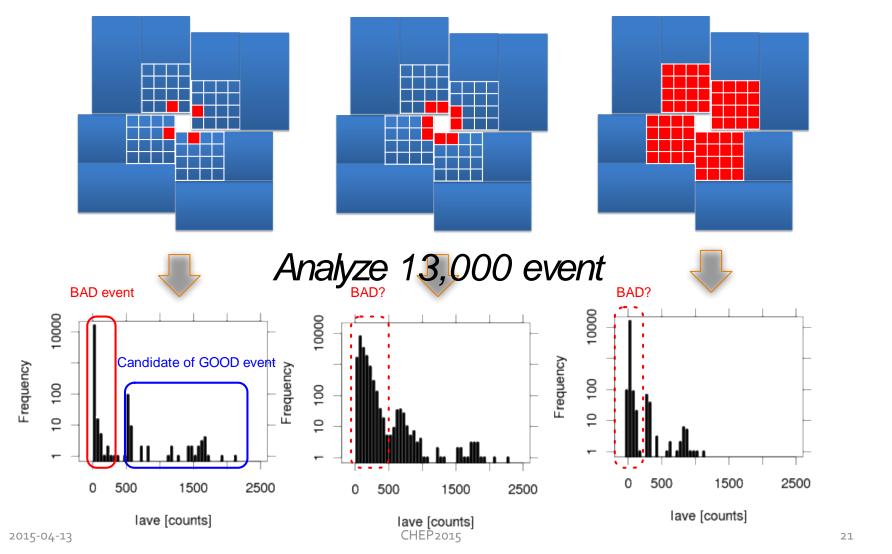


Prompt data analysis: Low-level filtering

To reduce latter computing load, we want to identify candidates of GOOD event. Intensities in the mesh are recorded on event-synchronized DB.

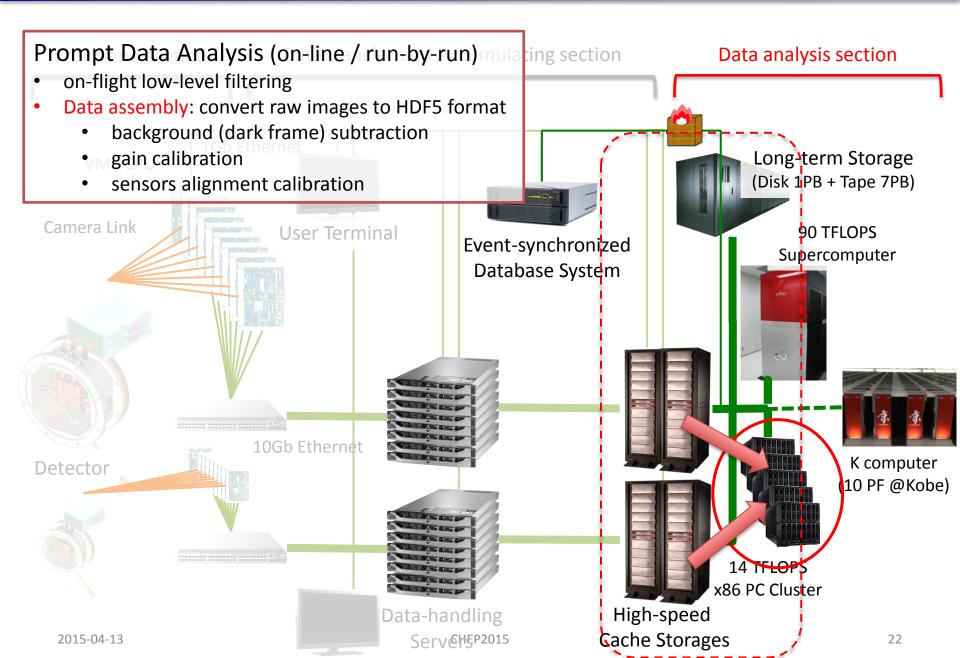
By selecting ROI, we can distinguish GOOD event from bad event.

Region of Interest (ROI)



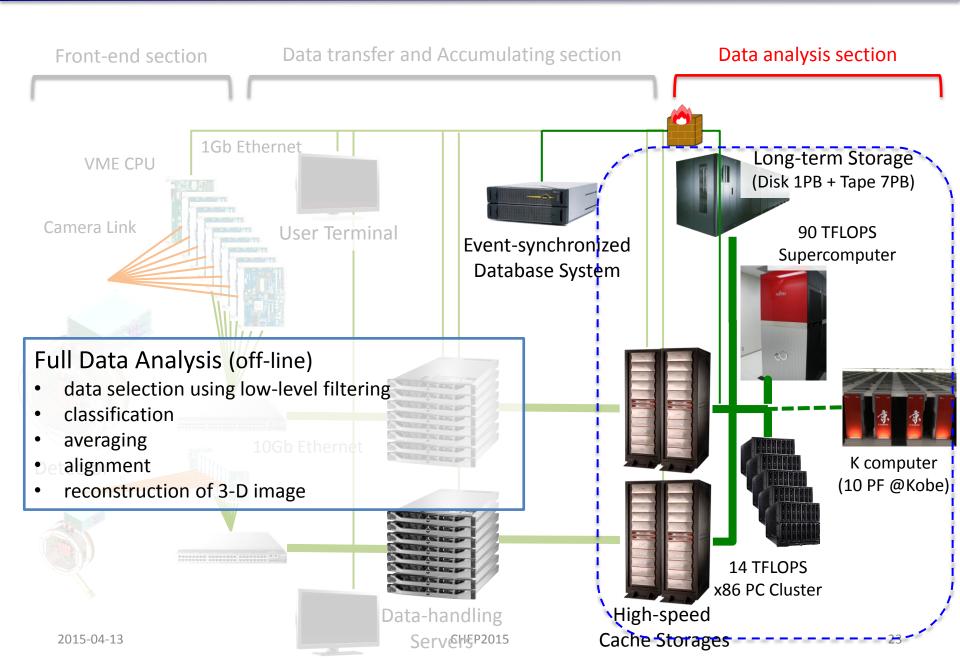


Prompt data analysis



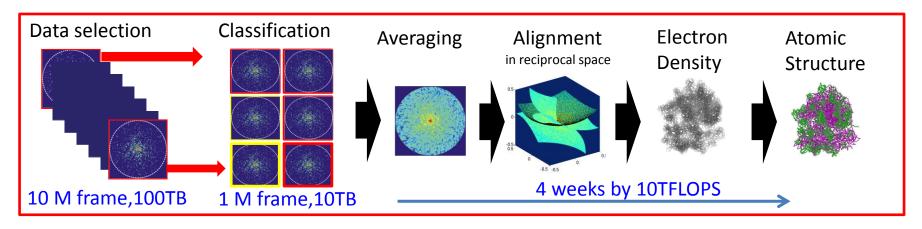


Full data analysis

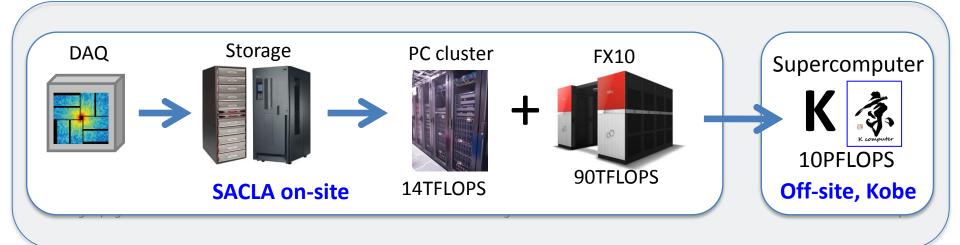




Full data analysis scheme



- Big data processing is indispensable for 3D analysis
 - $ightharpoonup 10^6$ frames for analysis $\Rightarrow 10^7$ frames for data taking (including junk).
 - 4 days data taking at 60Hz ~ 100TB storage size.
- Estimated CPU time for phase retrieval to get the atomic structure
 - 4 weeks by 10 TFLOPS PC cluster.
 - > 3 days by 100 TFLOPS supercomputer.
 - -> Experimentalist requires more computing power.





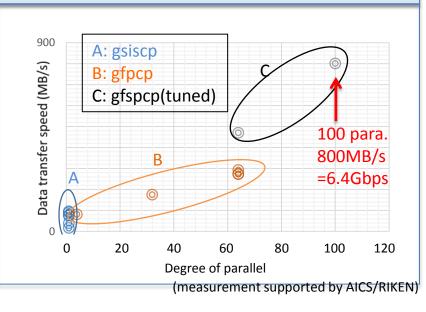
Data analysis plan using supercomputer 'K computer'

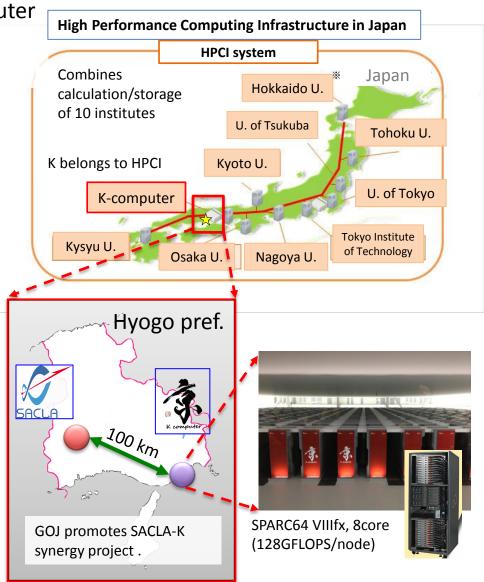
We just started feasibility study to use K computer for not only full data analysis, but also prompt

data analysis.

SACLA-K synergy

- We carried out a preliminary datatransfer and job submit from SACLA to K computer.
- We achieved 6.4Gbps bandwidth, which satisfy the experimental data rate.







- SACLA provides new opportunity to study complex targets using XFEL, especially, reveal 3-D structure of proteins.
- We developed DAQ and Analysis system for SACLA to satisfy experimental requirements:
 - 6 Gbps data rate
 - 120 TBytes data storage per protein sample
 - short experimental period (a few days cycle)
 - high computing power
 - 10 TFLOPS for run-by-run prompt analysis
 - 100 TFLOPS for full data analysis
 - ... and more computing power!
- We also started feasibility study of joint analysis using K computer, which is 10 PF supercomputer at Kobe.



backup