



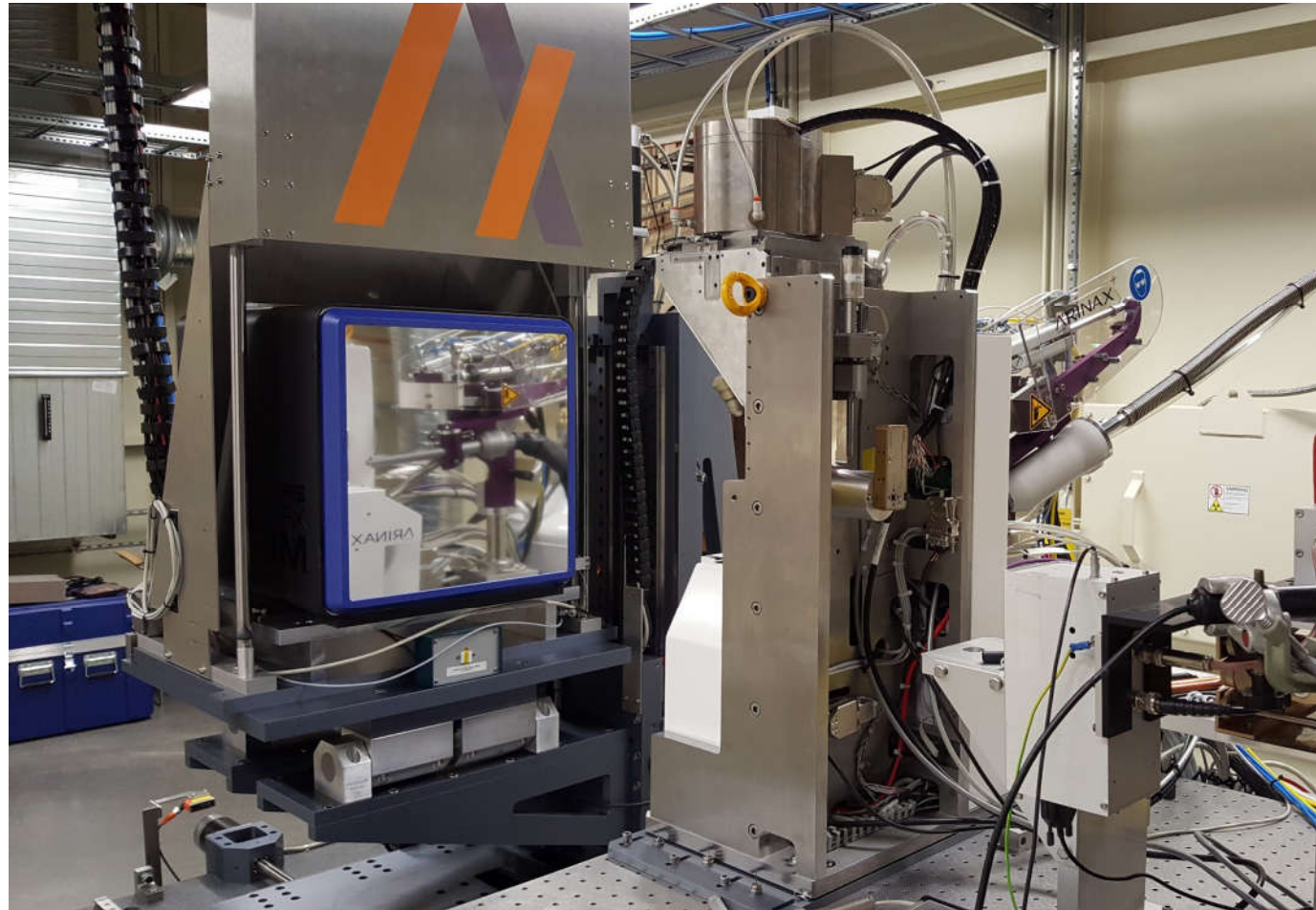
MXCuBE at MAX IV

Ph.D. Jie Nan

Jan. 16th, 2017

MX Beamline at MAX IV - BioMAX

- MAD Station, 5 – 25 keV
- In-vacuum undulator
- MD3 Micro-Diffractometer
 - Mini-kappa
 - Crystallization Plate holder
- Detector, Eiger 16M
- Cryojet5, HClab and REX
- Amptek fluorescence detector
- BCU
- ISARA, Sample Changer



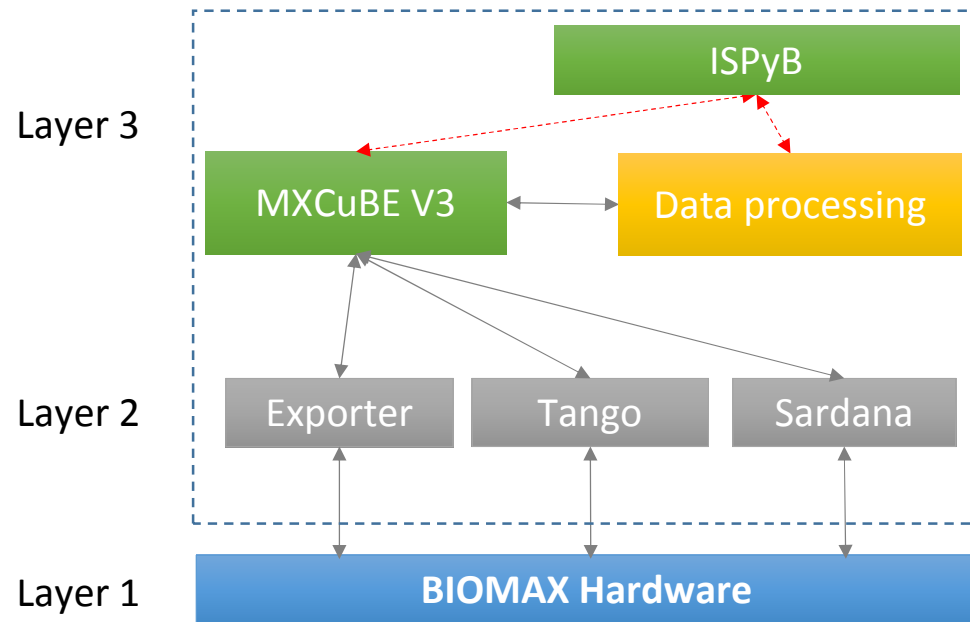
ISARA – Sample Changer

Four grippers

- Unipuck (Double)
- Unipuck (Single)
- SPINE (Single)
- Plate



BioMAX Software Overview



MXCuBE3 at BioMAX

- Sample View
 - 3-click centering,
 - Loop Centering
 - 2D Centring
- Data Collection
 - Std. Data collection
 - Characterization
 - Helical scan
 - Support queue
- Beamline/machine info.
- Processing

The screenshot displays the MXCuBE-3 control interface. On the left, there are control panels for Omega (10.16), Kappa (-0.5), Phi (0), Y (0.08), Z (0.21), Focus (-0.27), Samp-X (0.22), and Samp-Y (0.15). The top right shows beamline parameters: Energy: 12.66 keV, Resolution: 4112.333 Å, Transmission: 100%, and Detector Distance: 597.909 mm. The central image shows a sample with a yellow line indicating a scan path through points P1, L0, and P2. The right panel shows data collection settings and a parameters summary.

HardwareObjects

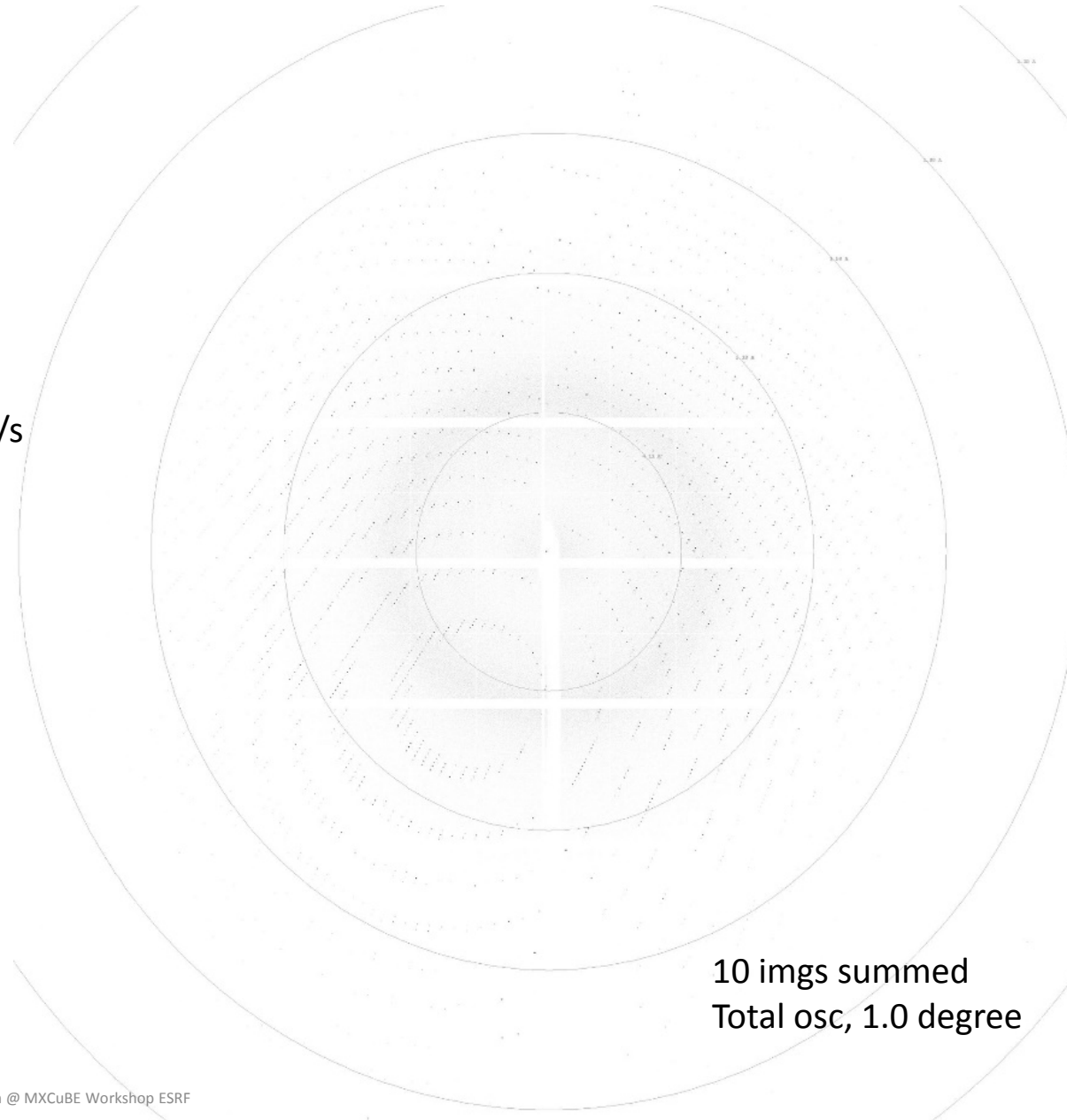
- 2.2 branch
- new hwobjs
 - BIOMAXEiger.py and BIOMAXEigerMockup.py
 - BIOMAXCollect.py (AbstractCollect.py)
 - BIOMAXMD3.py (GenericDiffractometer.py)
 - BIOMAXResolution.py
 - BIOMAXBeamInfo.py
 - BIOMAXAperture.py

Data Collection at BioMAX

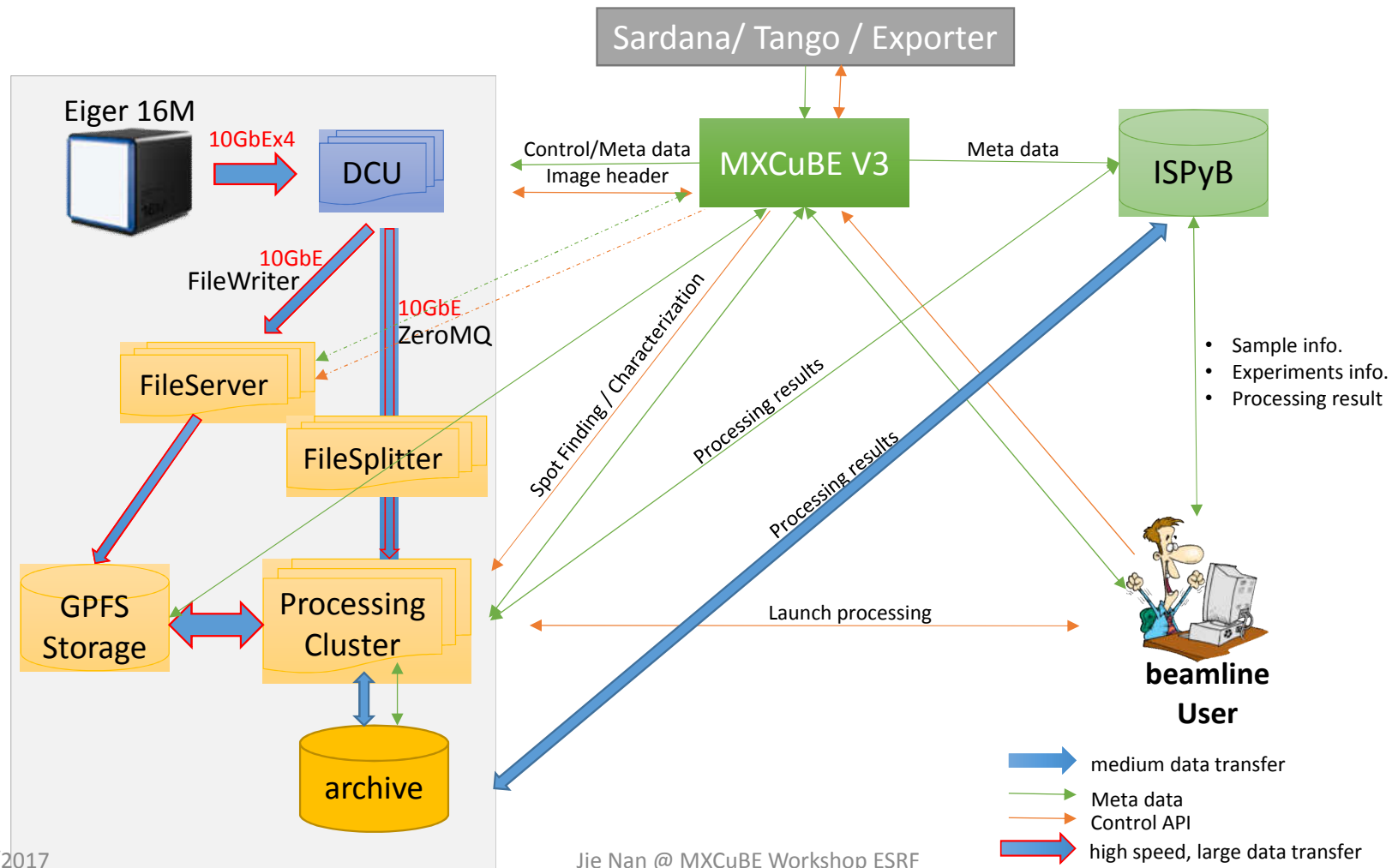
- Ring Current, 20-50 mA
- Beam size, 20 x 5 μm^2 (hor x ver)
- Estimated photon flux (sample): over 10^{11} ph/s
- Fixed energy: 12.7 keV
- User operation: Commissioning users
- Sample mount: manual

Thaumatin crystal

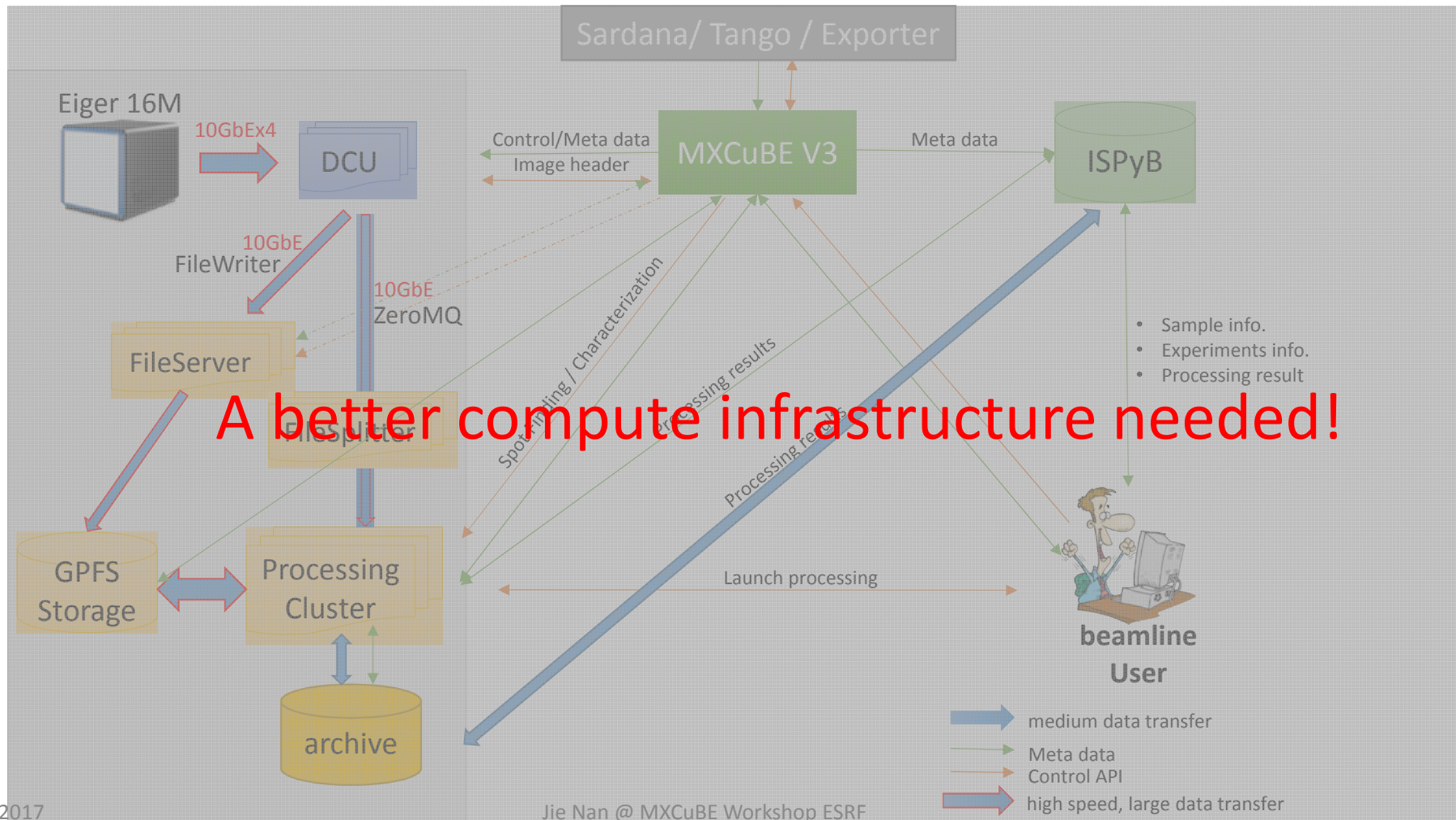
- Osc. 0.1 degree
- Exposure, 0.1 s
- Helical scan



BioMAX Data Flow Overview



BioMAX Data Flow Overview



Upgrade of Compute Infrastructure

Current Status

- 1 Gb/s to 10 Gb/s
- Buffer server at BioMAX
- GPFS storage
- HPC Cluster with 8 nodes (shared)

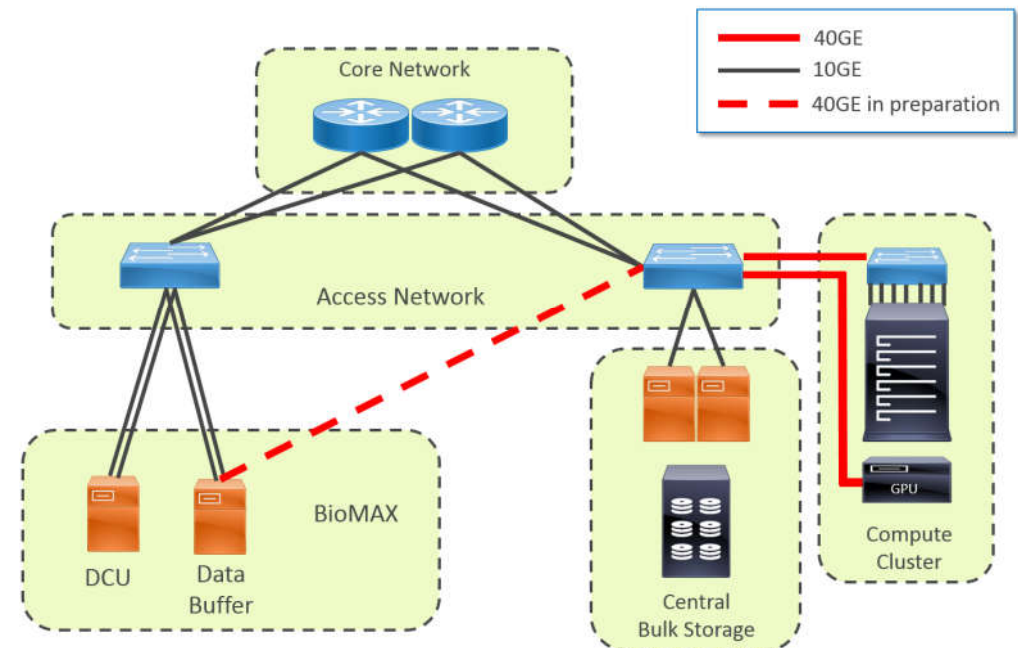
Further upgrade of 40GE based network

- Buffer Server to GPFS storage
- Buffer Server to Cluster

Computation

- New 8 nodes, 128 GB RAM, 20 cores per node
- Continuously grow

Blue Data Network: BioMAX to Central Resources



Future work

- Hardware integration
 - Sample changer
 - Fluorescence detector
 - Cryojet5, HClab and REX
- Experiments
 - Mesh scan
 - Element analysis & XANES
 - MAD data collection
 - Plate manipulation
 - RT collection (HClab) & crystal freezing (REX)
 - New collection protocols
- Integration with ISPyB
- Data processing
 - Spot finding
 - Reduction

Acknowledgement

MX Group

Uwe Mueller
Thomas Ursby
Johan Unge
Roberto Appio
Christopher Ward
Ross Friel

Marjolein Thunnissen

ESRF

Matias Guijarro
Marcus Oscarsson
Daniele De Sanctis
Antonia Beteva

KITS Group

Darren Spruce
Mikel Eguiraun
Fredrik Bolmsten
Antonio Milán-Otero
Artur Barczyk
Zdenek Matej

many others from MAX IV

Vicente Rey Bakaikoa