



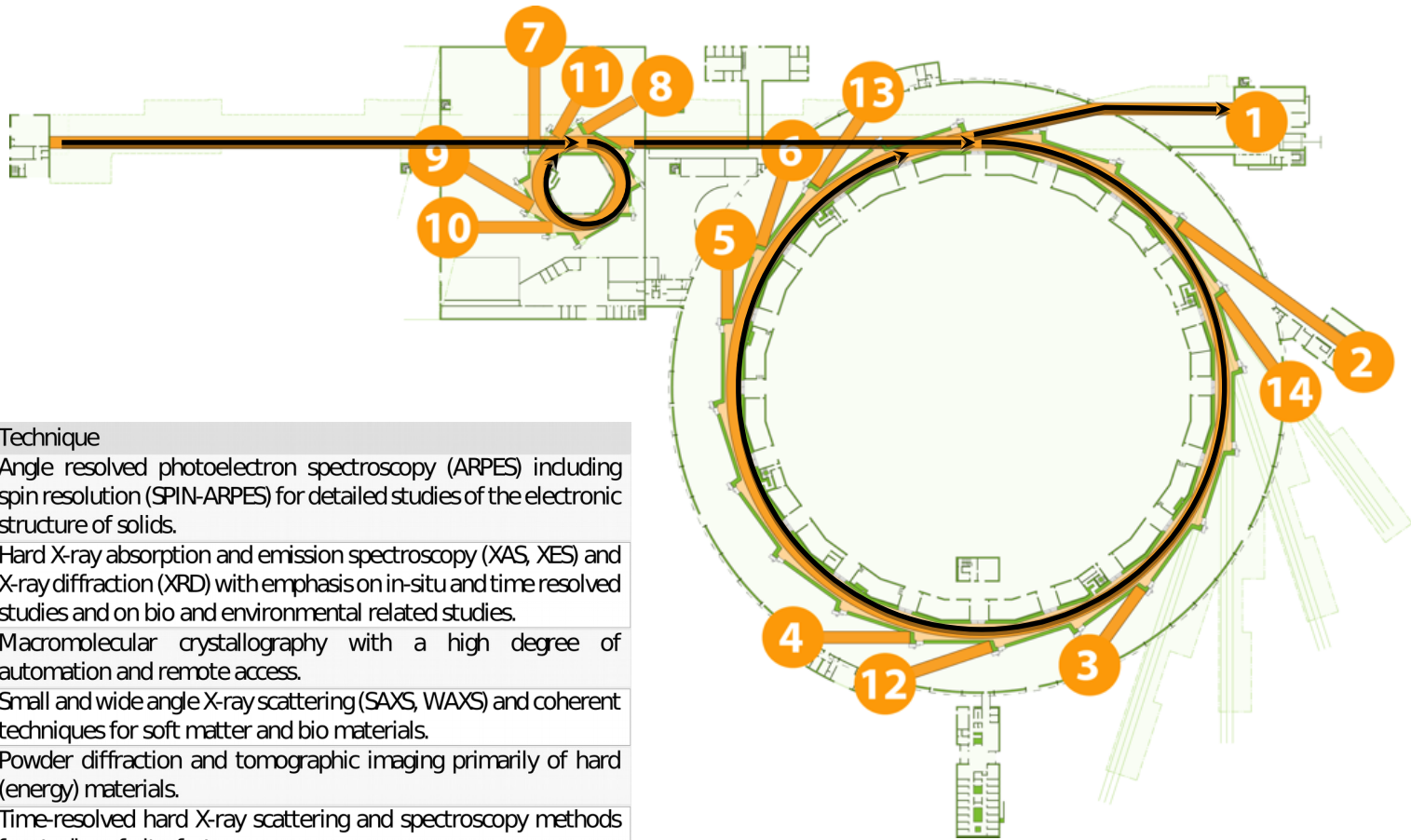
MXCuBE @ MAX IV: Status Report

Johan Unge

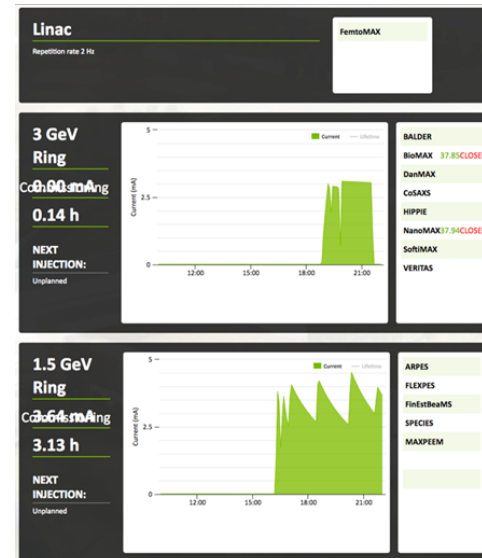
January 31, 2018



1 Linac 2 Rings 15 beamlines



Beamline	Accelerator	Technique
ARPES	7 1.5 GeV	Angle resolved photoelectron spectroscopy (ARPES) including spin resolution (SPIN-ARPES) for detailed studies of the electronic structure of solids.
Balder ★	3 3.0 GeV	Hard X-ray absorption and emission spectroscopy (XAS, XES) and X-ray diffraction (XRD) with emphasis on in-situ and time resolved studies and on bio and environmental related studies.
BioMAX ★	4 3.0 GeV	Macromolecular crystallography with a high degree of automation and remote access.
CoSAXS	12 3.0 GeV	Small and wide angle X-ray scattering (SAXS, WAXS) and coherent techniques for soft matter and bio materials.
DanMAX	14 3.0 GeV	Powder diffraction and tomographic imaging primarily of hard (energy) materials.
FemtoMAX ★	1 Linac	Time-resolved hard X-ray scattering and spectroscopy methods for studies of ultrafast processes
FinEstBeaMS	8 1.5 GeV	Electron spectroscopies and luminescence methods for studies of low density matter and solids.
FlexPES	11 1.5 GeV	Soft X-ray spectroscopies for studies of low density matter and solids.
HIPPIE ★	6 3.0 GeV	Near ambient pressure photoelectron spectroscopy on solids and liquids.
MAXPEEM	10 1.5 GeV	Aberration corrected photoelectron microscopy for investigation of surfaces and interfaces.
NanoMAX ★	2 3.0 GeV	Imaging with spectroscopic and structural contrast techniques and nanometre resolution.
SoftiMAX	13 3.0 GeV	Scanning transmission X-ray microscopy and coherent imaging methods.
SPECIES	9 1.5 GeV	Resonant inelastic X-ray scattering (RIXS) with high resolving power and near ambient pressure photoemission.
VERITAS ★	5 3.0 GeV	Resonant inelastic X-ray scattering (RIXS) with unique resolving power and high spatial resolution.

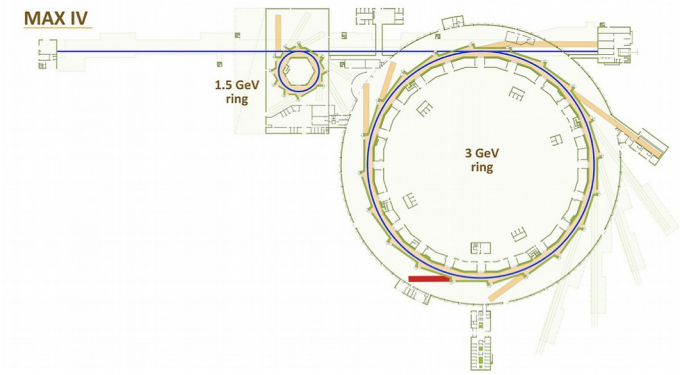


Linac:
Injector 1.5 & 3 GeV & SPF
Injects both rings & delivers light to FemtoMAX

3 GeV:
200 mA, >5 Ah
≈ (3400±30) pm·rad
≈ 8.5 mA single bunch
Delivers light to users

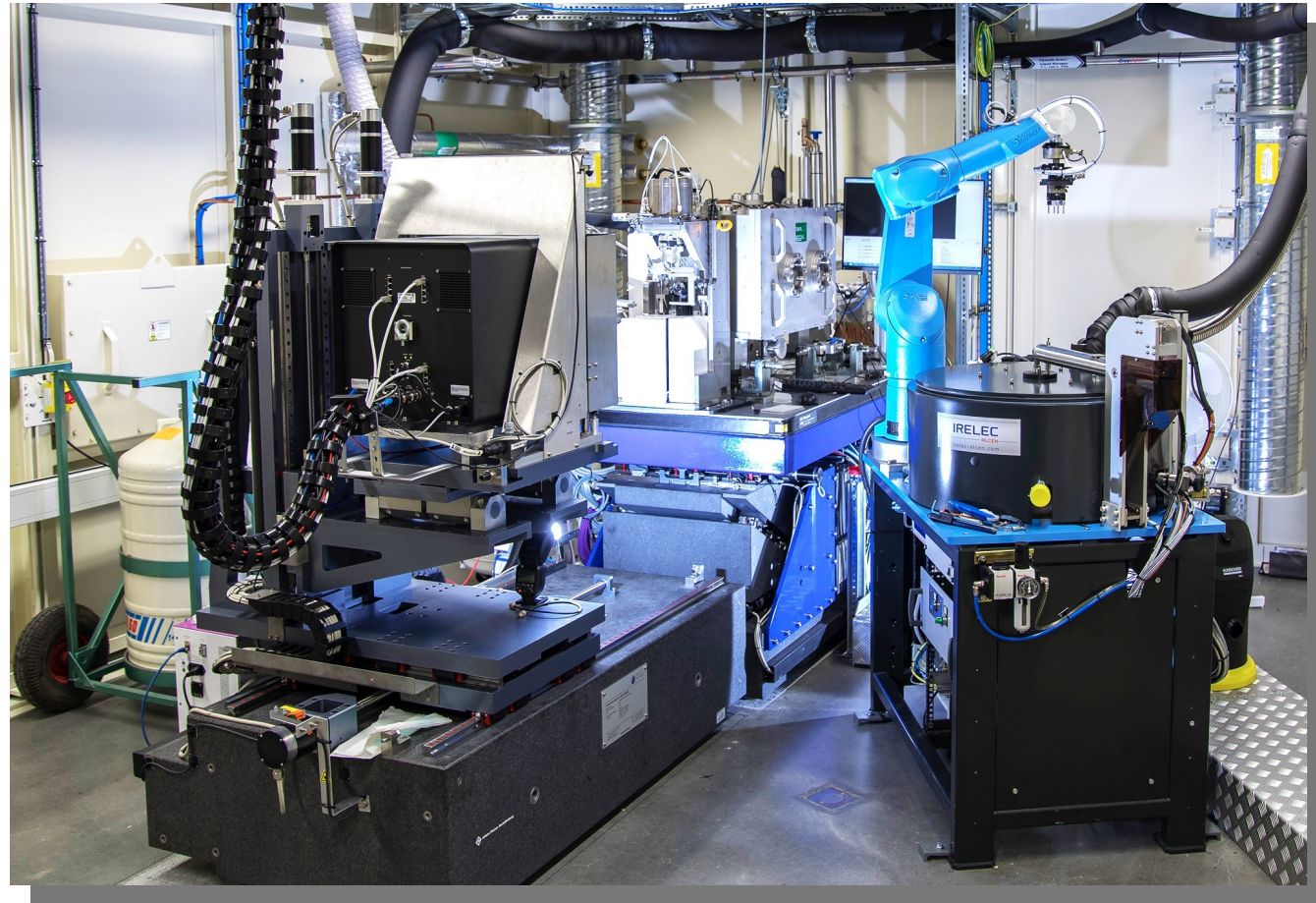
1.5 GeV:
< 200 mA
Commissioning

BioMAX experimental station



Experimental environment

- MD3 micro-diffractometer
- Eiger 16M hybrid pixel detector
- ISARA sample changer
- HCLab humidifier
- Beam Condition Unit incl. XBPM
- Amptek fluorescence detector

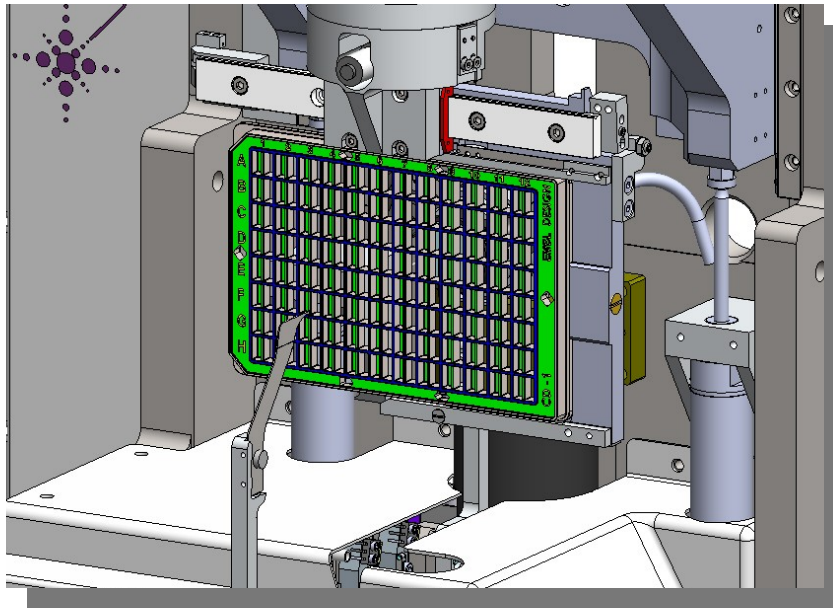


BioMAX experimental station

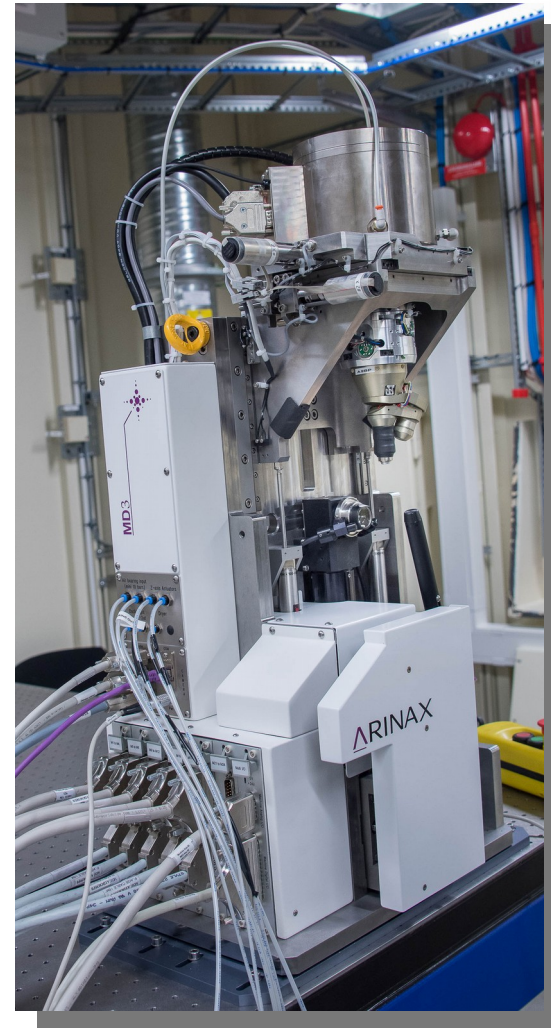
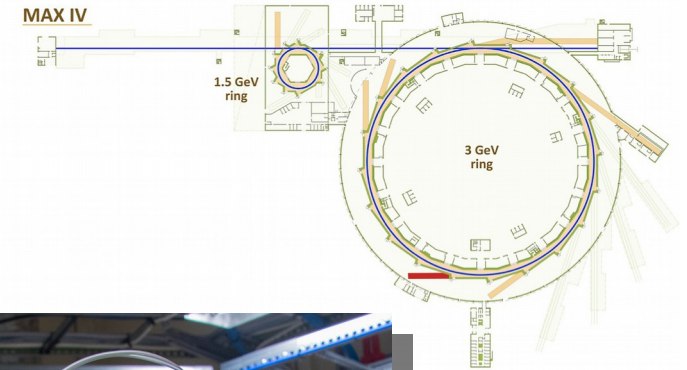
Goniometry

MD3 diffractometer:

- Ultra precise Omega-axis
 - Sphere-of-confusion $r=100$ nm
 - Max. speed 800 deg/sec
- MK3 Mini-kappa goniometer
- Improved on-axis-microscope
- Crystallization plate holder



MD3 plate holder

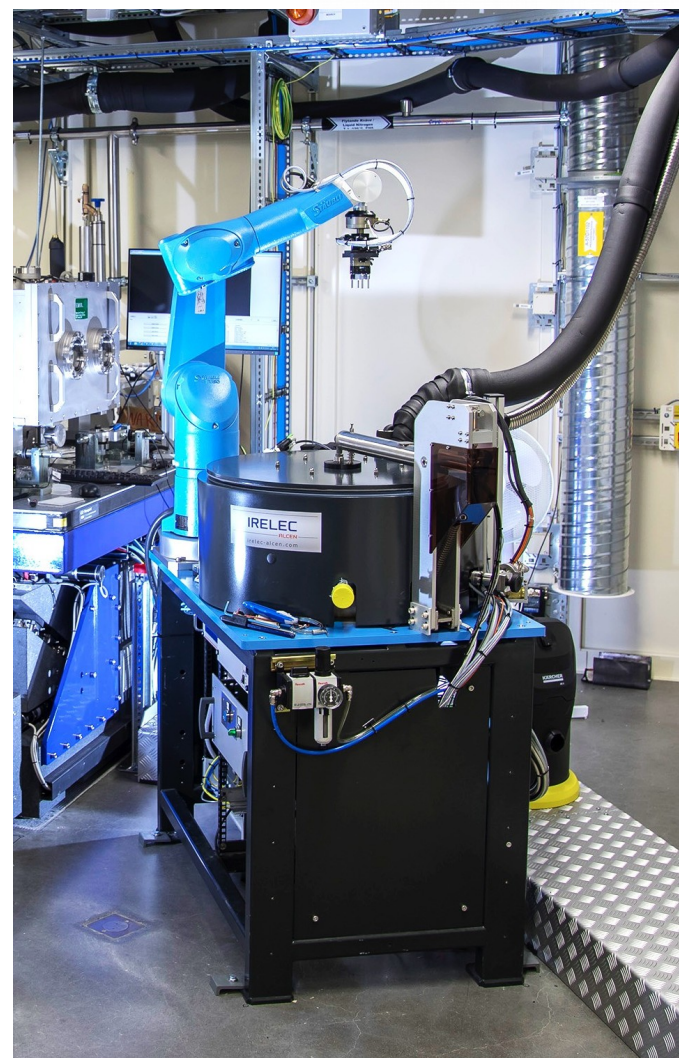
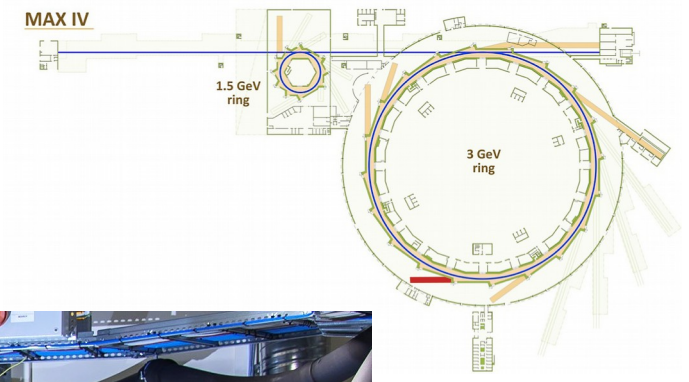
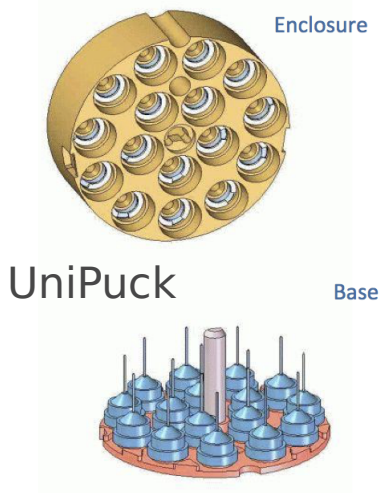


MD3 at BioMAX experimental hutch

Sample changer

ISARA sample changer:

- Dewar capacity
 - 100 samples in SPINE pucks
 - 304 samples in UniPuck
- 4 crystallization plates
- 18 sec sample transfer time

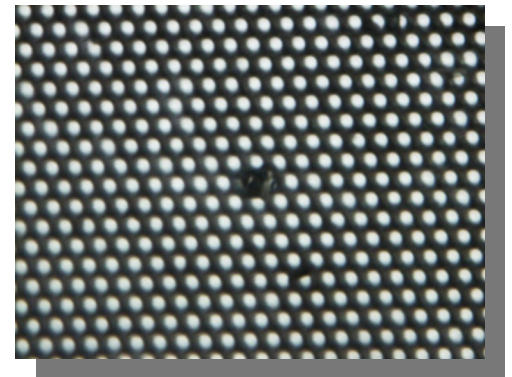
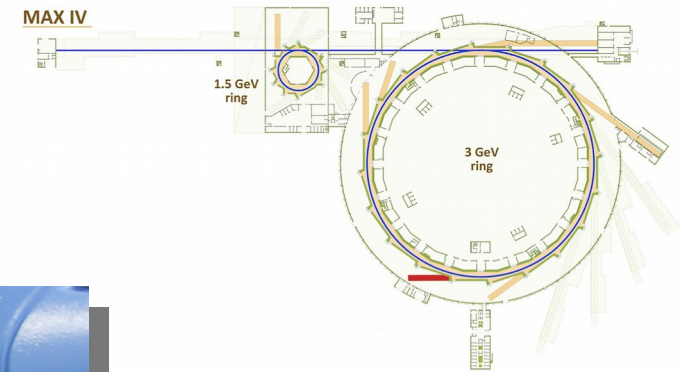


Status

- Ice formation issues with new design
- Currently Unipuck is working
- Integrated into mxcube3, operation, maintenance page and hwobj

ISARA sample changer (IRELEC)

HCLab - Room temperature



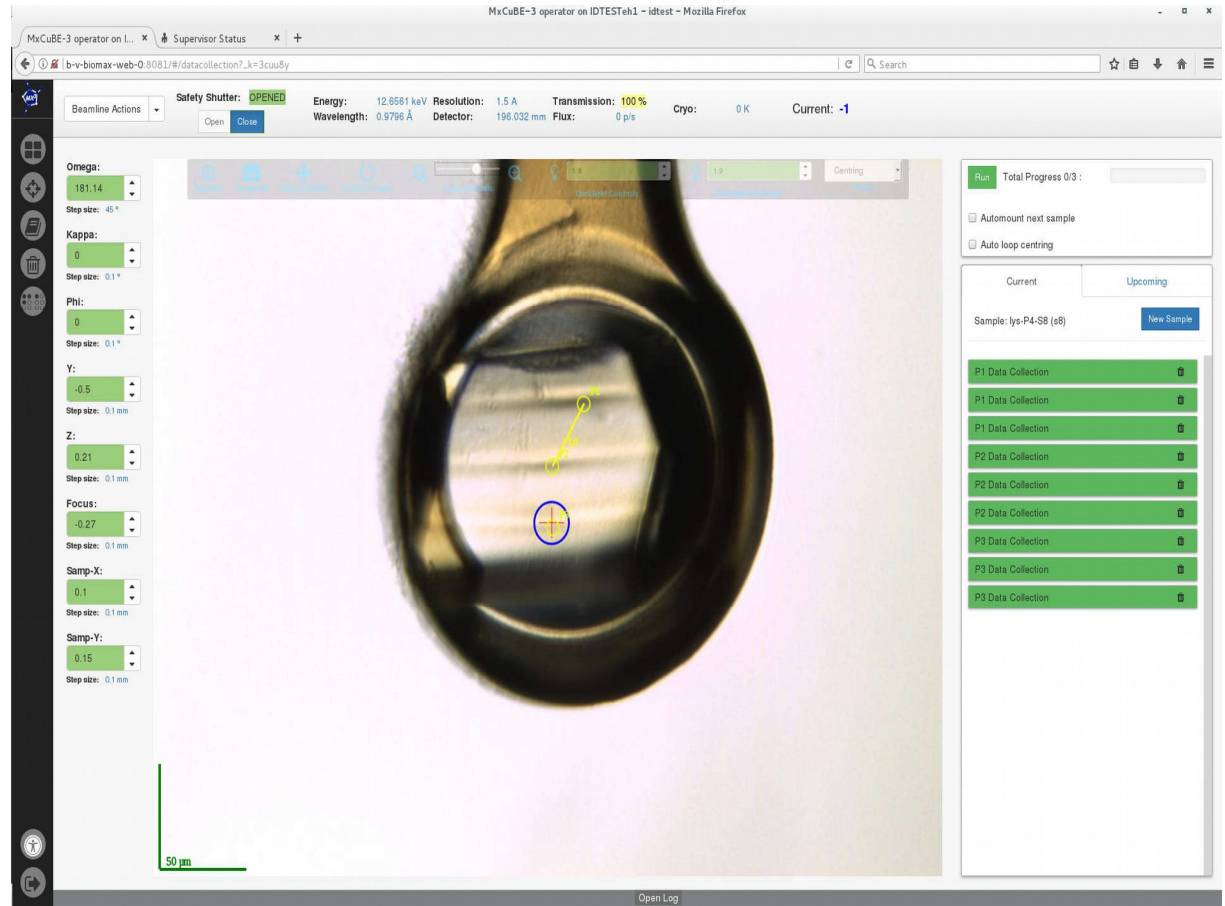
HCLab facility operational

First Si chip operation at BioMAX,
(B. Vestergaard, UC)

MXCuBE3

MXCuBE3 data collection view

- *Current Status, major features*
 - Sample Centering
 - Standard Oscillation
 - Characterization
 - Helical Scan
 - Sample Changer
 - ISARA adaptation for Tango device and HWO
 - Mesh Scan
 - In Situ
- *In progress and todo*
 - Stability maturation
 - XRF, XANES
 - Parallel processing for spot finding for MESH scan
 - ISPyB and SDM integration

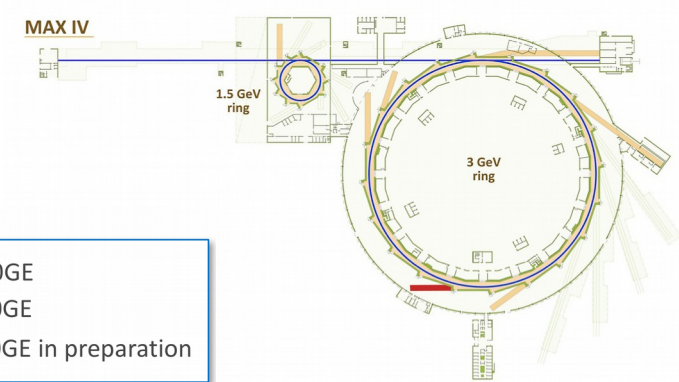


- *Advantage*
 - Joint collaboration with ESRF
- *Challenges*

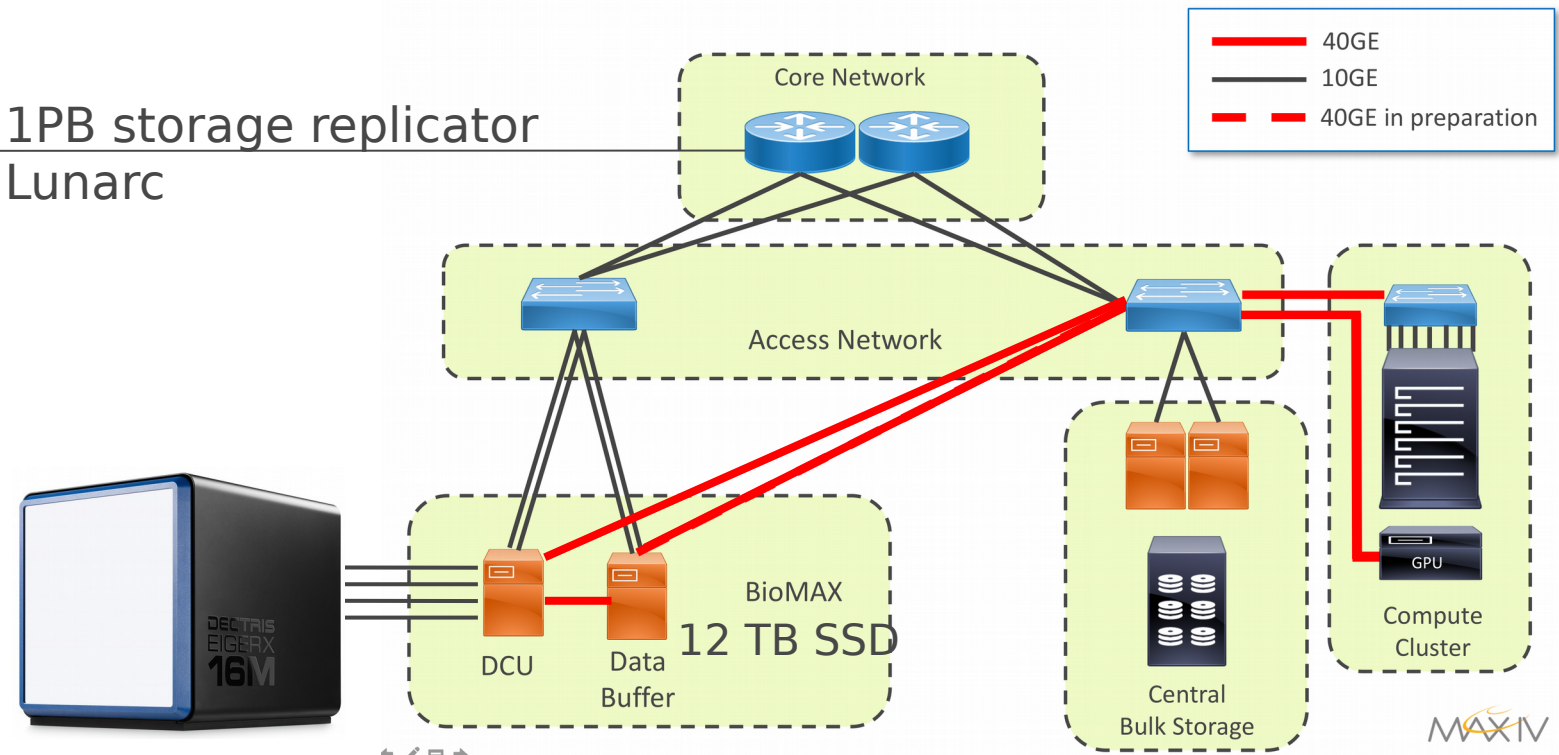
Software stability, user-friendliness and integration with of various hardware + software

IT environment

Eiger 16M IT-environment



1PB storage replicator
Lunarc



~ 1100 CPU cores

60 nodes 2017
25 dedicated to BioMAX

Courtesy of Artur Barczyk

250 TB installed
GPFS file system

- Alternative data retrieval via Globus (www.globus.org)
- DUO (proposal management) in place and will be used soon
- Account and data management system (SDM) → talk by Fredrik

Data reduction and performance

- Three pipelines currently used
 - fast_dp, autoPROC, BioMAX_Pipeline
- System to install all MX software on cluster, PreSTO, collaboration with National Supercomputer Centre and Lunarc.
 - Thinlinc to create simple interface to cluster
 - XDS and central GPFS file system? XDS makes many small I/O operations...testing alternatives

MAX IV, GPFS system (1)	Lunarc, Same blades type (1)	4 Nodes, NFS mount 8 XDS-jobs 24 processes (2)	8 Nodes NFS mount 16 XDS-jobs 24 processes (2)
134.0 sec	52 sec	56.2 sec	45.6 sec

(1) 900 images, thaumatin, 0.1 deg/image, (2) 900 images, Dectris data

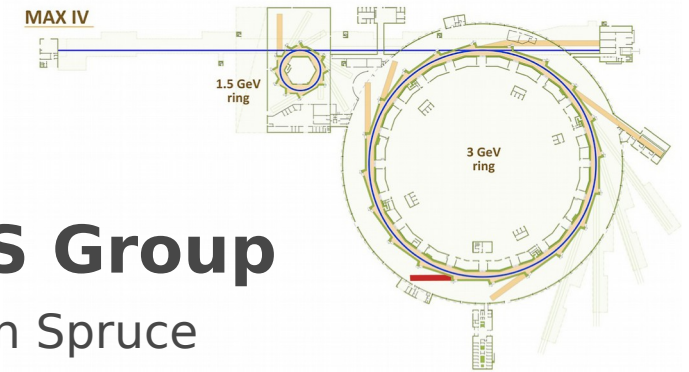
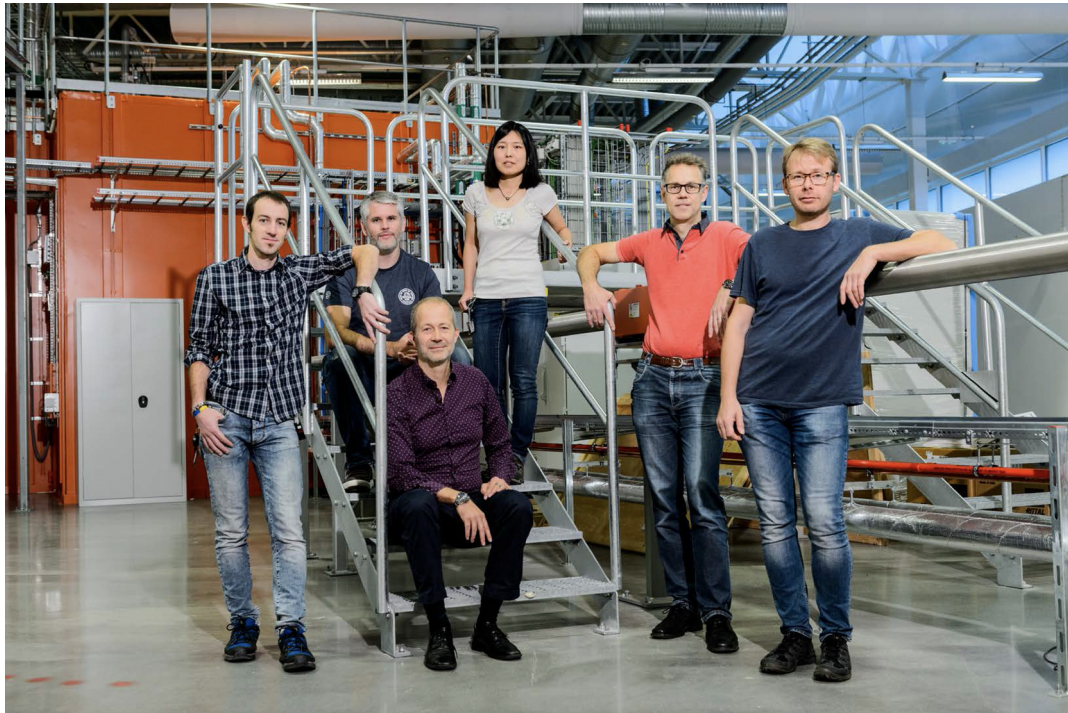
Outlook

- Remote operation using web based MXCuBE. This will be critical during 2018. Involves MAXIV IT systems.
- ISARA sample changer: all functionalities and a lot of testing -> ready to users!
- Remaining equipment to integrate...
- Extend beamline capabilities (energy tuneability, beam stability feedback systems, MESH scan, XRF, minikappa...

Acknowledgement

MX Group

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KITS Group

Darren Spruce
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Antonio Milán-Otero
Artur Barczyk
Zdenek Matej

User Office

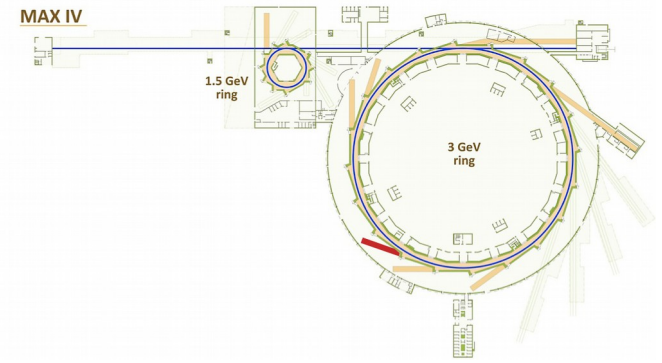
Marjolein Thunnissen

ESRF

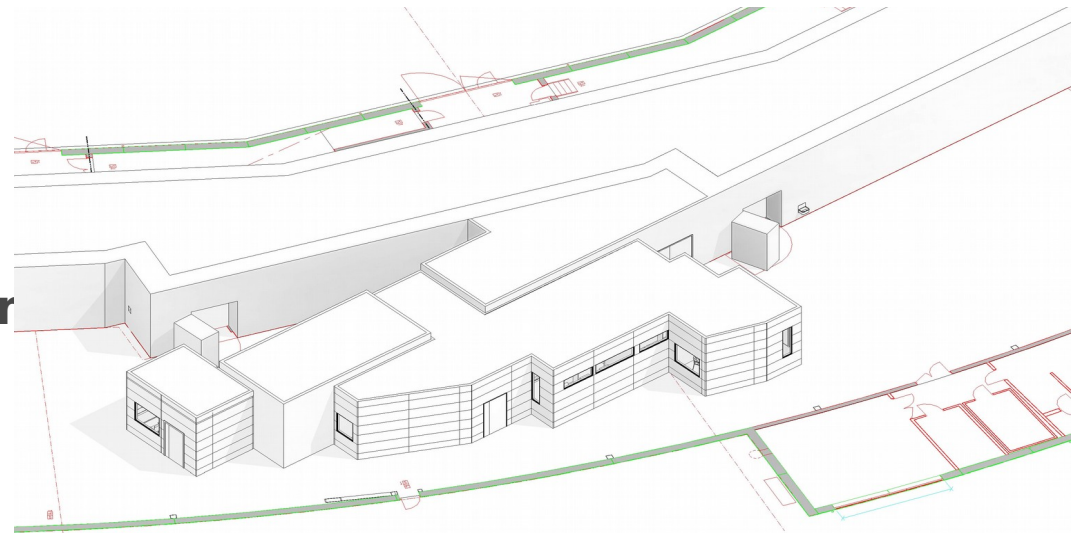
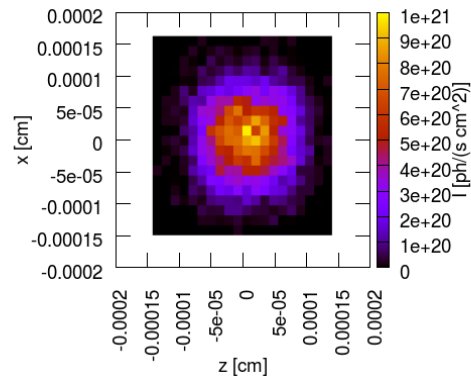
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MicroMAX



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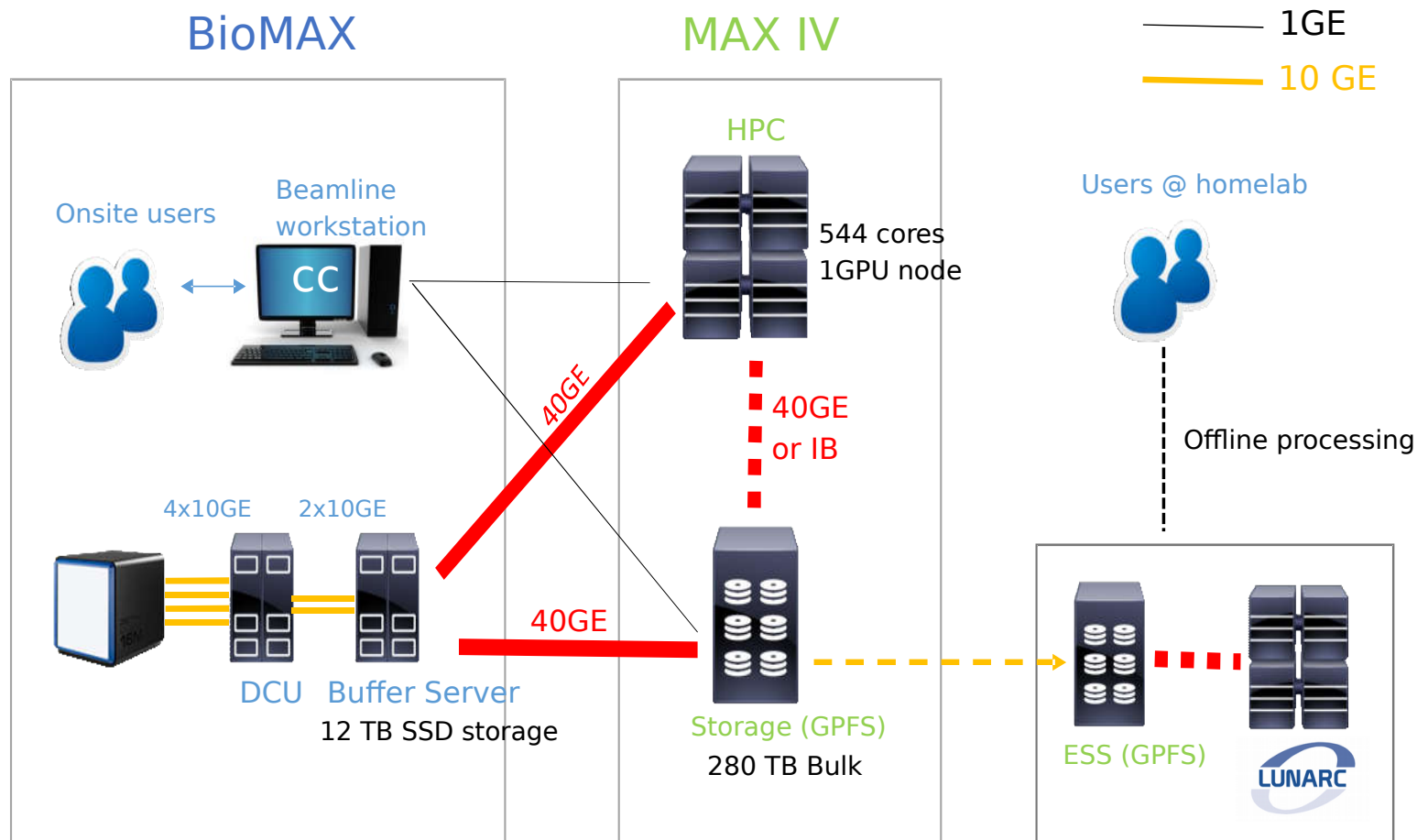


Plans for the second MX beam

- Microfocus beamline
- **1 x 0.7** μm^2 beam at sample
- Photon flux **10^{13} - 10^{15}** phot/sec
- Traditional setup (goniometry, sample environment)
- Exploratory setup (serial crystallography, fixed target single shot)
- Optimal source for most demanding projects (large complexes, membrane proteins)

Status: funded by Novo Nordisk Foundation

BioMAX - IT Infrastructure



Data reduction and performance

- Automatic processing

- Fast_dp
- autoPROC
- Biomax_pipeline

	HPC - 4 nodes (24 cores, CPU E5-2650 v4 @ 2.20GHz)	HPC - 8 nodes (48 cores, CPU E5-2650 v4 @ 2.20GHz)	BufferServer (5-cores, CPU E5-2643 v4 @ 3.4GHz)	Key32 cpus, HT	SOLEIL - 4 nodes (16 cores, E5-2690 @ 2.70 GHz)
Time*	53s	36s	136s	120 s	76.6 s

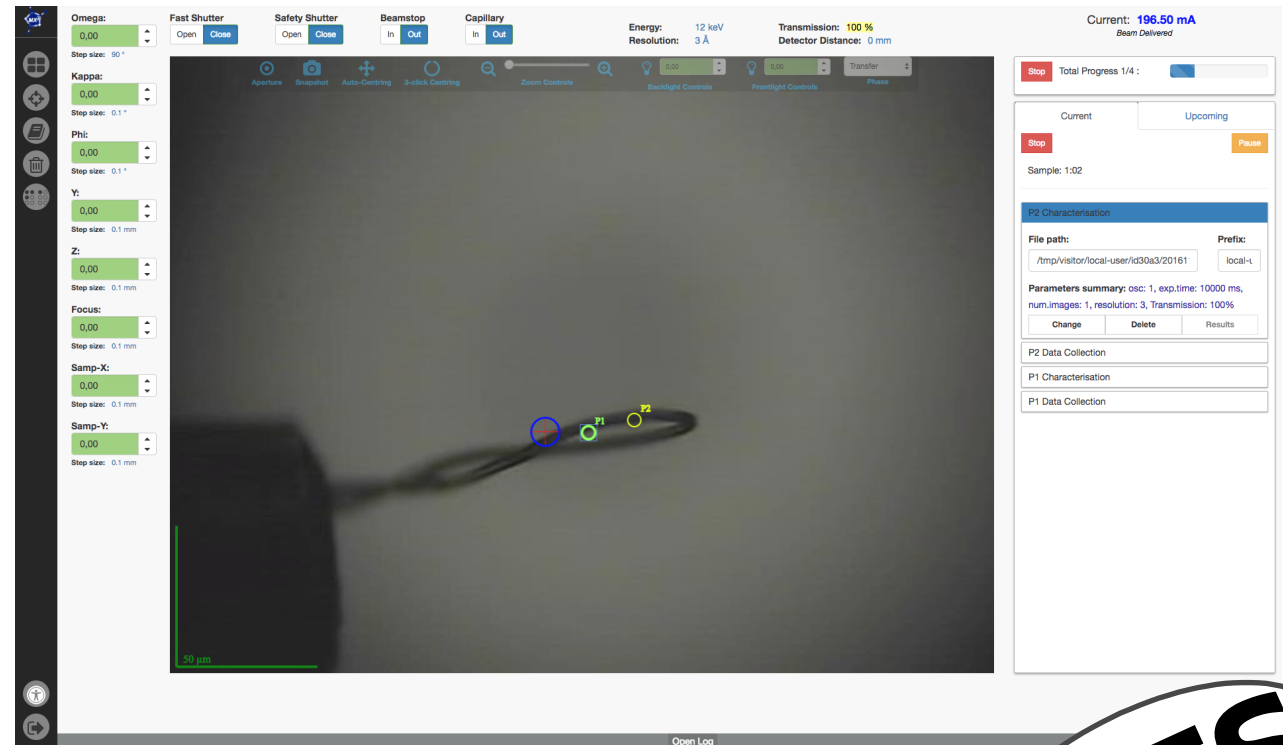
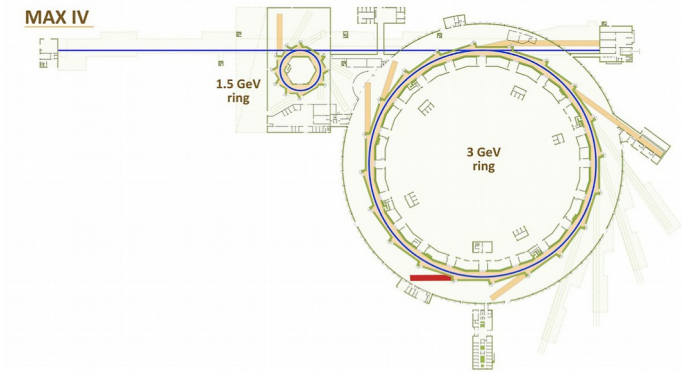
EIGER_16M_Nov2015.tar.bz2, 900 images in 9 containers, 0.1 osc, from Dectris website

- Run XDS with SLURM on HPC

https://github.com/JieNanMAXIV/xds_slurm

MXCuBE3

- *Current Status, major features*
 - Sample Centering
 - Standard Oscillation
 - Characterization
 - Helical Scan
 - Sample Changer
 - Mesh Scan
- *Todo*
 - Stability maturation
 - XRF, XANES
- *Advantage*
 - Joint collaboration with ESRF
- *Challenges*
 - Integration with of various hardware, software,
 - Software stability and user-friendliness



MXCuBE3 data collection view

KITS

MAX IV

MXCuBE3 at BioMAX

MxCuBE-3 operator on IDTESTeh1 - idtest - Mozilla Firefox

b-v-biomax-web-0.8081/#/datacollection?_k=3cuu8y

Safety Shutter: **OPENED**

Energy: 12.6581 keV Resolution: 1.5 Å Transmission: 100 %
Wavelength: 0.9796 Å Detector: 196.032 mm Flux: 0 p/s Cryo: 0 K Current: -1

Omega: 181.14
Step size: 45 °

Kappa: 0
Step size: 0.1 °

Phi: 0
Step size: 0.1 °

Y: -0.5
Step size: 0.1 mm

Z: 0.21
Step size: 0.1 mm

Focus: -0.27
Step size: 0.1 mm

Samp-X: 0.1
Step size: 0.1 mm

Samp-Y: 0.15
Step size: 0.1 mm

50 µm

Manual mounting
Over 40 samples, 500GB / 8h shift

Run Total Progress 0/3 :
 Automount next sample
 Auto loop centring

Current Upcoming
Sample: lys-P4-S8 (s8) [New Sample](#)

P1 Data Collection
P1 Data Collection
P1 Data Collection
P2 Data Collection
P2 Data Collection
P2 Data Collection
P3 Data Collection
P3 Data Collection
P3 Data Collection

Open Log

Main features

- Sample Video
 - 3 click Centering
 - 2D Centring
 - Auto-Loop Centri
- MD3 controls
 - Motors
 - Zoom
 - Light
 - phase
- Data Collection
 - Standard collecti
 - Characterization
 - Helical Scan
- Processing
 - Launched on HPC
- Queue