

# Unified EM setup: the benefits of SerialEM for facilities

Wim Hagen



# CBB unit / EM core facility



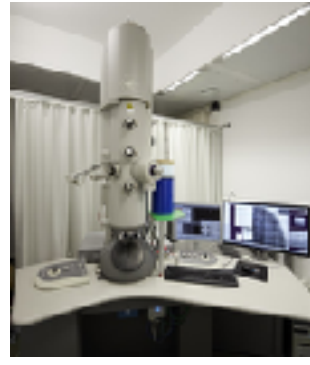
Morgagni  
MegaView



CM120  
SIS



JEOL 2100+  
Ruby camera



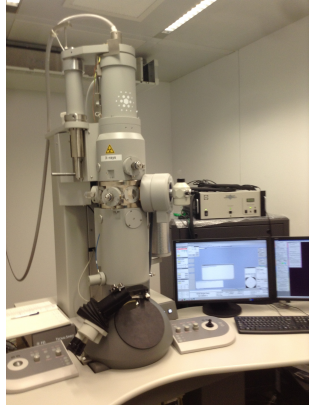
Tecnai F30  
Gatan OneView



Zeiss Crossbeam

- Yannick Schwab, Team Leader and Head of Electron Microscopy Core Facility
- Mandy Boermel, Scientific Officer
- Pedro Machado, Technical Officer
- Rachel Mellwig, EM Facility Operations Manager
- Giulia Mizzon, Scientific Officer
- Paolo Ronchi, Scientist in EM Facility
- Martin Schorb, Application Engineer

# SCB unit / cryo-EM service platform



Tecnai 12  
US 4000



Polaris  
Falcon 2  
GIF2002



Arctica  
Falcon 3



Krios  
Quantum K2

- Felix Weis, Cryo-Electron Microscopy Specialist
- Wim Hagen, Senior Engineer in Electron Microscopy

One third guests over iNext and Direct Access.



CORE FACILITIES

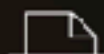
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CRYO-ELECTRON MICROSCOPY SERVICE  
PLATFORM

# Microscopy



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### CRYO-ELECTRON MICROSCOPY SERVICE PLATFORM

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RECENT PUBLICATIONS

MEMBERS

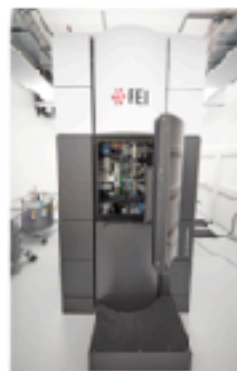
CONTACT

EMBL Heidelberg hosts a cryo-electron microscopy (cryo-EM) service platform, available for use by external scientists with both single particle and tomography projects.

The platform gives access to state-of-the-art cryo-EM equipment for structure determination projects using the latest technology and methods. The platform hosts a FEI Titan Krios G3 equipped with a phase plate and a Quantum-X2 camera and a 200 kV FEI Talos Arctica with an electron coating Falcon 3 camera. Both microscopes are equipped with an automatic sample loading system and configured for automatic data acquisition.

Experts are on hand to help and support researchers during microscope handling, data acquisition and optimisation of imaging conditions.

Thanks to a continuous test program with Thermo Fisher, we ensure the microscopes have the latest hardware and software features.



Titan Krios G3 equipped with phase plate and Quantum-X2 camera.





# EMBL Imaging Center



Why SerialEM?

## Tecnai 12

- Windows XP, US4000, SerialEM.
- Windows 7, US4000, SerialEM.

## Polara

- Windows 2000, GIF2002/film, SerialEM, UCSF Tomo.
- Windows XP, GIF2002/US4000, SerialEM, EPU, FEI Tomo.
- Windows 7, GIF2002/Falcon 2, SerialEM, ~~EPU, FEI Tomo.~~

## Talos Arctica

- Windows 7, Falcon 3, SerialEM, EPU.

## Krios

- Windows XP, GIF2002/US4000/film, SerialEM.
- Windows XP, GIF2002/US4000/film, SerialEM, EPU, FEI Tomo.
- Windows XP, GIF2002/Falcon 2, SerialEM, EPU, FEI Tomo.
- Windows XP, Quantum K2, SerialEM, ~~EPU, FEI Tomo.~~
- Windows 7, Quantum K2, SerialEM, EPU, FEI Tomo.
- 2019: Quantum K3, SerialEM -> EPU, FEI Tomo?



## Latest FEI EPU/Tomo

- Only Thermo Fischer microscopes with supported camera's on Windows 7 software.

## Latest Gatan Latitude(S)

- Only Gatan camera's supported by GMS3 Windows 7 software, regardless of microscope type.

## Leginon

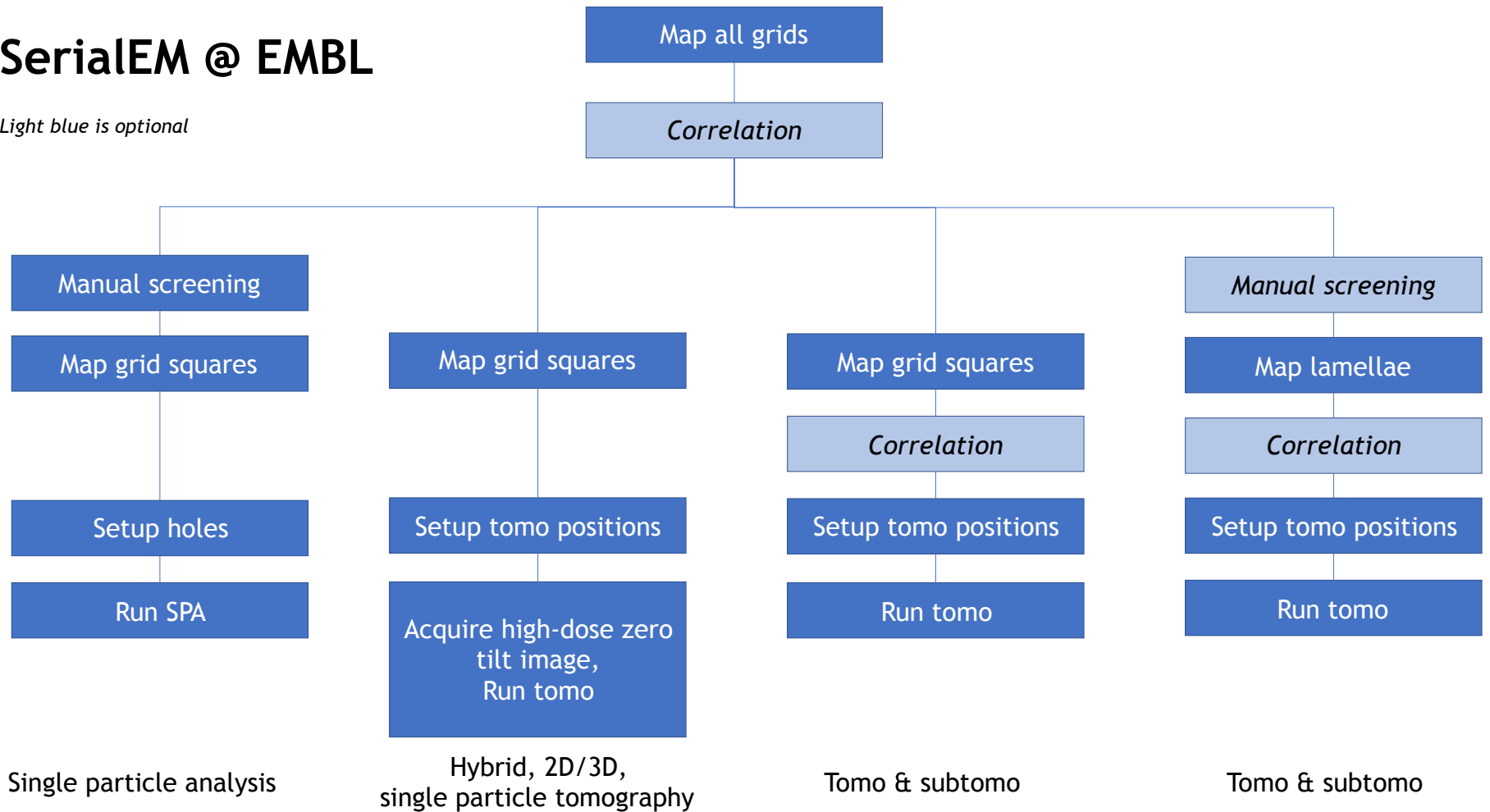
### SerialEM

- Windows 2000, Windows XP, Windows 7
- Thermo Fischer Tecnai, Titan, Talos, Glacios.
- JEOL
- Hitachi
  
- Gatan
- TVIPS
- FEI
- AMT
- DirectElectron
- EMSIS (formerly Olympus)
- JEOL Ruby
- STEM from Gatan's DigiScan or Thermo Fischer's STEM imaging.

**FEI Maps?????**

# SerialEM @ EMBL

*Light blue is optional*



# Any session

- Load grids, map all grids in lowest usable LM mode -> email.

## SPA

- Screen for good squares.
- Select squares/lamellae, map montage in SA mode -> email.
- Setup acquisition.

## Tomo

- Select squares/lamellae, map montage in SA mode -> email.
- Search for targets, setup acquisition.

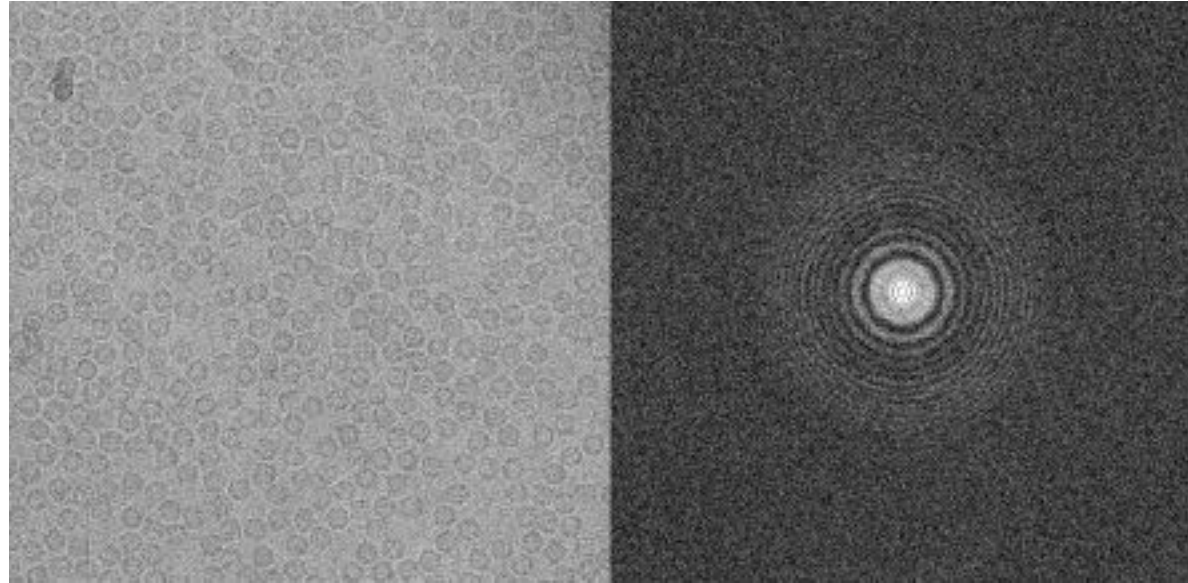
# SPA acquisition

Gatan pc SerialEM:

- Post-actions.
- EarlyReturnNextShot.
- Save uncorrected LZW compressed tif.

Support pc with GPU:

- IMOD FrameWatcher.
- IMOD AlignFrames.
- Data stored on EMBL network.



## Raw Data

SAVE SETTINGS | LOAD SETTINGS

### Input

Input: W2000003\_01m000001-1.mrc  
Pixel Size: 0.0400/0.0400 Å @ 0.03°  
Bin: 3.00x (0.0400 Npix)  
Mode: [2D \(4/1/Frame\)](#)

### Preprocessing

Correct gain using [LSD \(see reference\)](#)

High X axis  High Y axis  Transpose

### CTF

Window: 1024 px Range: 0.00-0.40 Hz  Use Minus Sum  
Average: 1000 Hz Cp: 0.00 mm Cp: 0.00 mm  
Amplitude: 0.07 Bk Amplitude: 0.00 AD: 0.00 eV  
Default: 0.0-0.5 μm  Phase Shift

### Motion

Window: 200-200 px Weight: 0.00 = 0.00 Å²

### Model



### Pick Particles

Use [DockerHub](#): 20200320

Input: [1.0](#) & [0.0](#) particles per micrograph

Maintain a minimum distance of 0.4 Å from

Green: 100 particles, 0.000 Npix,  Green:  Normal

### Output

skip first 0 set of frames

Average  
 Decimated average strength = 100, offset = 1.00  
 Rigid: each, collapse every 5 frames

# STOP PROCESSING

## Overview | LouverSpace | RealSpace

[REPORT MICROGRAPH LIST](#) | [AGONY PARTICLES \(LOCAL\)](#) | [EXPLORE PARTICLES](#) | [IMPORT PARTICLE COORDINATES](#) | [MATCH TIMESTAMPS](#) | [EXPORT BOXSET \(COMPLEX\)](#)

### Processing Status



### Acquisition (see up to 4.0 c)



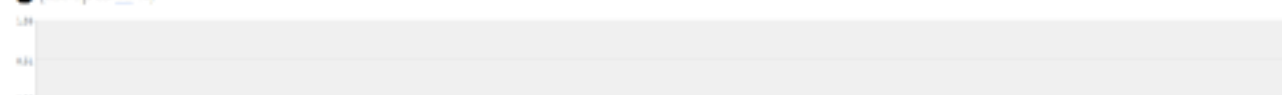
### Defocus (see 0.00-5.50 μm) — average CTF: 0.00



### Estimated resolution (see better than 10.0 Å)



### (see up to 50%)



# Tomo



## Journal of Structural Biology

Volume 197, Issue 2, February 2017, Pages 191-198



### Implementation of a cryo-electron tomography tilt-scheme optimized for high resolution subtomogram averaging

Wim J.H. Hagen, William Wan, John A.G. Briggs  

 **Show more**

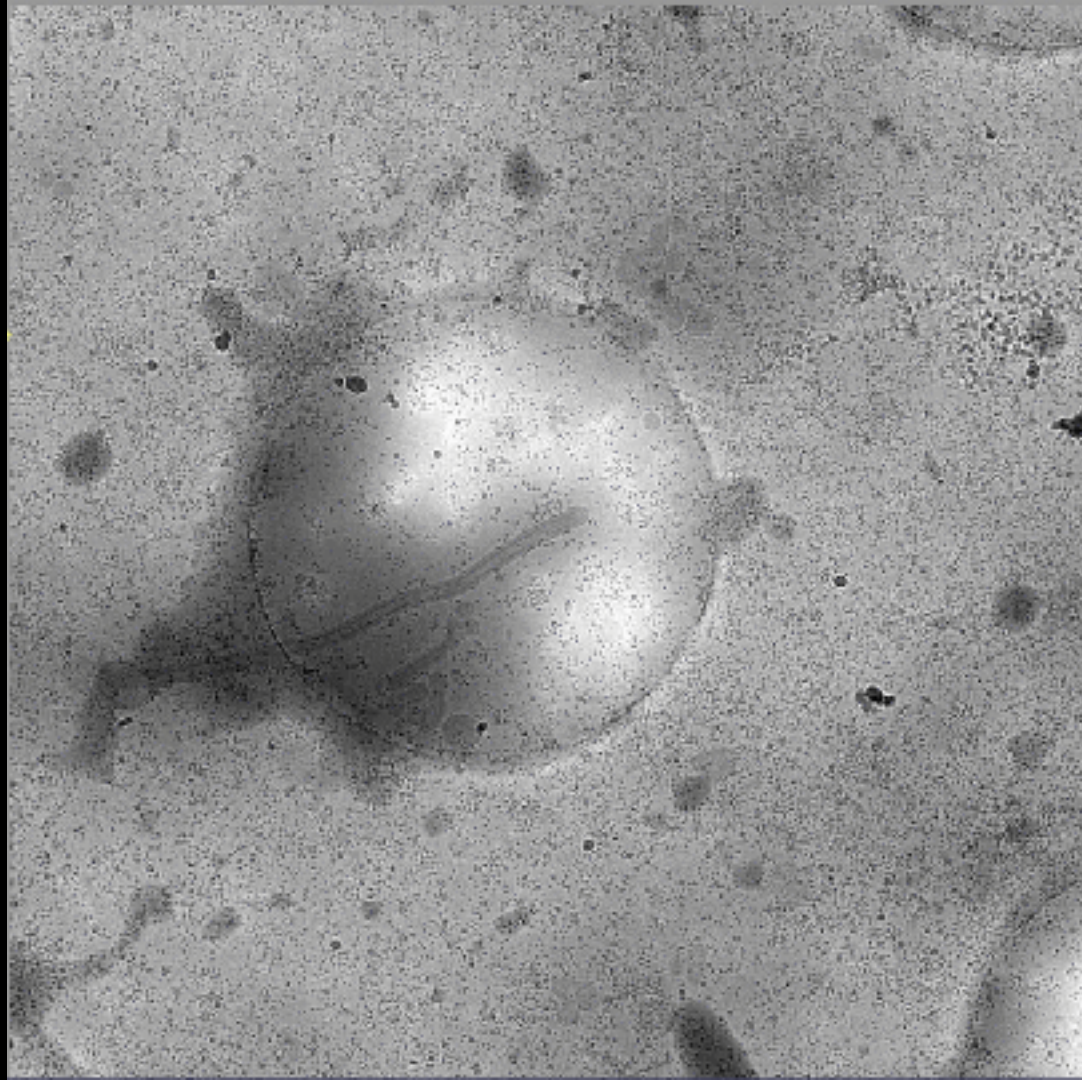
<https://doi.org/10.1016/j.jsb.2016.06.007>

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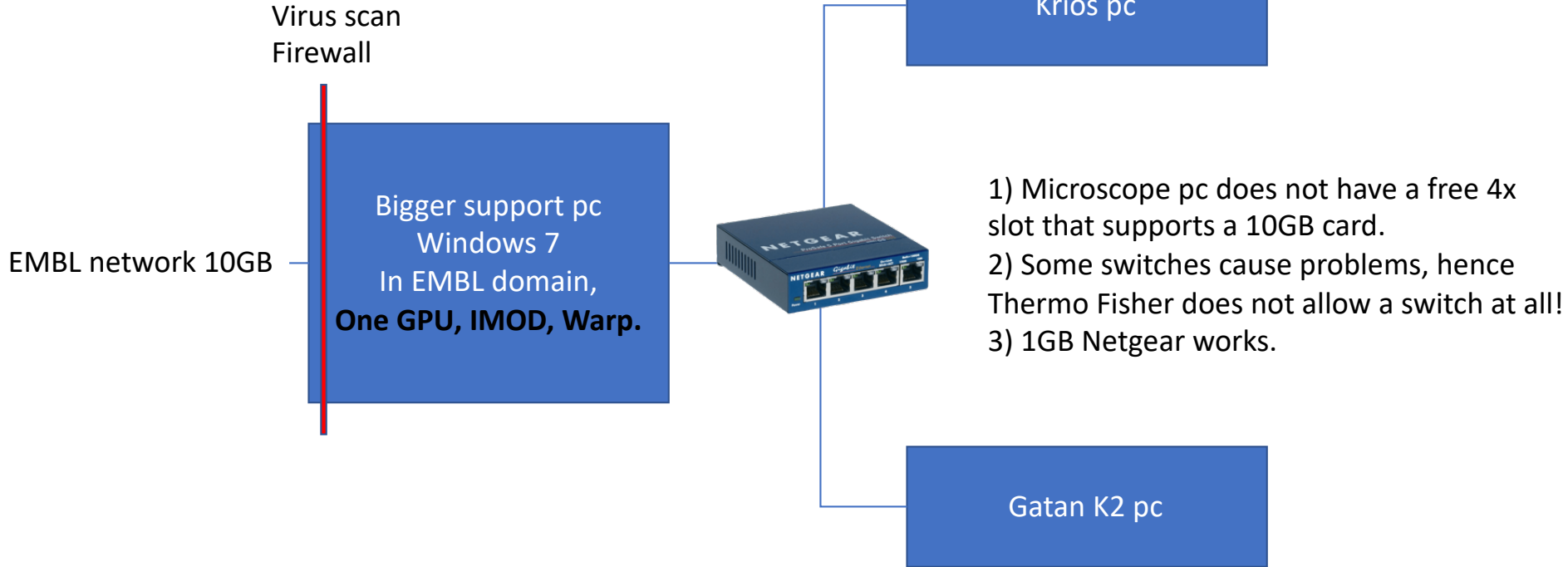
# SerialEM Tomo acquisition

Gatan pc SerialEM:

- Gatan pc has GPU.
- Dose-symmetric tilt scheme.
- On-the-fly frame alignment.
- Also save raw frames as uncorrected LZW compressed tif.
- Currently +/- 60°, 3° steps, 41 images, 27 minutes.
- IMOD FrameWatcher & AlignFrames also possible.



# Remote & data transfer



Advantage: Remote to Microscope AND Gatan pc by VNC through Support pc with port forwarder.

# Remote & data transfer

- Internal users store data on group shares.
- No storage space, no data acquisition!
  
- 10Tb storage for external users.
- Typically for three months.
- Data transfer over EMBL Aspera server.
  
- Warp data?

# Internal training

- Data collection training only.
- EM training on request.

# Internal & external sessions

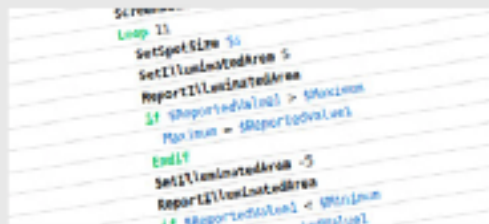
- Minimise need for operator help.



# SerialEM

- Maps, EPU, Tomo, Latitude and LatitudeS.
- Built-in AutoStigmat, AutoComa (uses CTFFIND4).
- Built-in active beam tilt compensation (so far SPA only).
  
- Very flexible.
- Fast support.
- Works with everything.
  
- **Steep learning curve!**

## SerialEM Script Repository



A repository for sharing SerialEM scripts (macros) allowing automation of TEM data acquisition is being provided by Nexperion.

[Read more](#)

## Solutions for Electron Microscopy



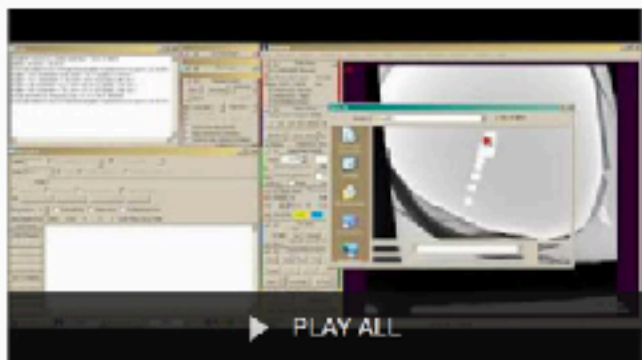
## SerialEM and IMOD



We provide support and training for SerialEM and IMOD, a freely available solution for acquisition and processing of (cryo)electron tomography data.


[Read more](#)

# www.nexperion.net




### SerialEM Lectures and Tutorials

23 videos • 14,694 views • Last updated on Nov 28, 2012

1  **SerialEM Introduction Lecture Part 1**  
BL3DEMC  
11:57

2  **SerialEM Introduction Lecture Part 2**  
BL3DEMC  
13:25

3  **Acquiring a gain reference with SerialEM**  
BL3DEMC  
1:57

[www.youtube.com/user/BL3DEMC](http://www.youtube.com/user/BL3DEMC)



BL3DEMC

SUBSCRIBE 279

5  **SerialEM Introduction Lecture Part 4**  
BL3DEMC  
11:53



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## FAQ

[Cryo-EM Training - Basic \(level I & II\)](#)[SerialEM Training - Basic, Tomography, Single Particle, Advanced](#)[Making Graphene Oxide Grid](#)[Single Particle Data Collection Using SerialEM](#)[Post Processing K2 Frames from SerialEM Data Collection](#)[Align Movie Frames with SerialEM and IMCQ Programs](#)[Monitor Data Collection in The Fly](#)[Rolltize and Distortion Info on K2 Camera](#)[SerialEM detaching and re-attaching sessions](#)[SerialEM Note: Installation and Calibration](#)[SerialEM Note: Have All LHM Maps Automatically](#)[SerialEM Note: Setup Dummy Instance](#)[SerialEM Note: Setup LD with Mk of MP and nP Modes](#)[SerialEM Note: More About Z-Height](#)[SerialEM Note: Speed, Speed and Speed](#)

## SerialEM Note: Installation and Calibration

AUTHOR:	Chen Xu
Contact:	< <a href="mailto:chenxu@umassmed.edu">chenxu@umassmed.edu</a> >
Date_Created:	2017-11-12
Last_Updated:	2018-08-28

### Abstract

When I helped a few sites to install and calibrate SerialEM, I had impression that most new users felt this process was very hard. I felt the same way when I initially learned to install and calibrate SerialEM by myself. I even got frustrated and had to call David for a few times. When I think back about all the troubles I had to install and calibrate SerialEM, I believe I would have an easier time if I had a brief guideline document for what steps to follow in order, and what to do in each step. The handbook from SerialEM is very complete to provide almost all information needed, but it is perhaps a lot to read and not clear where to start for a beginner.

I wanted to list some steps here to guide you through this initial installation and calibration phase. It is like a crash list. For more detailed information, you should always find it from helpfile.

### Installation

Here are steps to follow:

1. Ask David for the initial system file. Normally, you would fill out a "questionnaire" available at the ftp server - <http://bio3d.colorado.edu/ftp/SerialEM/questionnaire.txt> and send it to David. David will then create a framework file on the same ftp server for you to download. This framework file is a zip file, you can download it to local like Desktop and unzip it by double clicking on the file. Beside a sub-folder "Admin" created under C:\ProgramData\SerialEM, the most important file is the framework file one initial system file called "SerialEM\properties.txt". You must have this file to get started. Please refer to the SerialEM webpage for the latest information regarding this.
2. Make sure you have enough computer performance, computer are on the same network. For



## Acknowledgements:

- David Mastronarde
- Chen Xu
- The SerialEM community
- Felix Weis

# Questions?

# Latest development

## EM-Tools

- Map squares at 2250x.
- Mark positions.
- Cut positions and scale to 15Kx Low Dose View image: virtual maps.
- Align to virtual maps at 15Kx.

## Why?

- E.g. fibrils, a flock here and there...
- Standard approach:  
2000 images per day, 10% yield.
- EM-Tools approach:  
1000 images per day with 100% yield.
- Selection can be done during grid square mapping.

