
IOCB SERVICE DAYS

Presentation of services by:
Research-Service Groups / Service Groups / Core Facilities

20
23

Cryogenic Electron Microscopy

Tomas Kouba

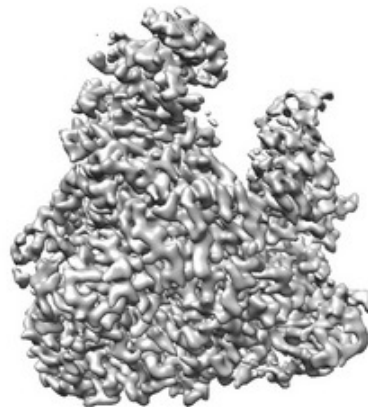
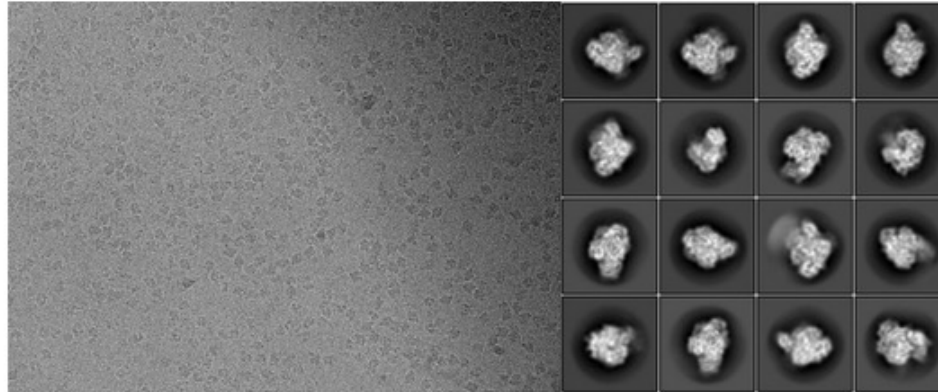


ÚOCHB AV
CR
IOCB PRAGUE



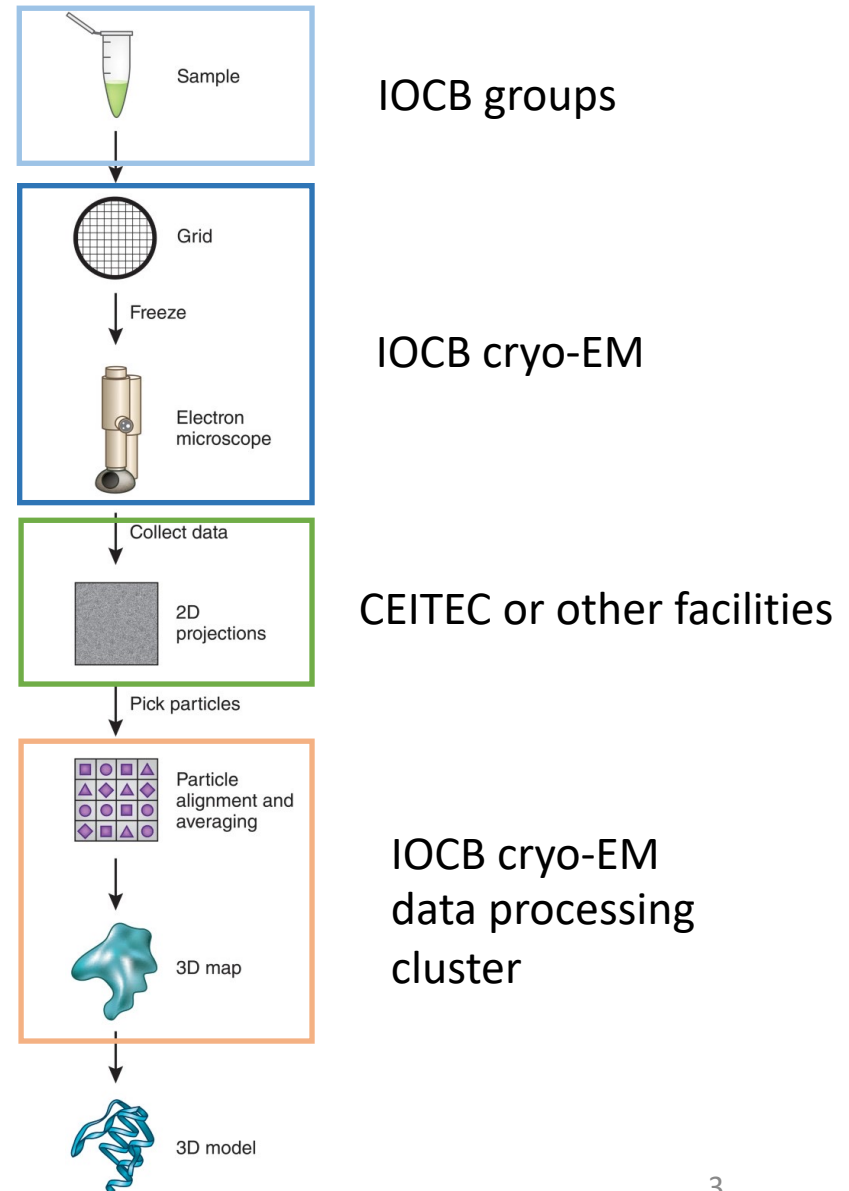
CRYO-EM at IOCB

TOWARDS 3D STRUCTURES OF BIOMOLECULES



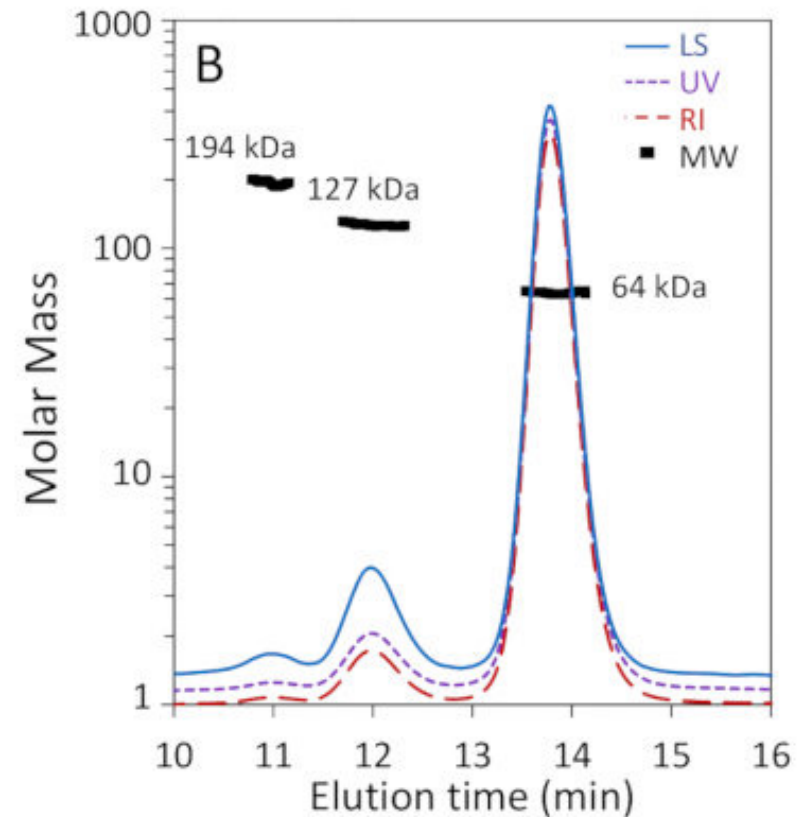
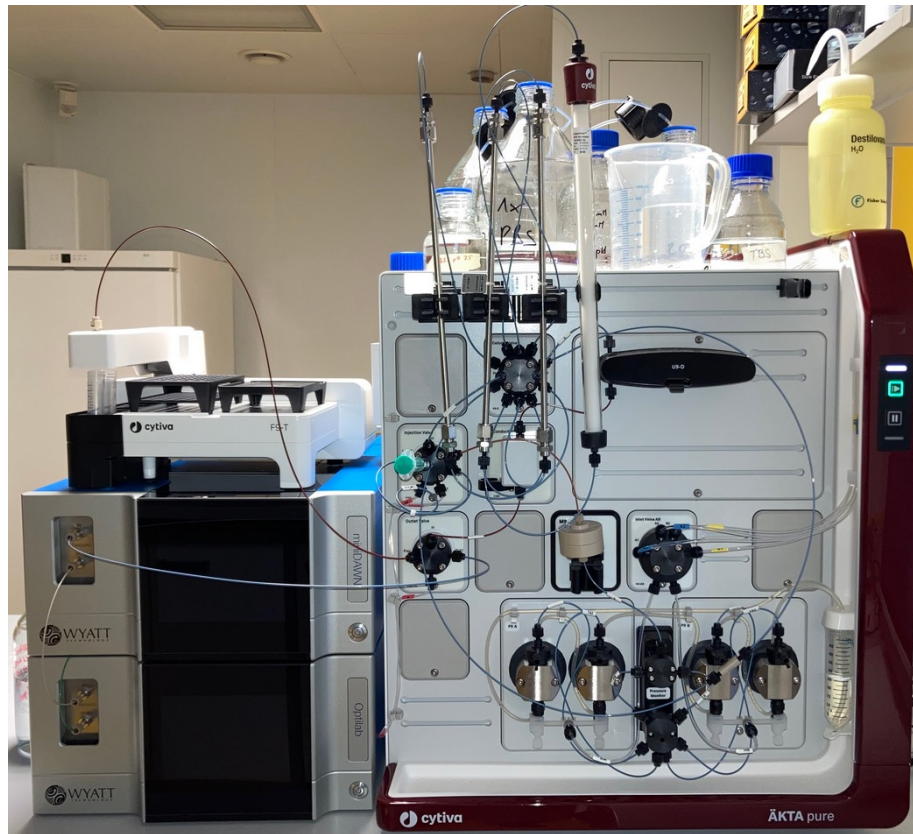
Provisional Cryo-EM facility at the IOCB

- Basic sample characterisation
- Negative stain analysis
- Cryo-EM grid preparation
- Basic cryo-EM screening
- Advanced screening in CEITEC
- Data collection at Europe facilities
- Data processing at IOCB cluster
- **Cryo-EM 3D structure**



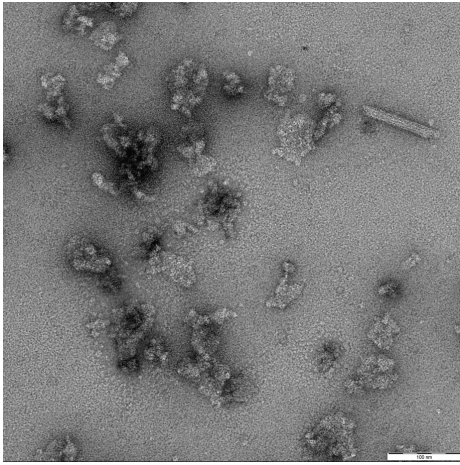
Cryo-EM sample preparation

- Collaboration with Zoll group
- SEC on micro-column (AKTA pure micro), 20 ul injection volumes
- SEC-MALS to characterise molecular weight, oligomer state
- Shodex PROTEIN KW40(3-5)-4F silica based columns – 15 min analysis

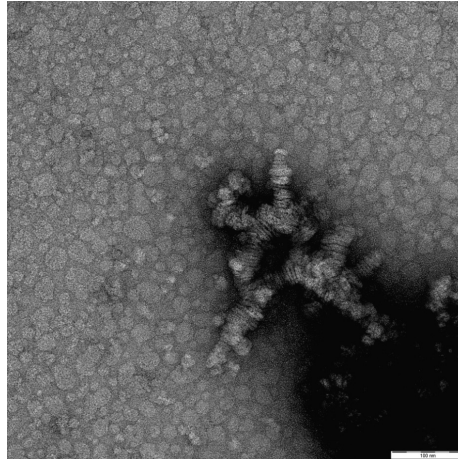


Negative stain analysis

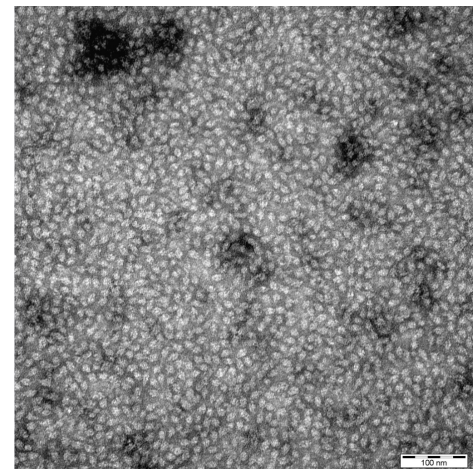
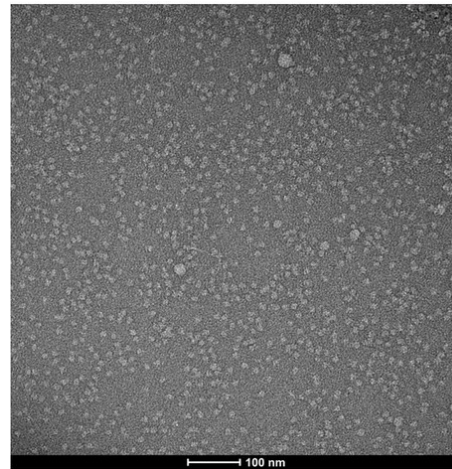
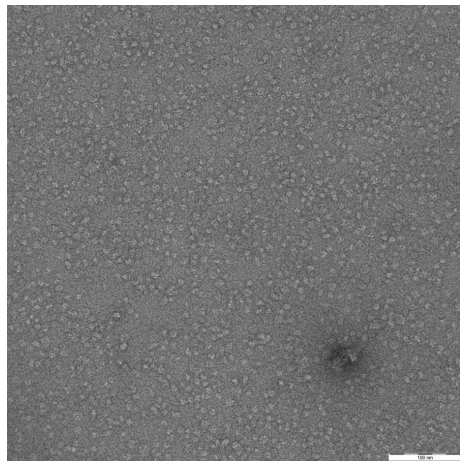
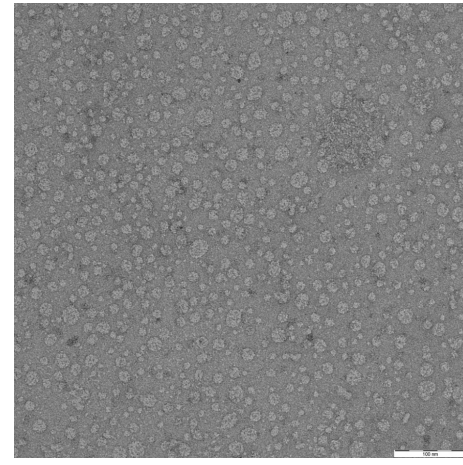
- Quick (half-day) sample quality check



Aggregates/contamination



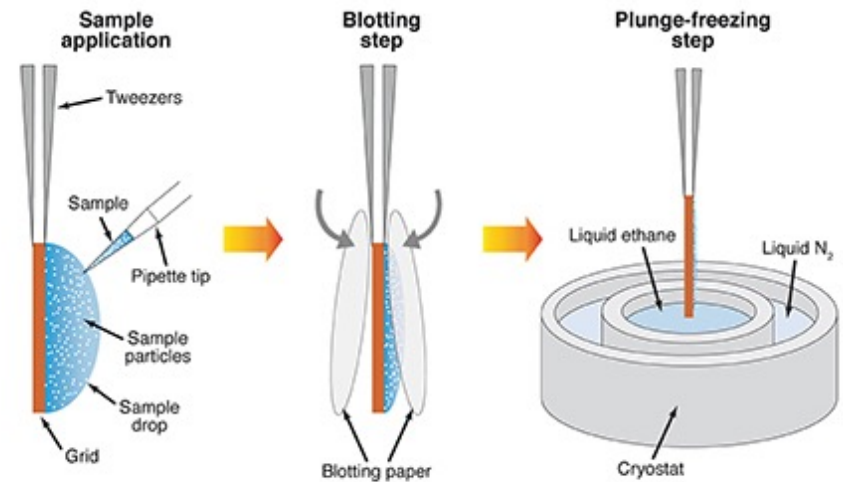
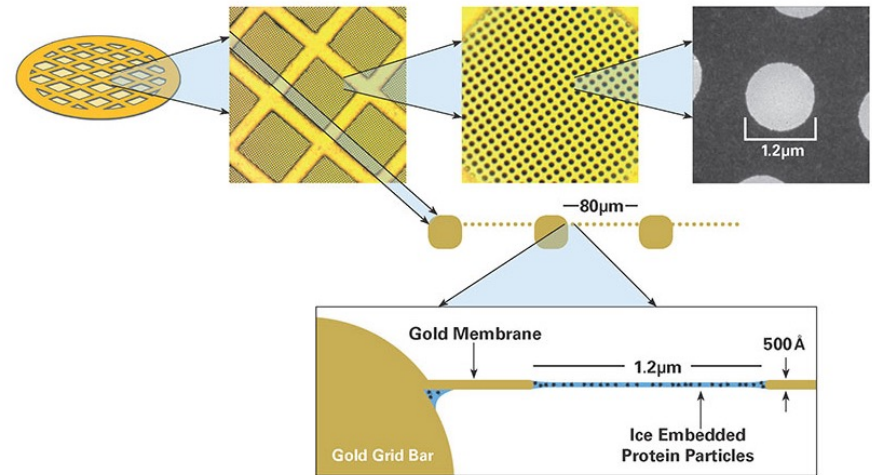
detergent artefacts/heterogeneous sample



Homogenous sample, ready for cryo-EM grid preparation/screening

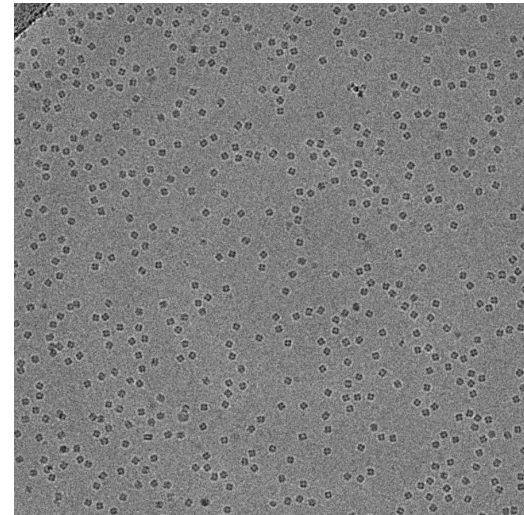
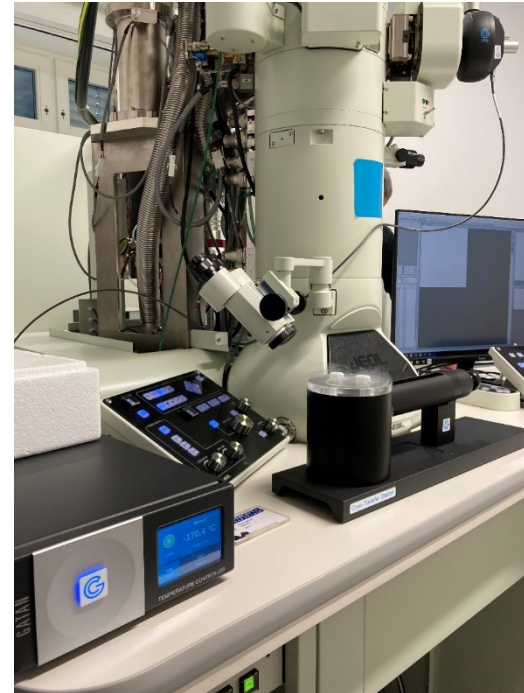
Cryo-EM grid preparation

- TFS Vitrobot Mark IV
- Gold standard in cryo-EM grid preparation
- Samples passing negative stain analysis
- Local cryo-EM storage



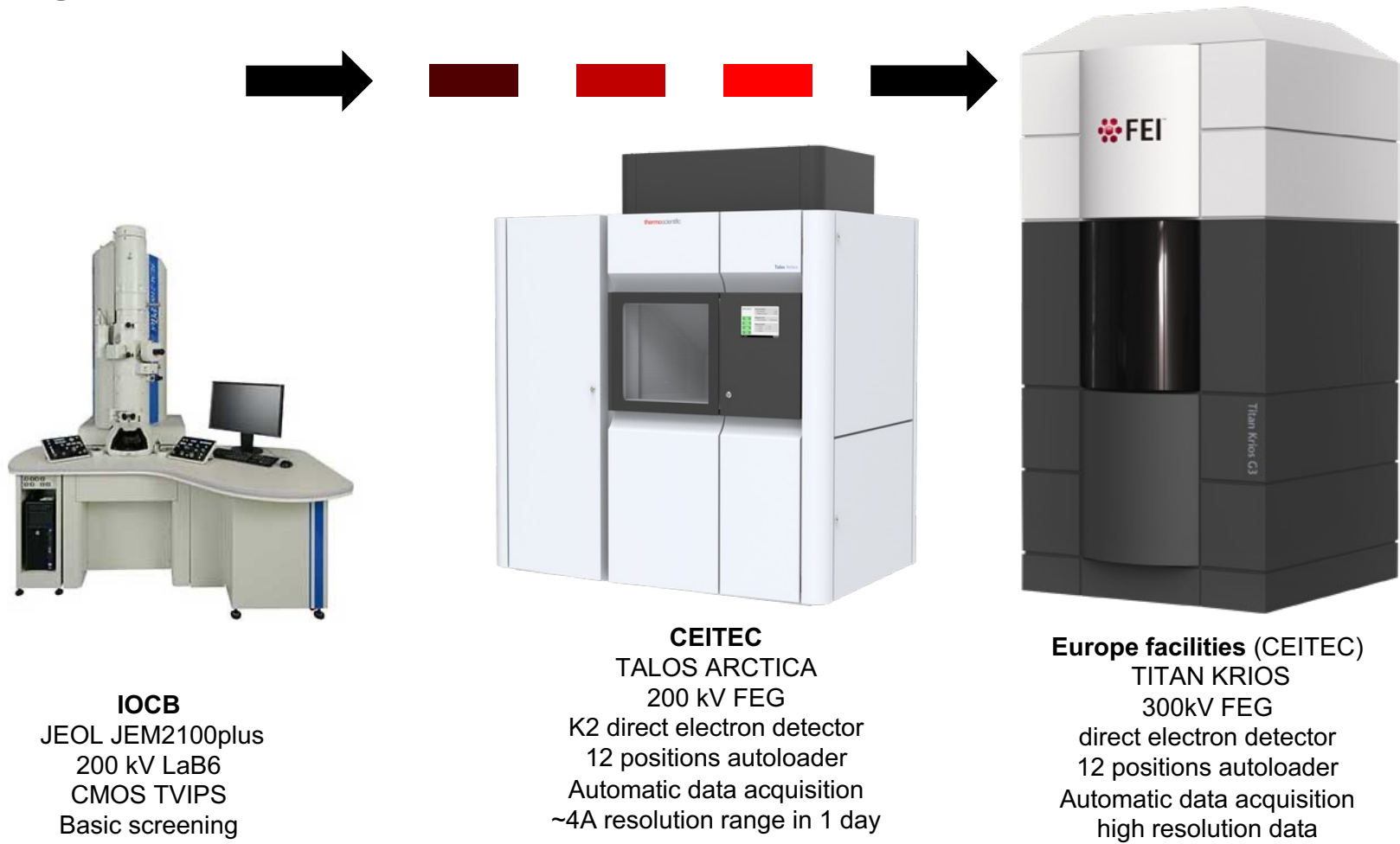
cryo-EM screening

- JEOL JEM2100plus
- 200kV, LaB6 source, cryo-polepiece, side entry holder
- TVIPS CMOS TemCam-XF416ES camera, 48 frames/s
- serial EM low dose software for SPA analysis and Tomography



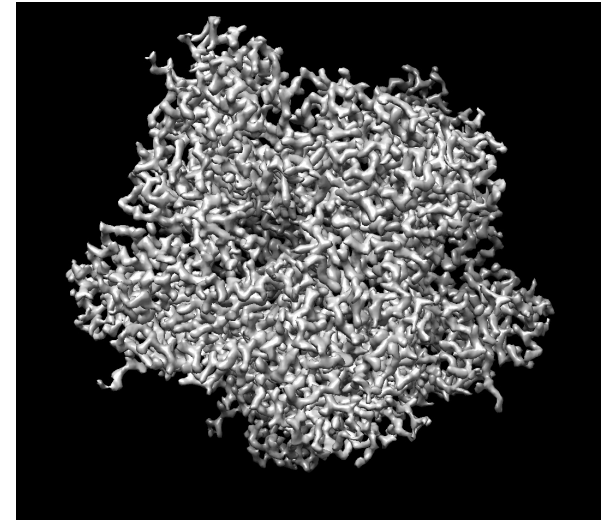
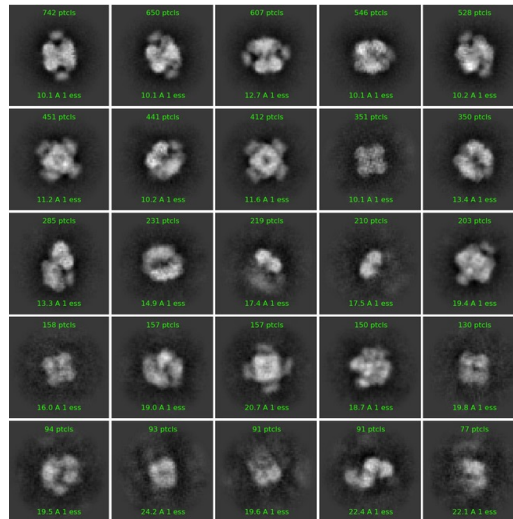
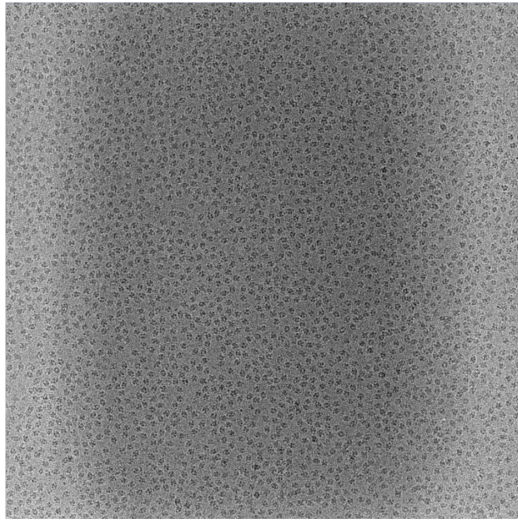
Advance screening and high resolution data collection in CEITEC, Brno

- Access through CIISB
- Peer reviewed proposal
- Regular remote access



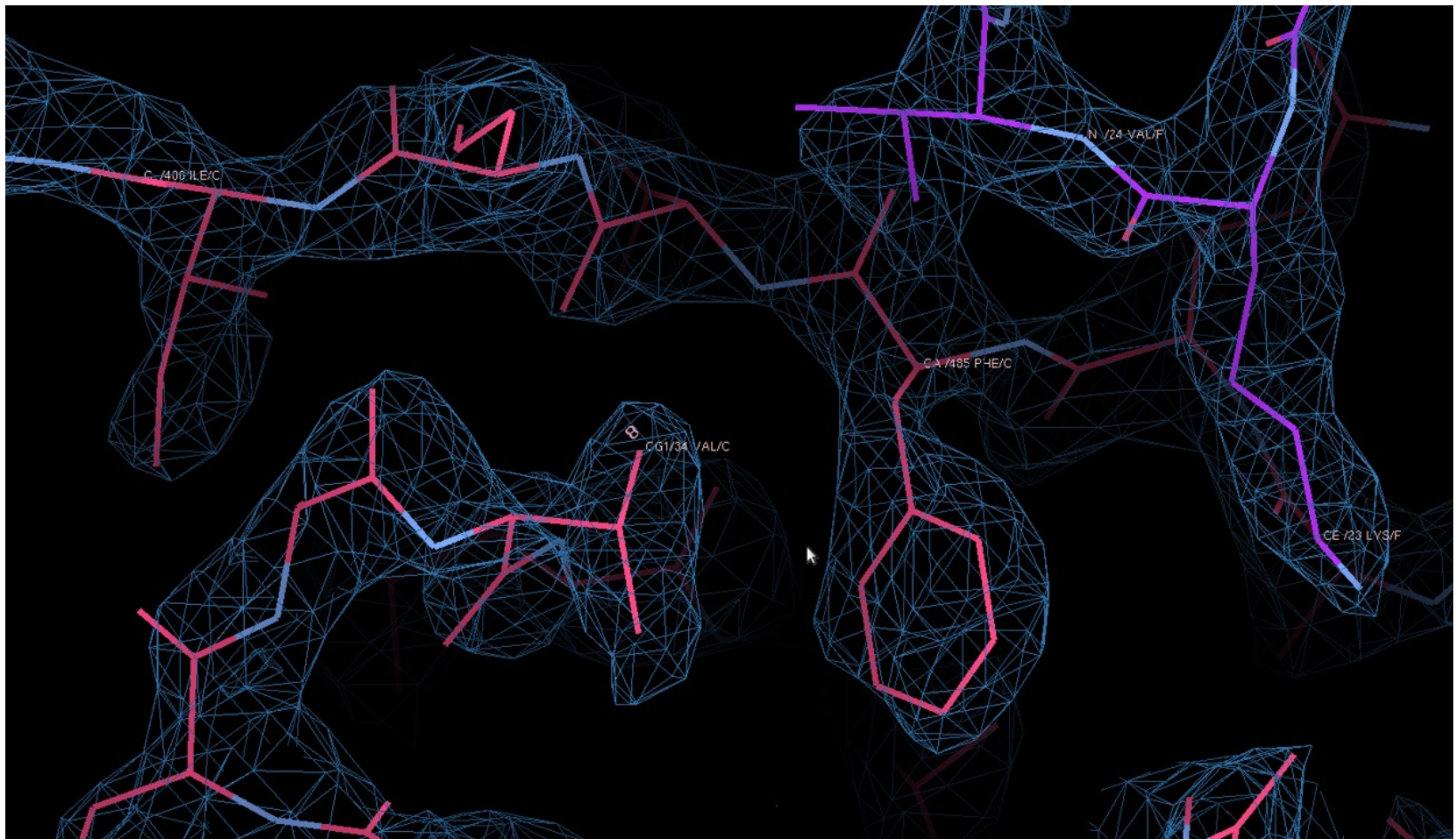
Cryo-EM data analysis

- 2Ds, 3Ds, 3D refinement



Cryo-EM data analysis

- 2Ds, 3Ds, 3D refinement
- ~2.5Å resolution cryoEM map (Iva Pichova group)
- ~6 months



Cryo-EM computing at IOCB

- Teeming up with IOCB computing core; Hector Martinez-Seara
- Cryo-EM WIKI on hpcg.uochb.cas.cz
- **SBGrid license available** sbgrid.org

To run RELION GUI

```
source /opt/uochb/soft/spack/20210921-git/share/spack/setup-env.sh
spack load relion@3.1.3
spack load fftw@3.3.9+mpi
spack load cuda@11.0.2
spack load openmpi@4.1.1~legacylaunchers

relion
```

https://hpc.nih.gov/apps/RELION/relion31_tutorial.pdf

[IMPORT_MOVIES](#)

[MOTION CORRECTION MC](#)

[CTFFIND](#)

[PARTICLE PICKING TOPAZ](#)

[PARTICLE_EXTRACTION](#)

[2D_CLASSIFICATION](#)

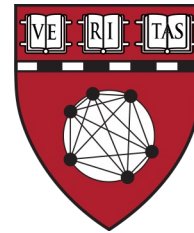
[PARTICLE_SELECTION](#)

[3D_CLASSIFICATION](#)

[3D_REFINEMENT](#)

TIPS from other HPC centers:

- [NIH](#)



SBGrid
CONSORTIUM

SERVICE vs TRAINING

‘Every user should get the maximum out of their data collection session’

SERVICE SCHEME

- Sample preparation
- Screening
- Data collection
- Data processing
- Collaboration based

cons:

- no knowledge and skill transfer
- depends on facility capacities

TRAINING SCHEME

- Training of basic users
- Training of independent super-users
- Knowledge and skill transfer
- Super-users can train basic users
- Motivated users
- Growing EM-community
- More time for facility development

cons:

- initial lag time

Cryo-EM team

Anatolij Filimoněnko

Cryo-EM sample preparation and screening

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Hana Šváchová

Cryo-EM data processing

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Kiran Kumar Telukunta

IT development and support

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EM team

Romana Hadravová

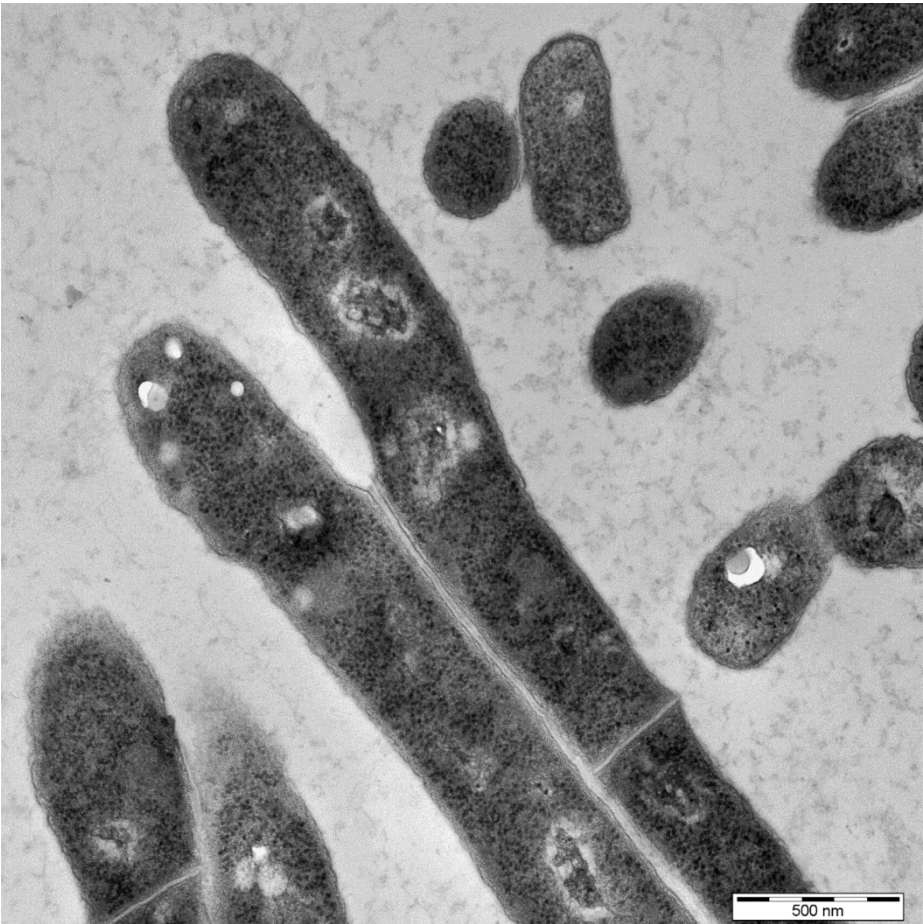
romana.hadravova@uochb.cas.cz



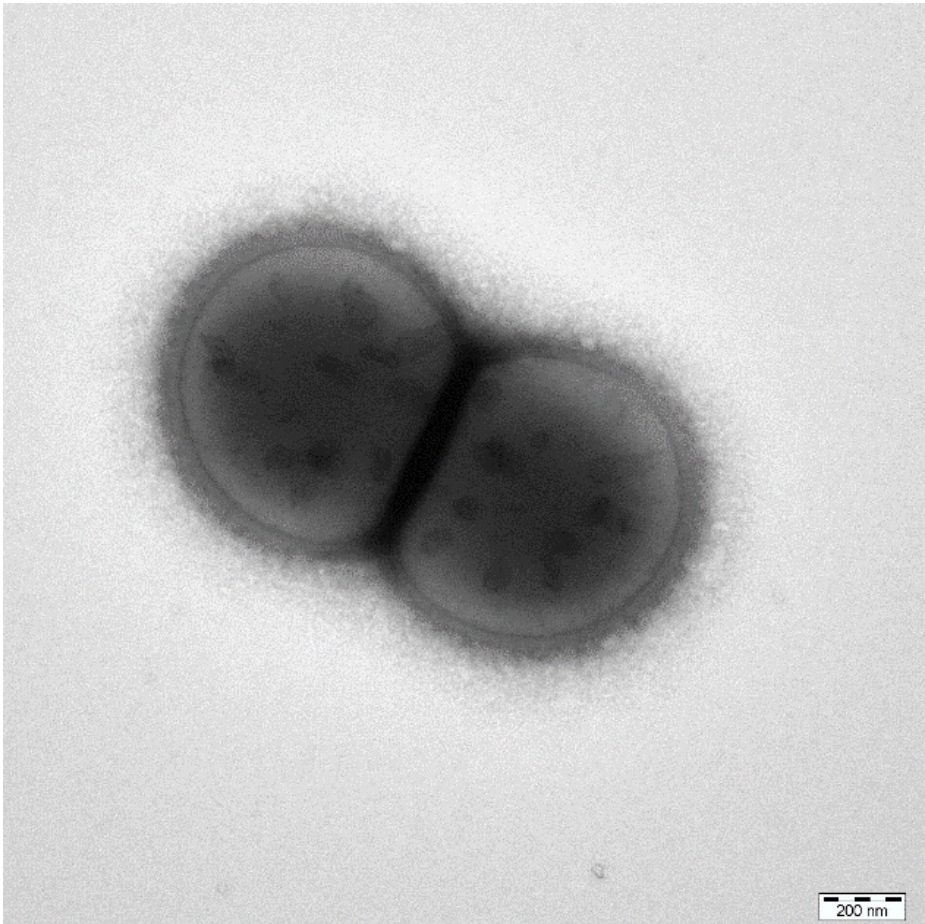
Ultrathin sections

- Cells
- Tissues
- Immunogold labeling

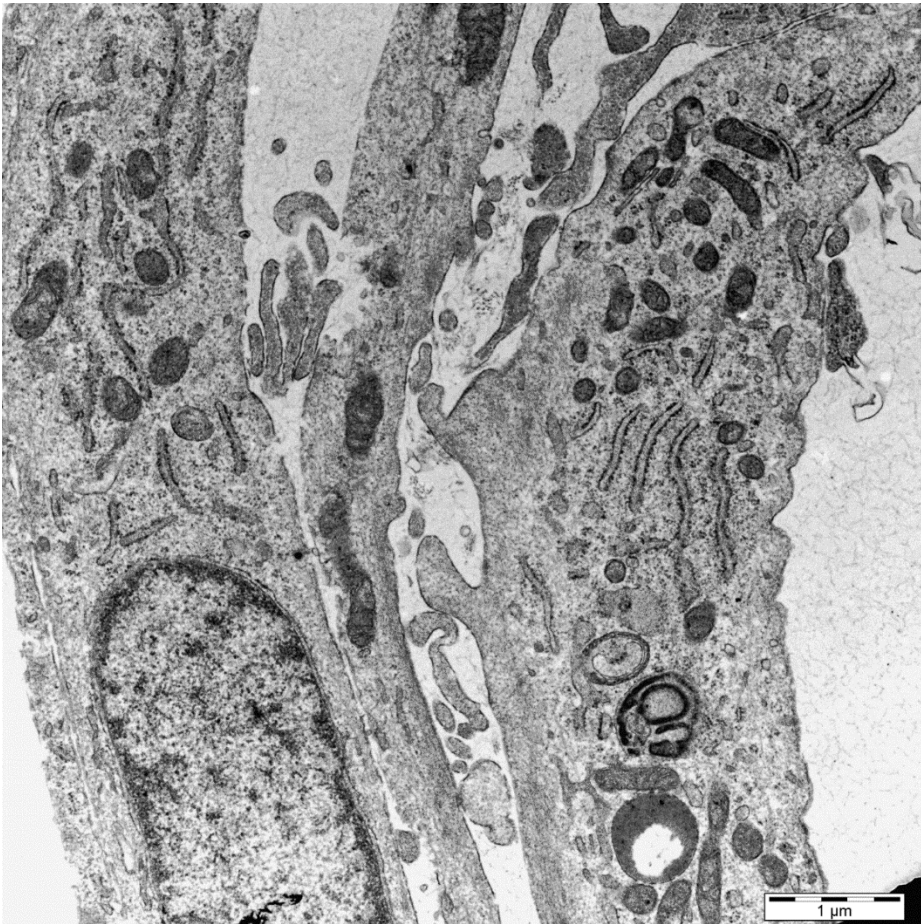
Materials



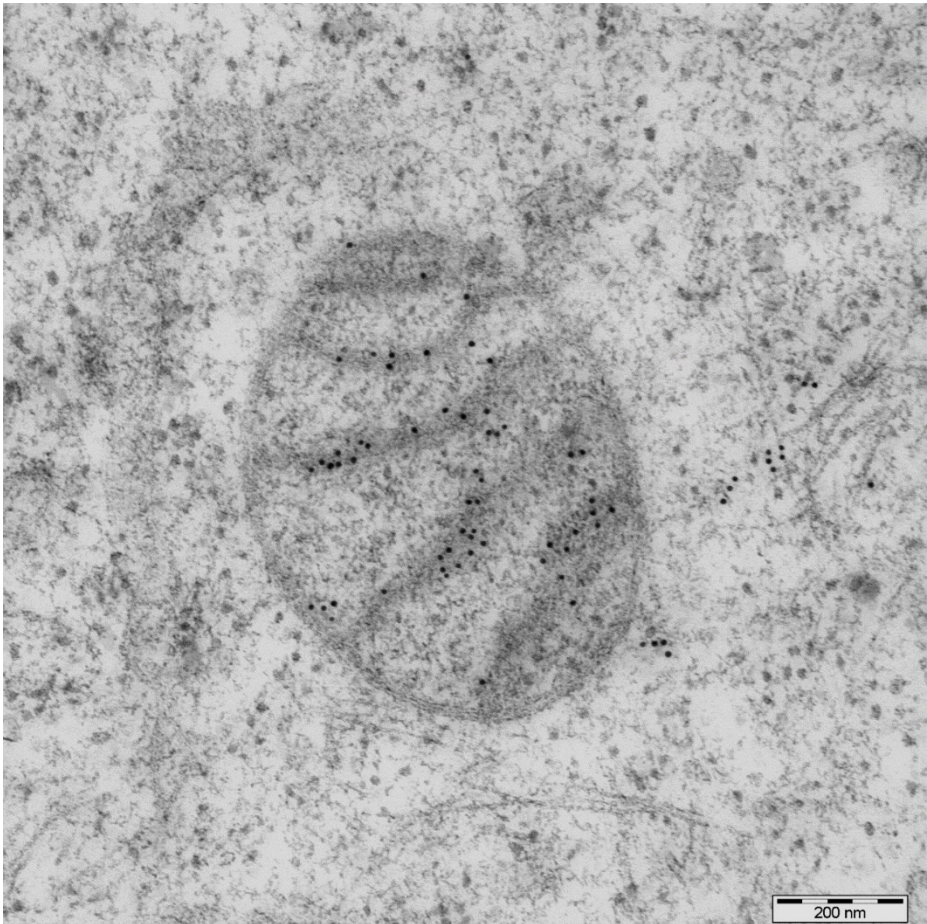
Mycobacterium



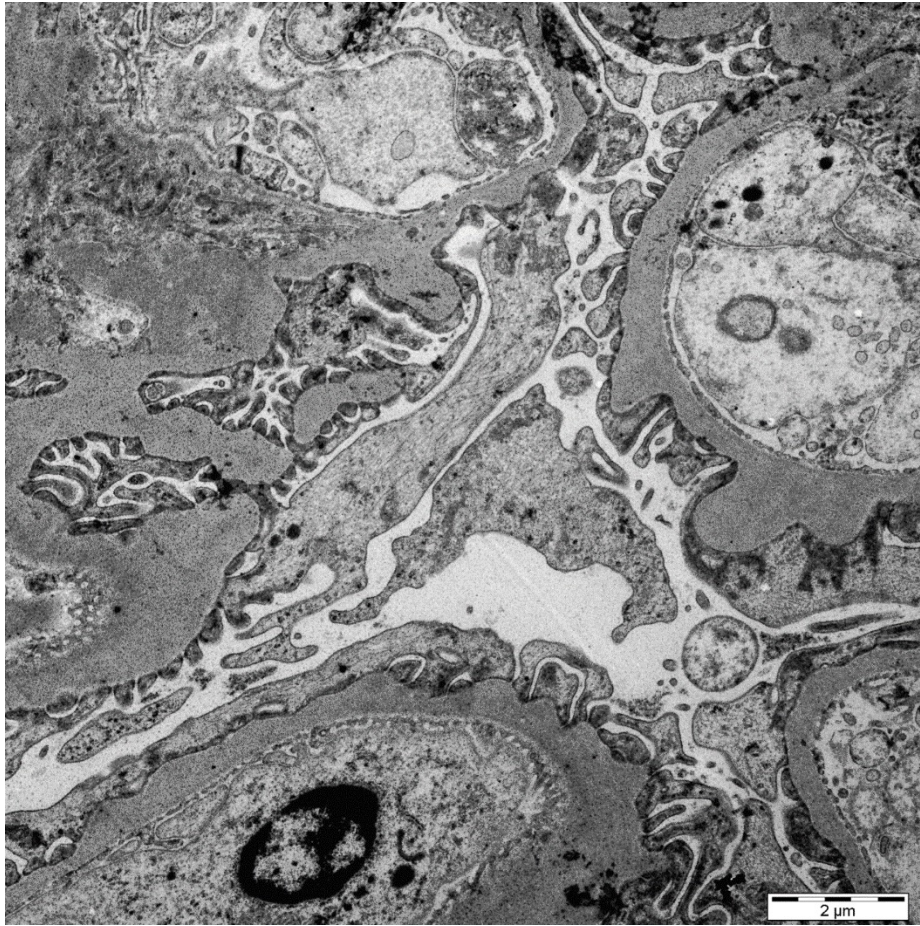
Staphylococcus Aureus



U251 cells

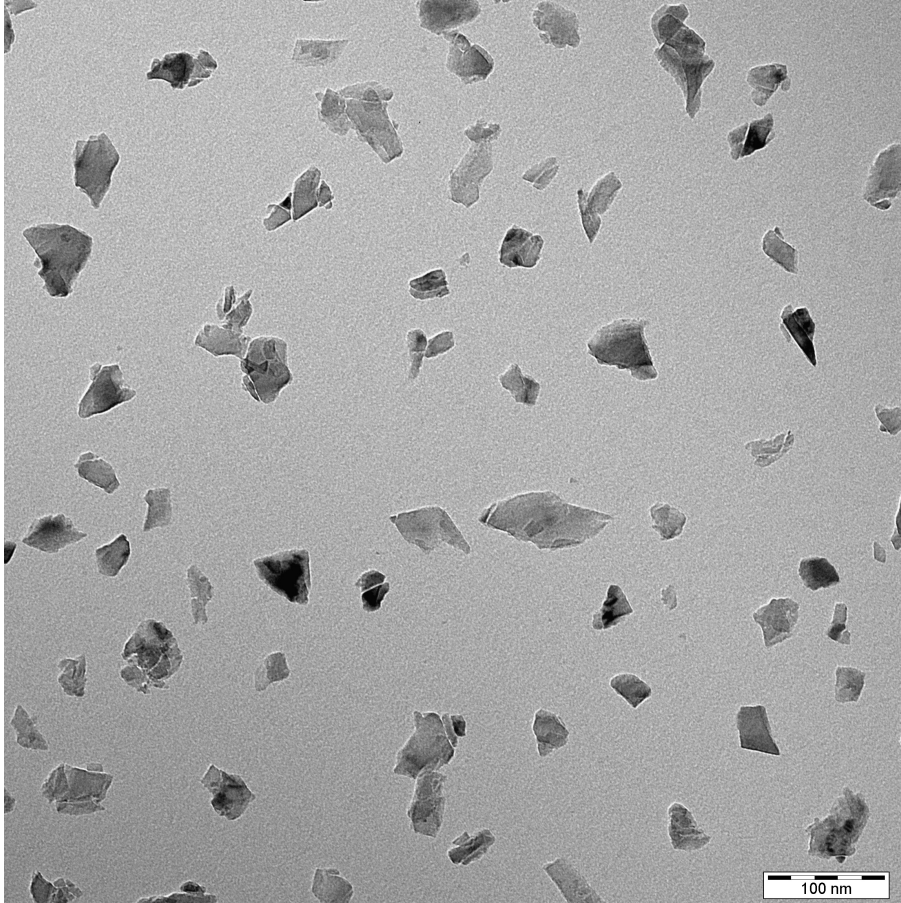


Gold immuno labelling

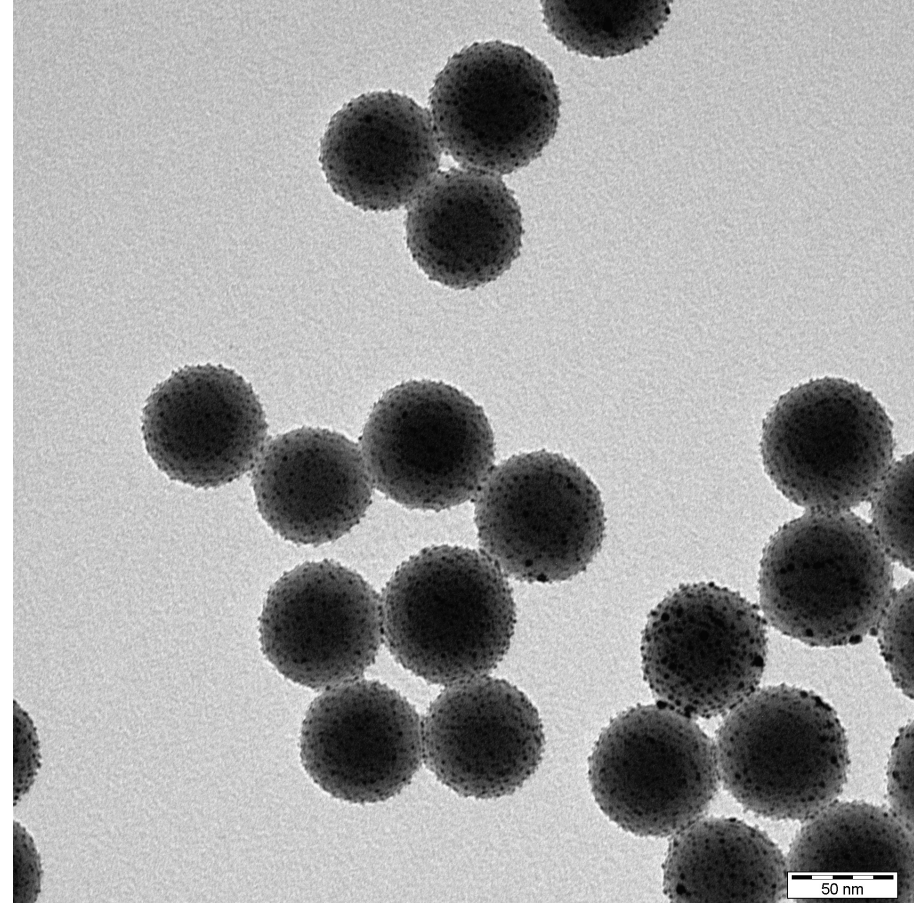


Kidneys

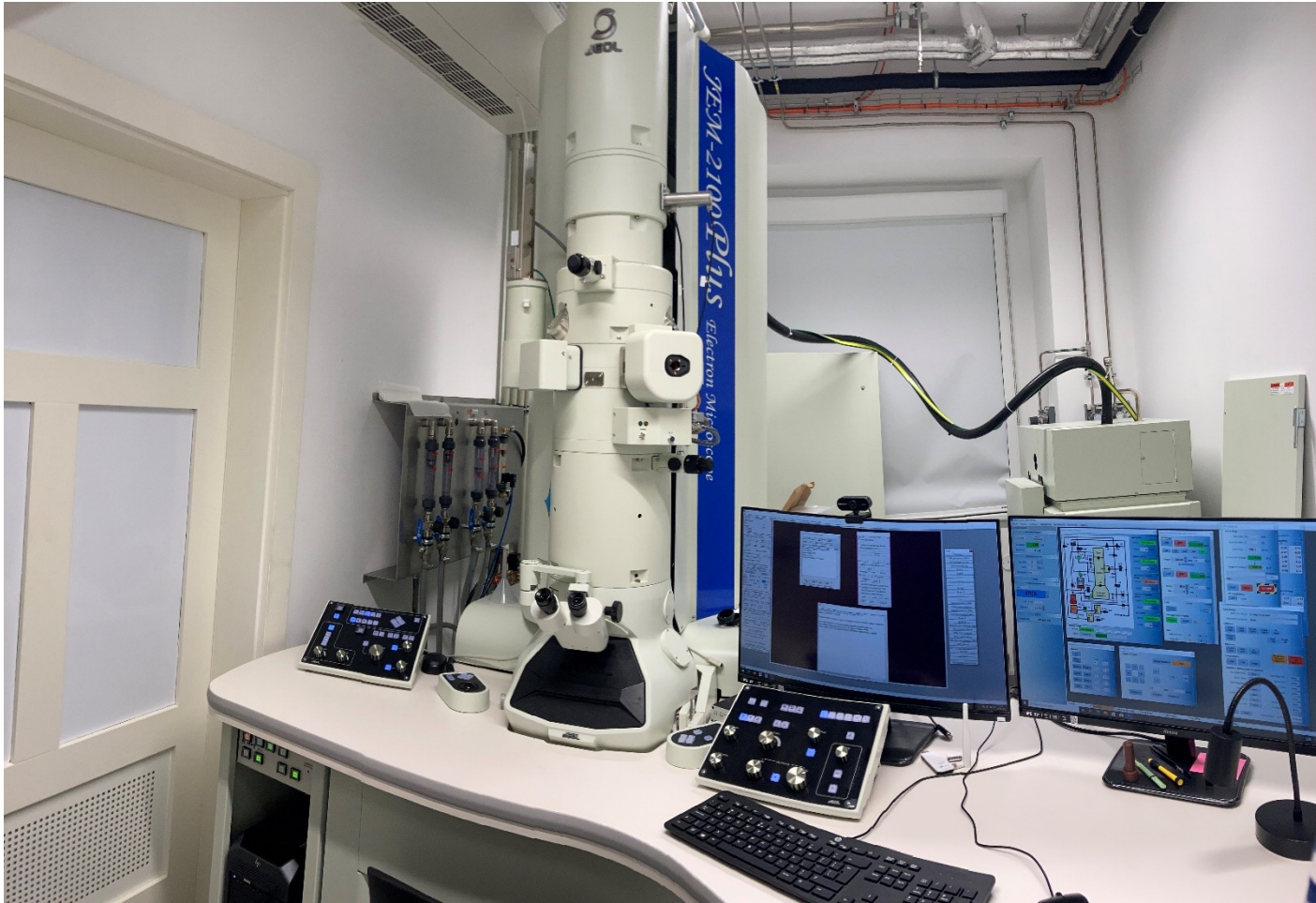
Materials



Diamond particles



Silicone particles
with gold nanoparticles



THANK YOU FOR YOUR ATTENTION !