



CRYO-ELECTRON MICROSCOPY AT THE ESRF



The European Synchrotron

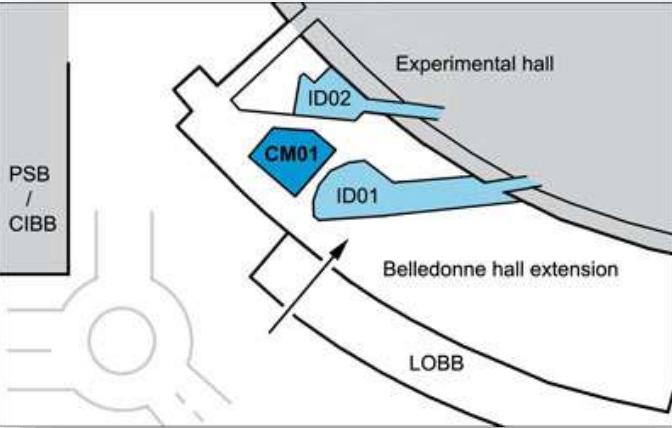
www.esrf.fr/home/UsersAndScience/Experiments/MX/About_our_beamlines/CM01.html 50% Search

 ESRF
The European Synchrotron

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CM01 CRYO-ELECTRON MICROSCOPE



PSB / CIBB



Our beamlines

CM01 CRYO-ELECTRON MICROSCOPE

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Information for users



Synopsis

The ESRF has acquired a Titan Krios cryo-electron microscope (cryoEM) for single particle experiments. It is equipped with a K2 Summit direct electron detector, a Quantum LS imaging filter and a Volta phase plate (VPP). The microscope is situated at the ESRF and is run as a beamline (CM01) with access for the international Structural Biology community given on scientific merit through [rolling-access applications](#). In the longer term, CM01 will be part of a cryoEM platform run within the [Partnership for Structural Biology](#) (PSB) in conjunction with the microscopes located at the IBS and the EMBL. A team of three scientists from the ESRF, the IBS and the EMBL will support user operation.

CM01 TEAM



Guy Schoehn,
Scientist, IBS

Grégory
Effantin,
Scientist,
IBS

Michael Hons,
Scientist,
EMBL

David Flot,
Engineer,
ESRF

Isai Kandiah,
Scientist, ESRF

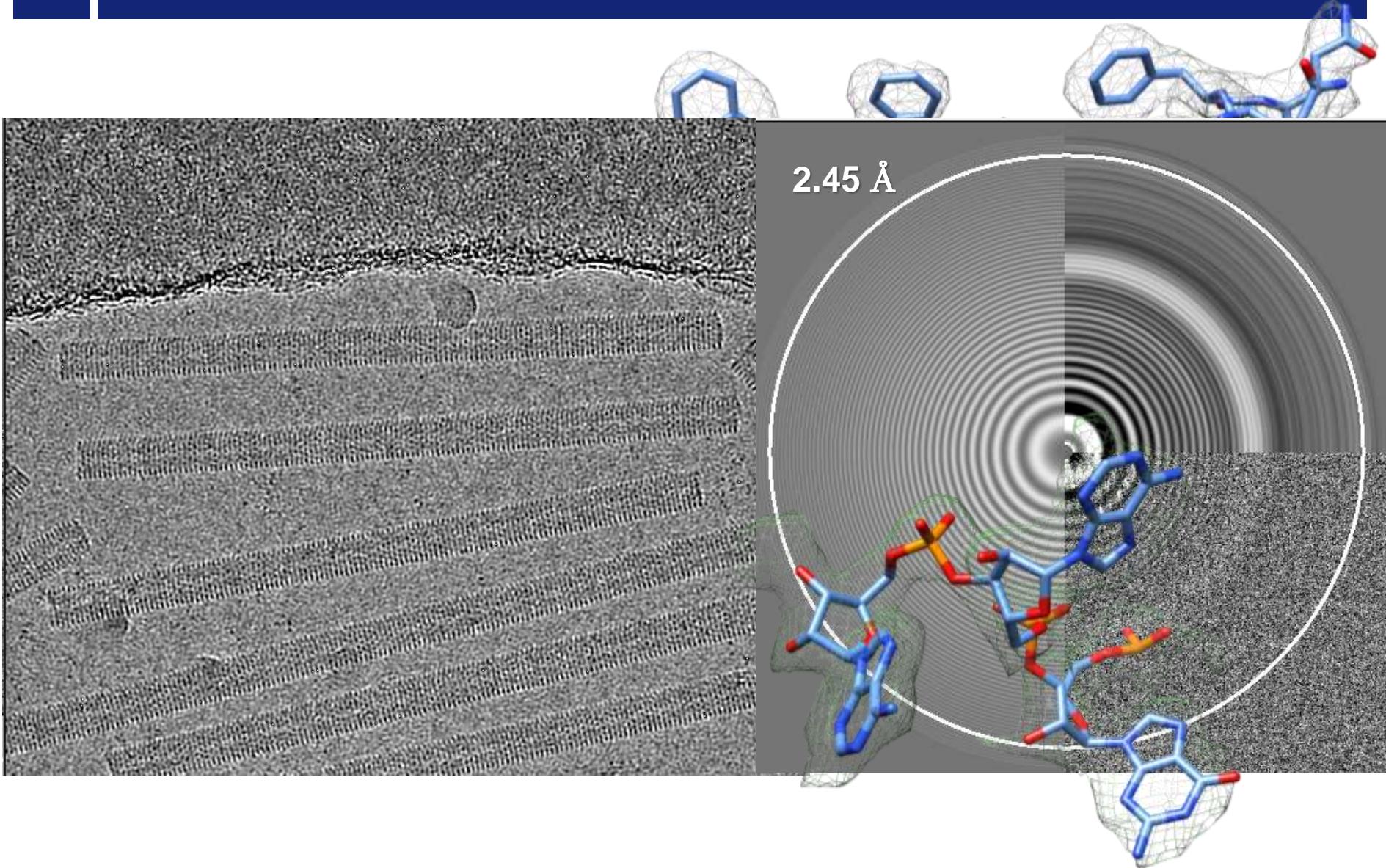
Christoph Mueller-
Dieckmann,
Scientist, ESRF

CM01 - INFRASTRUCTURE

Cryo-EM

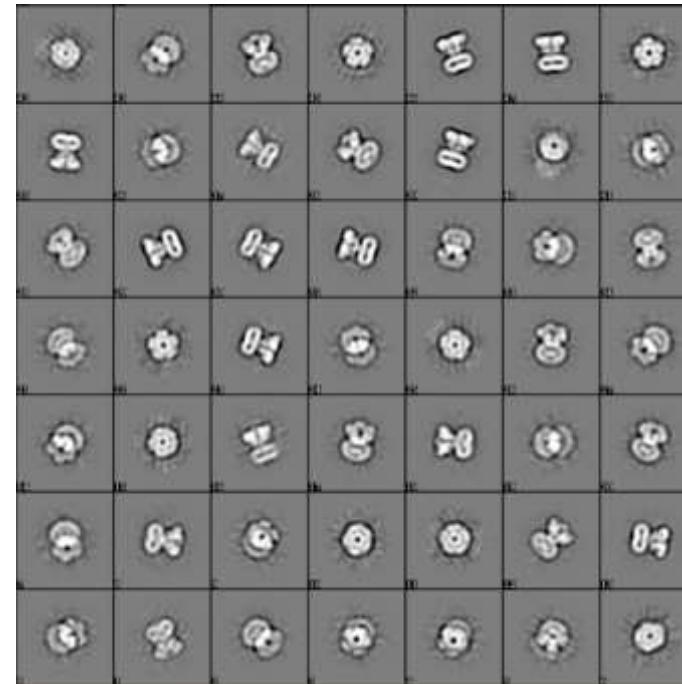
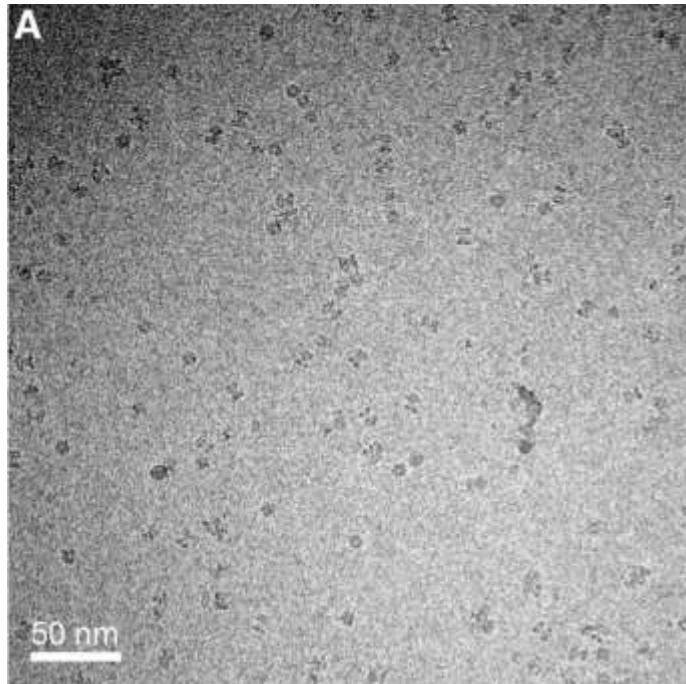


CM01 – BENCHMARK



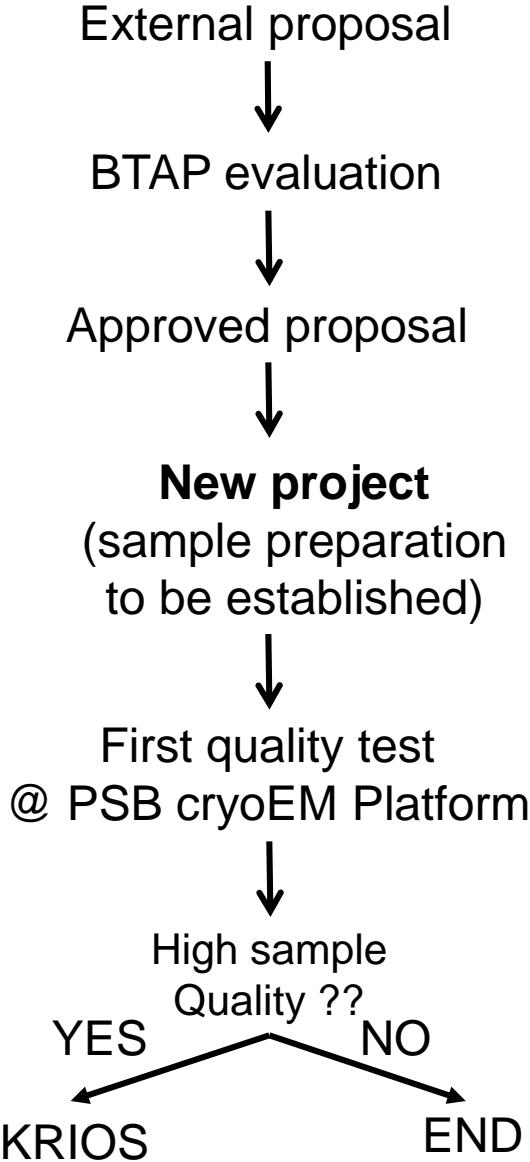
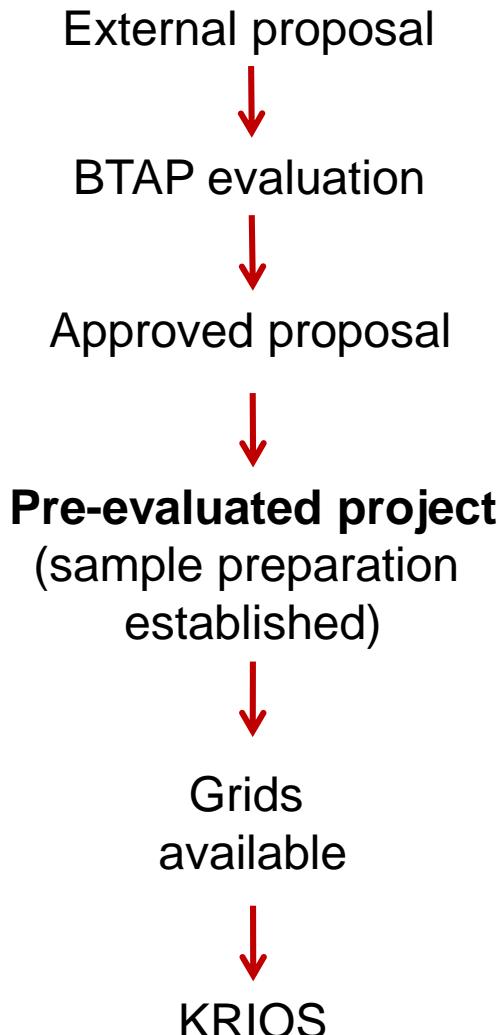
Rolling access proposals

➤ BTAP



doi: 10.1016/j.cell.2015.12.055

CM01 – WORKFLOW – FROM SUBMISSION TO DATA



- **600 shifts per year**
- **9 shifts, each of 8 hours, per proposal**
- **2 Proposals per week**
 - **Proposal 1 : Monday to Thursday (Tuesday MDT)**
 - **Proposal 2: Friday to Sunday**
- **9 shifts program:**
 - **1 shift for screening and setting up of data collection (User's presence is useful)**
 - **8 shifts of automated data collection (User's presence not needed)**

CRYO-EM PLATFORM @ EPN



Titan Krios @ ESRF
+ sample preparation



Cryo-EM
@ EMBL-Gr



Cryo-EM
@ IBS



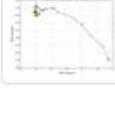
CM01 – ON-THE-FLY PROCESSING AND ISPYB UPLOAD

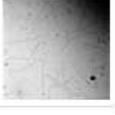
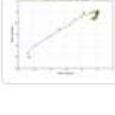
 **ExiMX** Extended ISPyB for MX BETA

Version: 5.4.4
Released: 20171211

Home Shipment ▾ Prepare Experiment Data Explorer ▾ Manager ▾ Help ▾ SMIS search by protein acronym Log out @ka

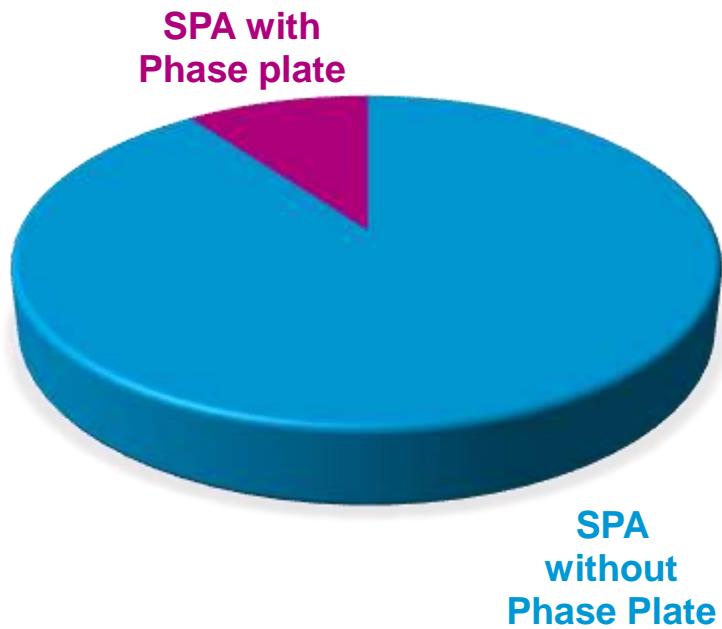
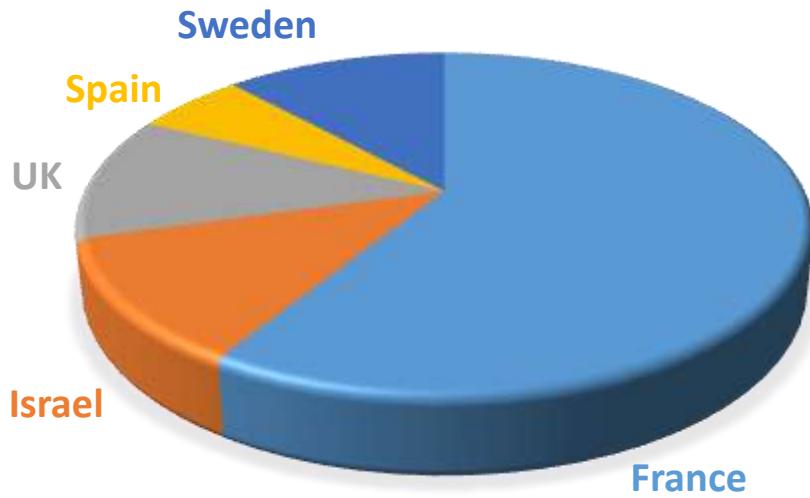
New Tab

Number	12224		Total Motion	195.2		Resolution Limit	3.111	Angle	79.03	
X, Y	-0.000, -0.000		Avg. Motion/frame	4.9		Correlation	0.124072	Estimated B factor	135.55	
Dose	0.7700		Frame Range	1 - 40		Defocus U	21742.38			
			Dose	1.1000		Defocus V	21535.78			

Movie #12223 03-02-2018 22:32:32		Instrument MR								
Movie		Motion Correction				CTF				
Number	12223		Total Motion	199.8		Resolution Limit	3.266	Angle	20.93	
X, Y	-0.000, -0.000		Avg. Motion/frame	5.0		Correlation	0.132302	Estimated B factor	134.27	
Dose	0.8000		Frame Range	1 - 40		Defocus U	17526.38			
			Dose	1.1000		Defocus V	17974.08			

CM01 – ON-THE-FLY DATA BACK-UP POSSIBLE

CM01 USAGE



- **CM01 will be operational during EBS shutdown**
- **ILL Beamline schedule will be followed for Guest house and Canteen support**
- **9 shifts, each of 8 hours, per proposal**
- **2 Proposals per week**
 - **Proposal 1 : Monday to Thursday (Tuesday MDT)**
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