

CROSS Cryo-EM

General Policy

Li Ka Shing Cryo-EM Laboratory

Cores in CPOS



**HKU
Med**

LKS Faculty of Medicine
Centre for PanorOmic Sciences
香港大學泛組學科研中心

Biobank Core

Genomics Core

Proteomics and
Metabolomics
Core

Bioinformatics
Core

Imaging and
Flow Cytometry
Core

Bioresearch
Support Core

Bioreagent Core

LKS Cryo-EM
Laboratory

HKUMed
Laboratory of
Cellular
Therapeutics

FMB Cores

Laboratory Block, 21 Sassoon Road

Online Platform



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Laboratory

HKUMed
Laboratory of
Cellular
Therapeutics

iLab

PPMS

Bioreagent Core
Online Purchasing
System

iLab



Li Ka Shing Cryo-EM Laboratory

Email: cryoem.cpos@hku.hk

Tel: 3910-2938

Opening hours: 9:00 am to 5:30pm

CPOS

enquiry.cpos@hku.hk | 3910-6600

HKU Med LKS Faculty of Medicine
Centre for PanorOmic Sciences
香港大學泛組學科研中心

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Li Ka Shing Cryo-EM Laboratory

Overview | Benefits | Applications | Process Steps | Imaging Equipment | Access Information During Soft Launch | Charges | Contact

Overview

Cryo-Electron Microscopy is the imaging of specimens frozen in vitreous ice and maintained at liquid nitrogen temperature using Electron Microscopes. In this method, specimens can be studied in their native state without dyes or fixatives, enabling the analysis of fine cellular structures, viruses, and proteins at molecular resolution. Despite being a decades-developed technique, Cryo-EM has been attracting interest since 2013 as a result of technological and algorithmic improvements that have driven a dramatic improvement in the resolution achievable using this technique (dubbed the 'resolution revolution'). In 2017, the technique won the Nobel Prize in Chemistry.

The Cryo-EM technique is becoming the first choice of many structural biologists when analyzing the protein structure experimentally. As a technique for determining the atomic structure of macromolecules that neither crystallize nor are difficult to crystallize under certain conditions, Cryo-EM has the same level of resolution as X-ray crystallography. Cryo-EM is the best way to study cell architecture, large proteins, membrane-bound receptors, or complexes of macromolecules.

General Rules and Security



The Core is under surveillance **24/7**



Do **NOT** lend account to other users



Only access booked equipment



The last user of the day must turn off light and lock the doors



No Drinking And Eating



Turn off the machine right after use



Must attend training before using instrument



Always wear appropriate protective clothing and glasses when working in the laboratories.

User Responsibility



User should strictly follow the standard operation protocol (SOP)



Please operate the instruments carefully and gently



Keep workspace / sample preparation bench tidy and clean



Report any problem related to instruments



Write down experiment settings in logbook

If you are uncertain about performing a particular procedure, please contact Cryo-EM core staff.

General Safety



Staff has the right to query and, if necessary, stop any activity that is considered unsafe.



Avoid working alone during non-office hours in the laboratory.



Plan your work well before getting started.

General Safety



Fire Extinguisher

First Aid Box



Safety shower at corridor



Fire Escape Route and First Aid box.

No gloves on computer and areas accessible by others.



Dispose biological waste in designated bins.

Dispose sharps / glasses in sharp box.



Emergency Exit



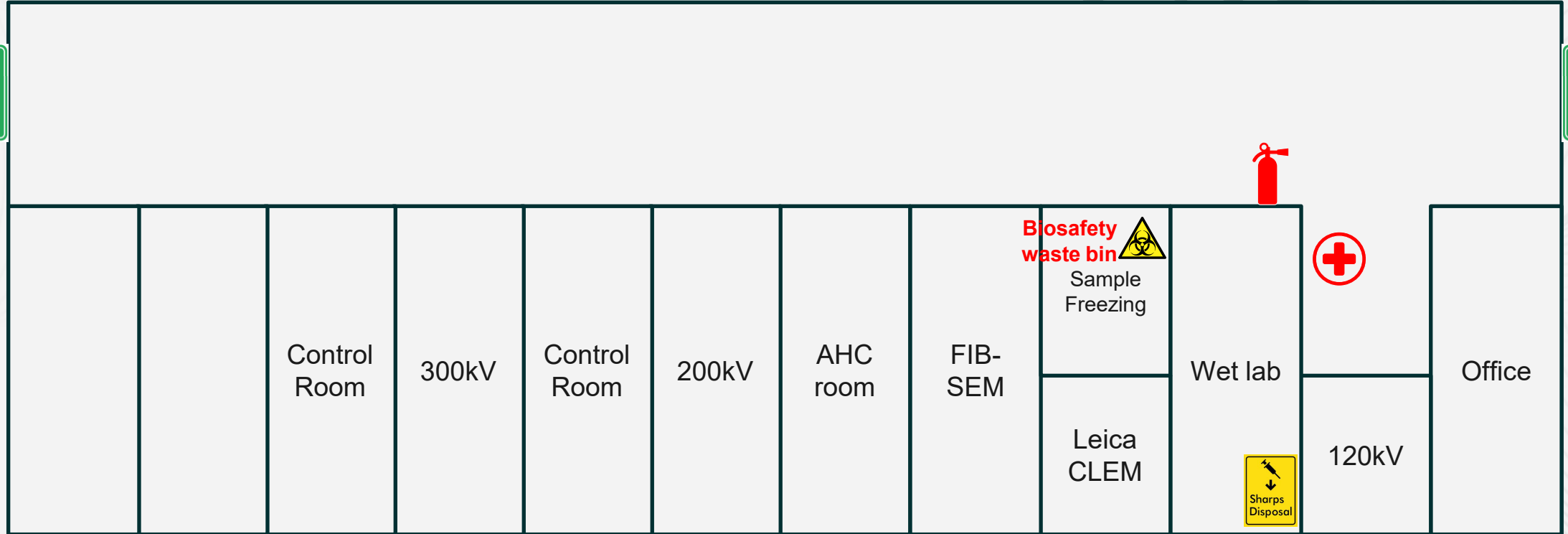
Biological waste bin



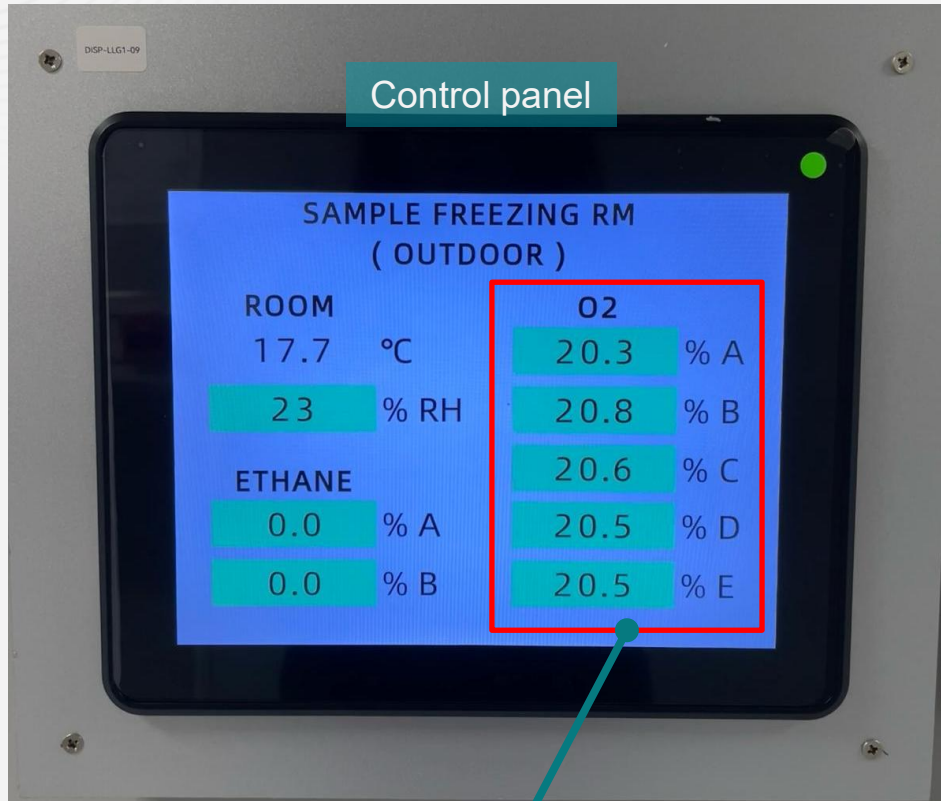
Sharp box



Safety (Floor Plan)

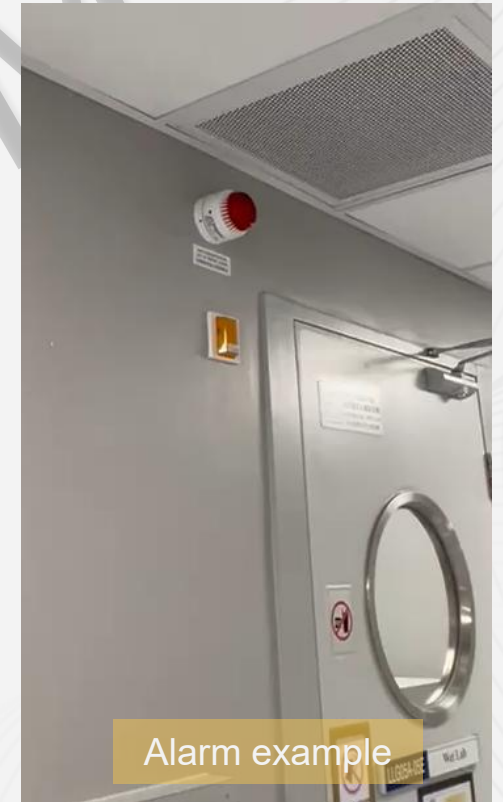
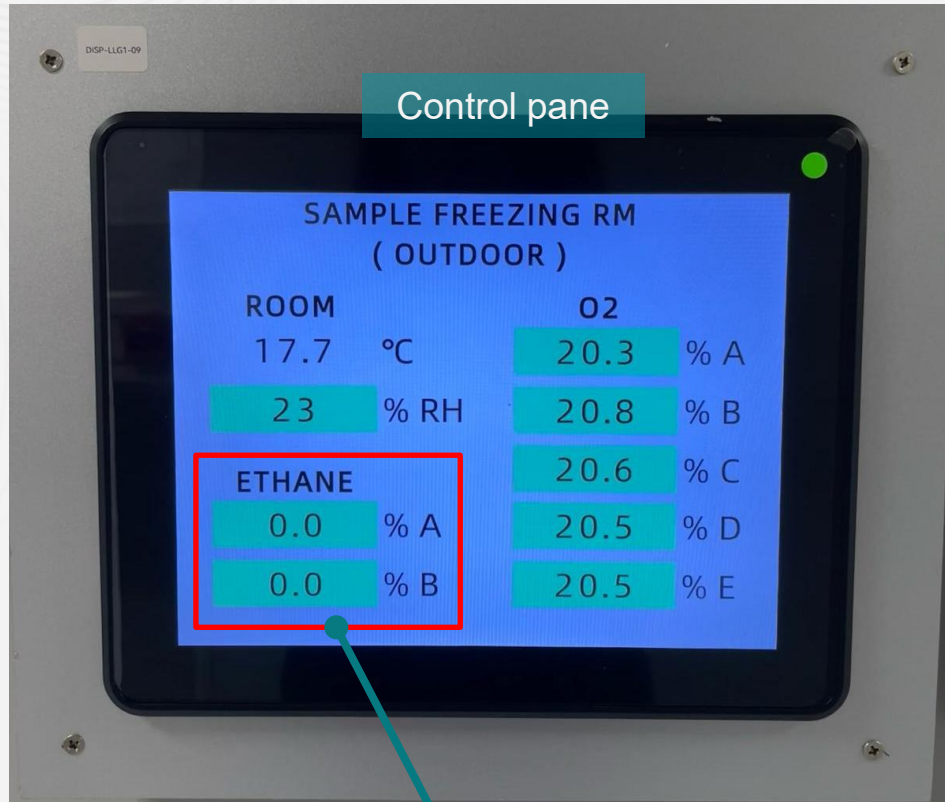


Oxygen Detector



If the O₂ VALUE shows red or alarm in the room, user should leave the room immediately and report it to staff.

Ethane Detector



If the ETHANE VALUE shows in red or alarm in the room, user should leave the room immediately and report it to core staff.

Getting Access to Cryo-EM Lab

1. Submit the following forms

a) Sample Safety Information e-Form (for new PI/ new project)

https://hku.au1.qualtrics.com/jfe/form/SV_4U6w9Xvill3F5gG

b) iLab Registration Form to enquiry.cpos@hku.hk and cc cryoem.cpos@hku.hk

<https://info.cpos.hku.hk/wp-content/uploads/2025/07/iLab-User-Registration-Form-202507.xlsx>

c) Training Application

Submit Training Request in iLab system

2. Training and booking for TwoMP

1. Register your HKU card for access

2. Training session

3. Self-booking (staff support is a must for 1st booking)

4. Email to cryoem.cpos@hku.hk to confirm the availability of staff

5. Transfer data (get a SFTP server from staff if you do not have HPC account)



Sample Information e-Form



iLab Registration Form

Charging

Training	HKU-Med	HKU
1 st training	Free	\$500
Re-training	\$700	\$1000

User should use the instrument **within 2 months** after training.

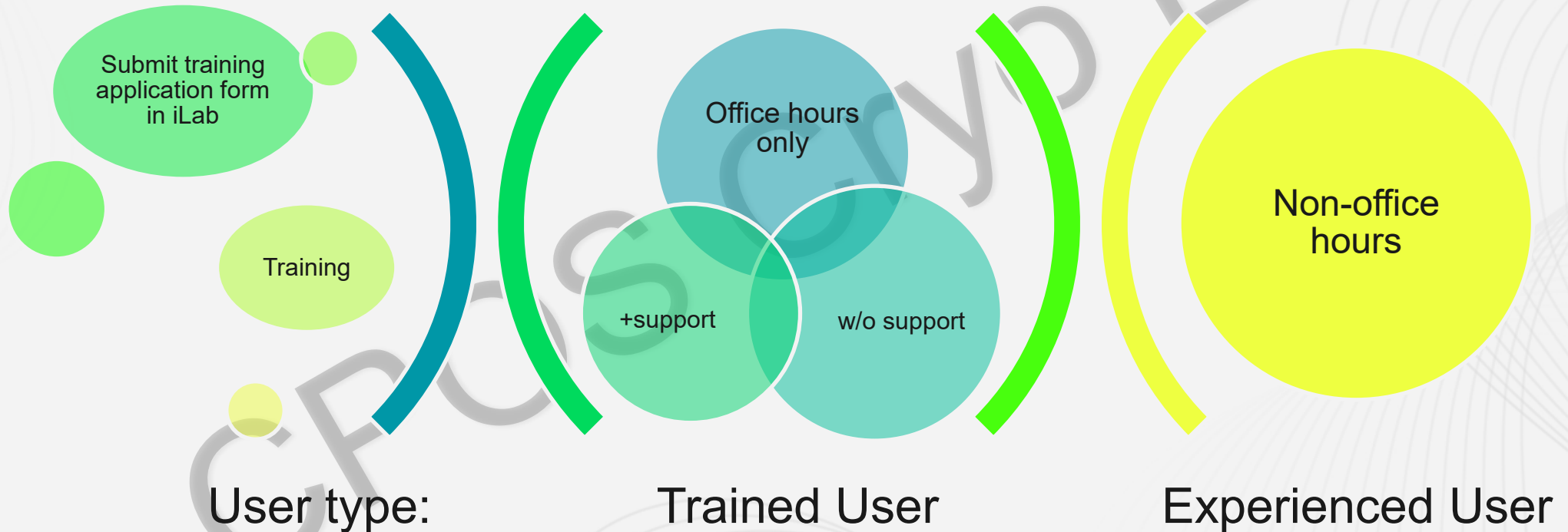
- If user has not booked within 2 months, **re-training** is required before using the instrument.

Usage	Min. usage / Session	HKU-Med	HKU
Refeyn TwoMP Mass Photometer	1 hour	\$100 / hour	\$110 / hour

Charge will be based on **(1) booking** or **(2) usage** whichever longer. Same charge will be applied for no-show (not recommended).

User Type

- Do not transfer your booking session without notifying CPOS staff.
- Fill in correct information (e.g. usage time) in the logbook.





TwoMP

Mass Photometer

Booking Sessions

Office Hour	
Anytime (Min 1 hour)	10:00 am – 5:00 pm

First time user should not book during 12:30 pm – 1:30 pm.

Cancellation Policy

- Before 24 hours → Free cancellation
- Within 24 hours → 50% charge
- Session starts → 100% charge

System Information

Please enter the HKU billing account number

%	HKU billing account number	Amount
1 100.0 %	test	

100.0% Total Allocated

Use the same payment information for all add-on charges

[Invite additional people to this event by email](#)

Please enter a comma separated list of valid email addresses

09:00 PM

System Information

Please enter the HKU billing account number

%	HKU billing account number	Amount
1 100.0 %	test	

100.0% Total Allocated

Use the same payment information for all add-on charges

[Invite additional people to this event by email](#)

Please enter a comma separated list of valid email addresses

09:00 PM



✕

Cancelling this event will result in a cancellation fee of \$150.00

Consumables

Items	Price
Sample Carrier Slide (glass slide) + Sample Well Cassette (gasket)	\$1000 / pack

Must buy whole package.

- 6 measurements can be done for each glass slide + gasket
- Total 60 measurements can be done for 1 pack
- Recommended to put in dry box



Glass slide and Gasket **CANNOT** be reused.

1 Pack contains:

Glass slide x 10



Gasket x 10

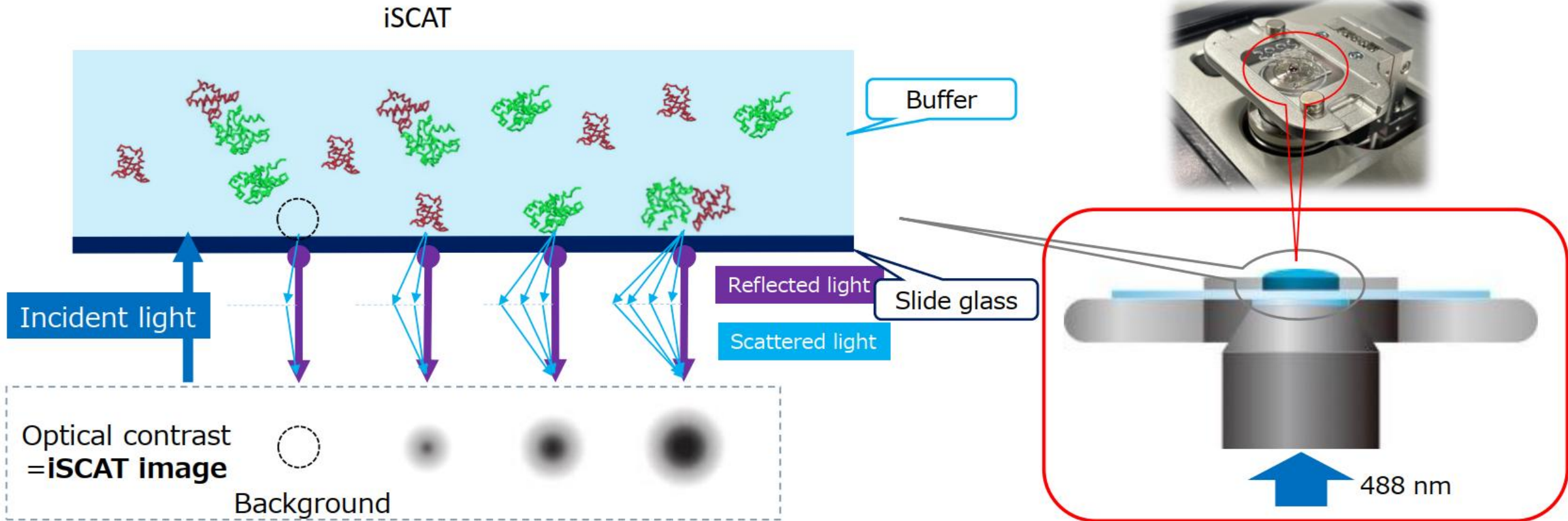


Introduction of TwoMP

Interferometric scattering microscopy

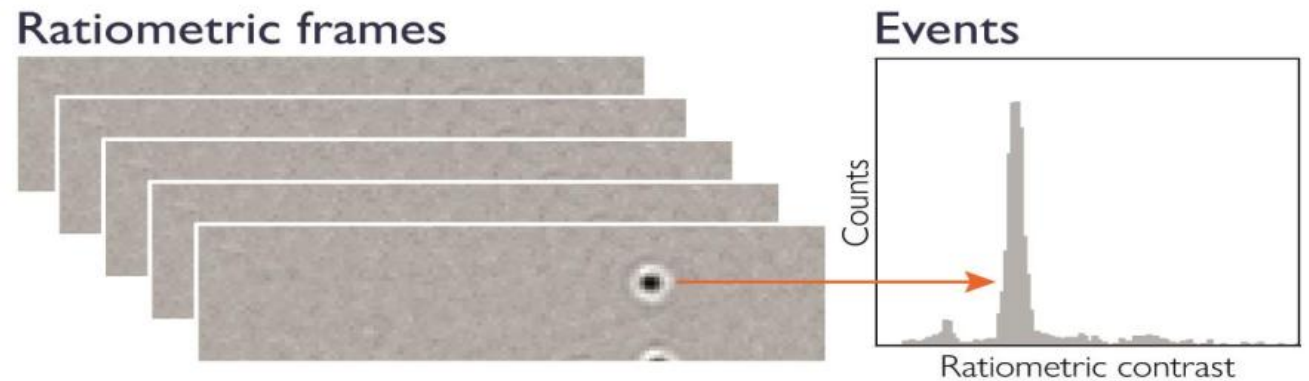
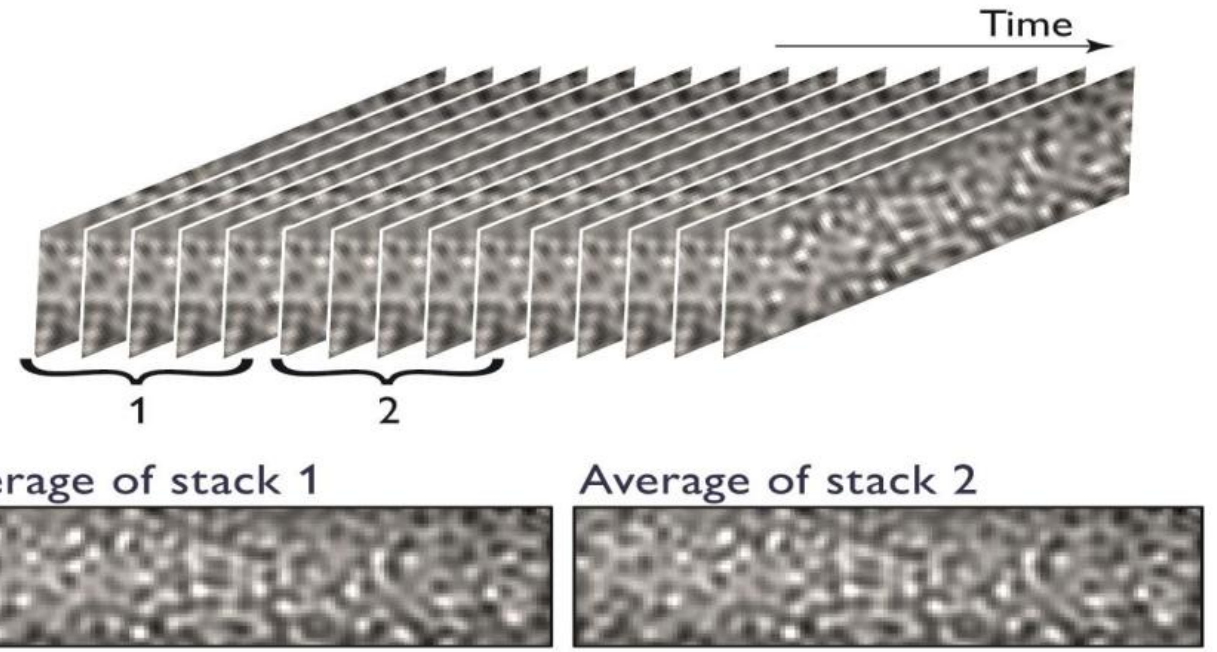
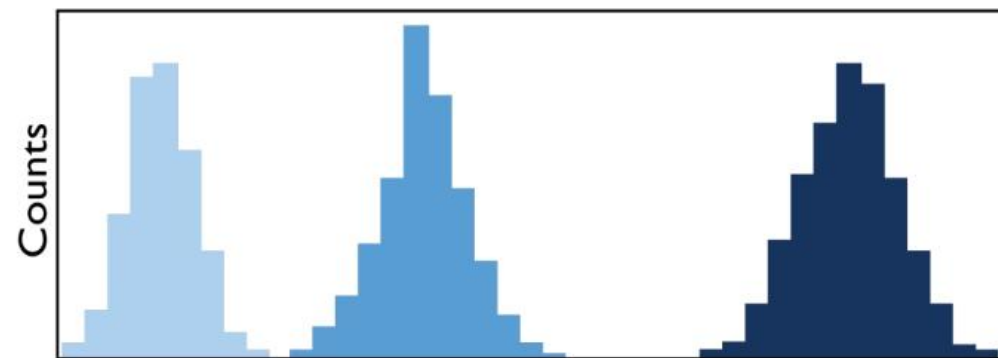
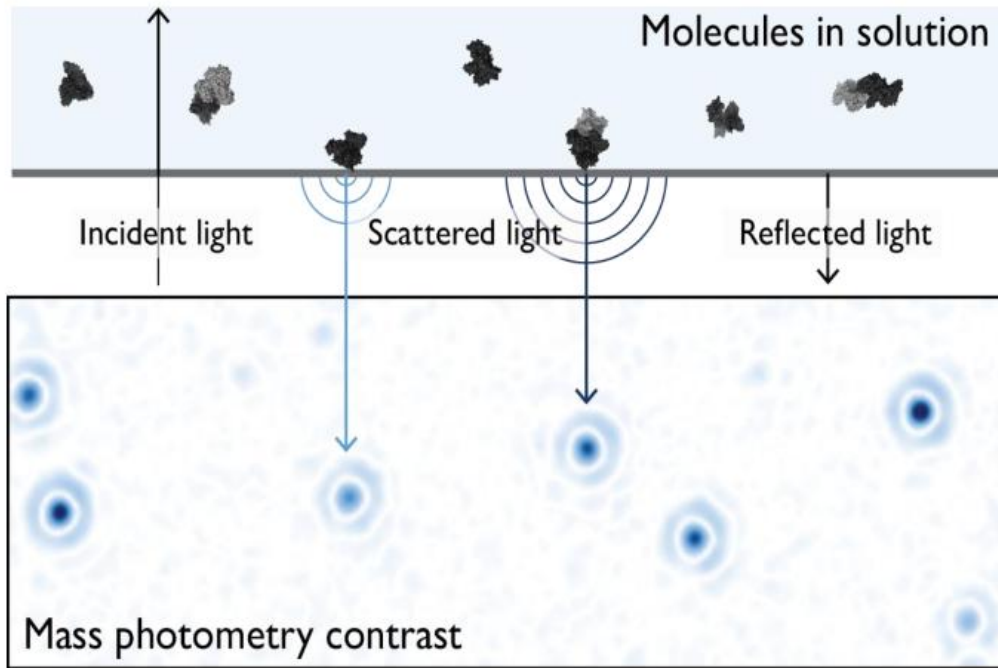


Do not look at the light source directly.



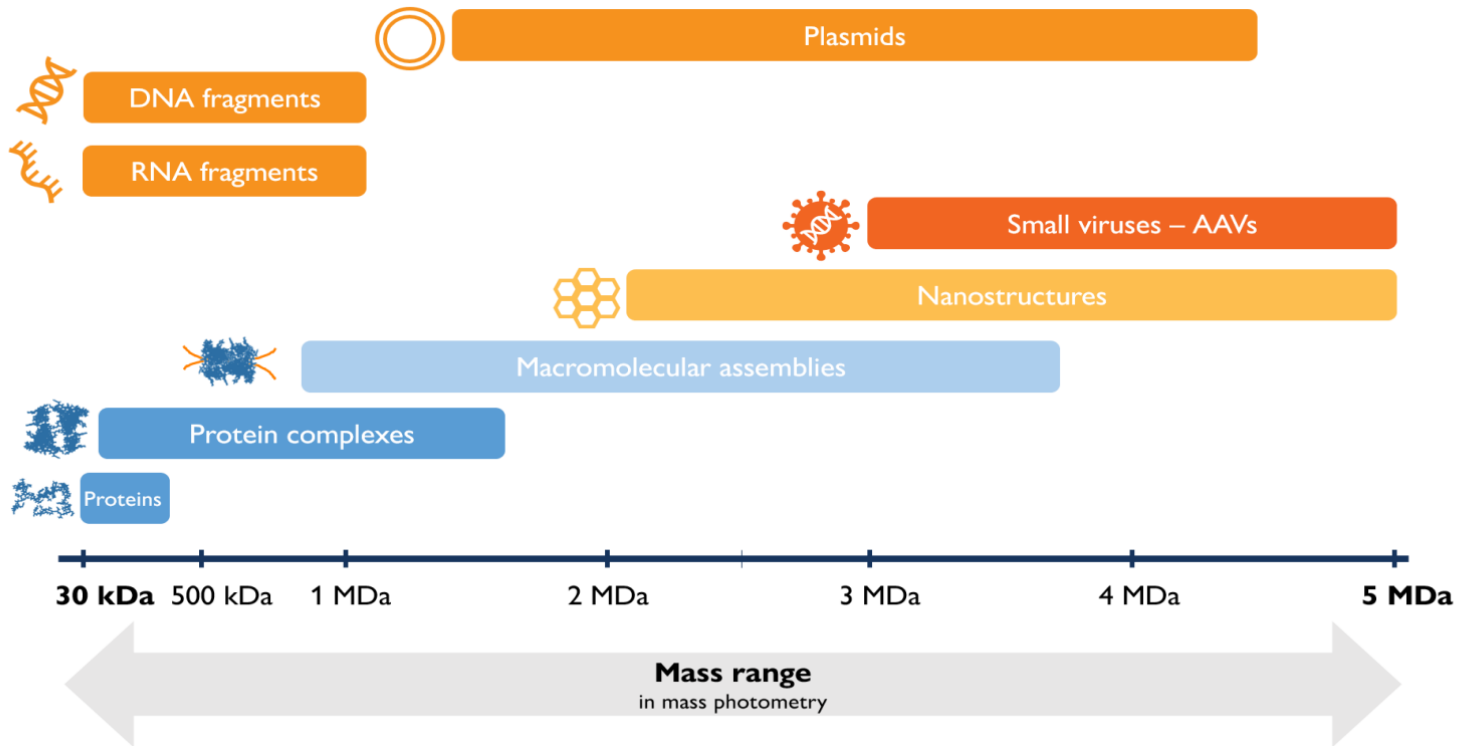
Introduction of TwoMP

Ratiometric analyzing method



Introduction of TwoMP

Mass range of TwoMP



Key Specifications

Refeyn Two^{MP} Mass Photometer Specifications*

Mass range	30 kDa – 5 MDa
Resolution (FWHM)	25 kDa @ 66 kDa 60 kDa @ 660 kDa
Mass precision	± 2%
Mass error	± 5% (single measurement)
Concentration range	100 pM – 100 nM 20 nM (recommend)
Sensitivity:	<< 1 ng of protein
Wavelength:	488 nm
Field of view:	Small: 3 x 11 μm (@ 700 Hz) Regular: 4 x 11 μm (@ 500 Hz) Large: 12 x 17 μm (@ 135 Hz)
Pixel size:	12 nm

* All specifications subject to meeting the installation requirements

Sample ,Calibrant and Buffer

- Bring your own sample, calibrant and buffer.

Detection range of TwoMP system:	30kDa to 5MDa
Concentration of standard	Protein standard marker (1:500) Protein with known mass (~1uM)
Concentration of the sample	20nM is preferred. The range is 100pM to 100nM.
Sample volume	100uL (actually only 10uL is required for 1 well)
Buffer	About 5mL (filtered using 0.22um filter, freshly prepare) <u>Avoid</u> using buffer contains: <ol style="list-style-type: none"> Detergent High concentration (mM) of small molecules Glycerol

Instrument	What to measure to measure	Calibrant	MW (kDa) / Size (bp or bases)
TwoMP OneMP	Small Proteins <90KDa	BAM	56, 112, 224 kDa
	Medium Size 90KDa-1MDa	BAM +TG (separate)	56, 112, 224 + 670 kDa
		Urease	90, 272, 545 kDa
	Larger proteins >500kDa	BAM + TG (separate)	112, 224 + 670 kDa
		Urease + TG (separate)	272, 545 + 670 kDa
	dsDNA	Invitrogen DNA ladder	100, 200, 400, 800, 1200, 2000 bp
	ssDNA	M13MP18 + Phix174	7249 and 5386 bases
	RNA	Agilent ssRNA Ladder	200, 500, 1000, 1500, 2000 bp
NEB ssRNA Ladder		500, 1000, 2000, 3000 bases	

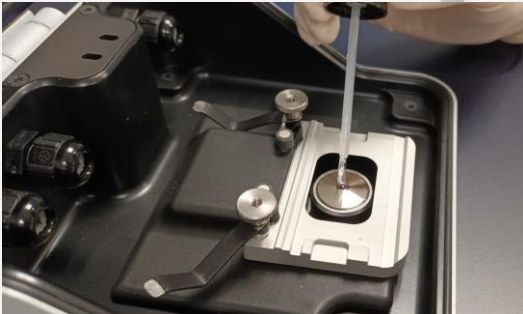


No live virus or bacterial samples allowed for TwoMP.

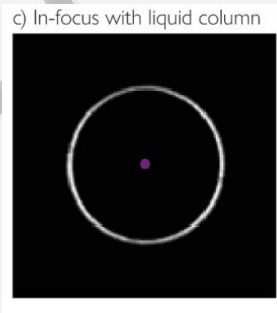
Steps

1. Warm up for 1 hour (done by staff)
2. Initialization
3. Focus
4. Measure
5. Clean up and shutting down TwoMP
6. Sample analysis
7. Data Transfer

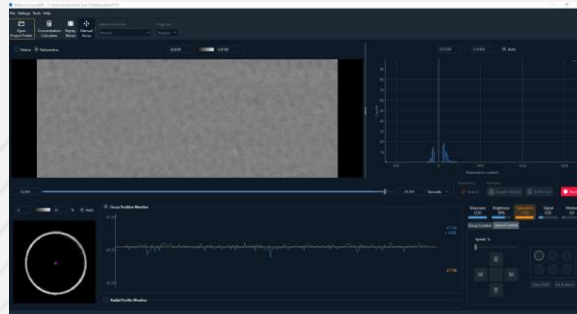
1. Initialization



2. Focus



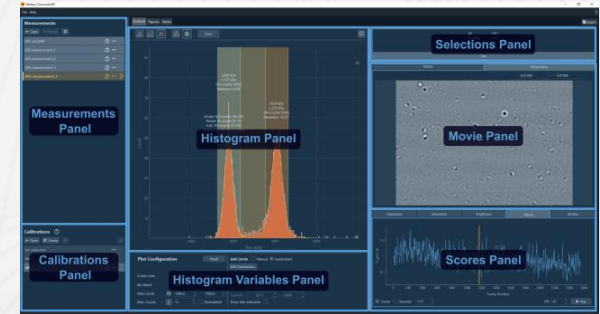
3. Measure



4. Clean up




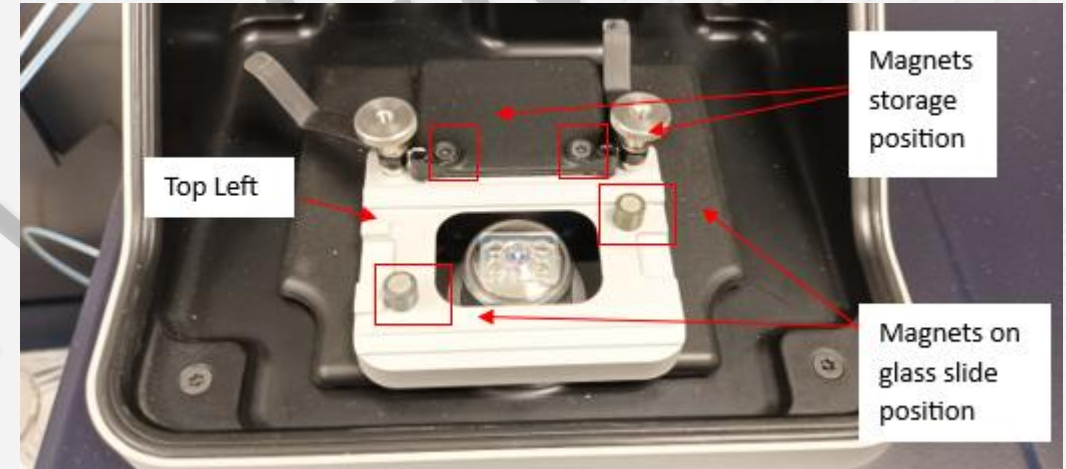
5. Sample analysis



Special Notice

- Discard the glass slide with gasket into the sharp box.
- Ensure that the stage plate is clean and oil free after clean up.
- Remember to transfer your data (using SFTP server or HPC account) to your personal computer.

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1. If you use SFTP server, please make sure you have successfully downloaded the data to your local computer within 14 days.
 2. The local data stored in the computer of TwoMP will be removed on the first working day of each calendar month.



Please handle the magnets with extra caution.
Do not drop them into the machine.