

An introduction to:



Platformfest – 17 Jan 2024

## CRBS is a Research Centre & a Platform:

CRBS's Mission is to exploit the power of structural biology and biophysics to produce the next wave of scientific breakthroughs in:

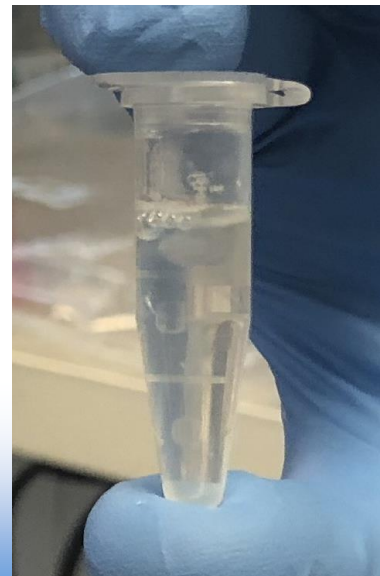
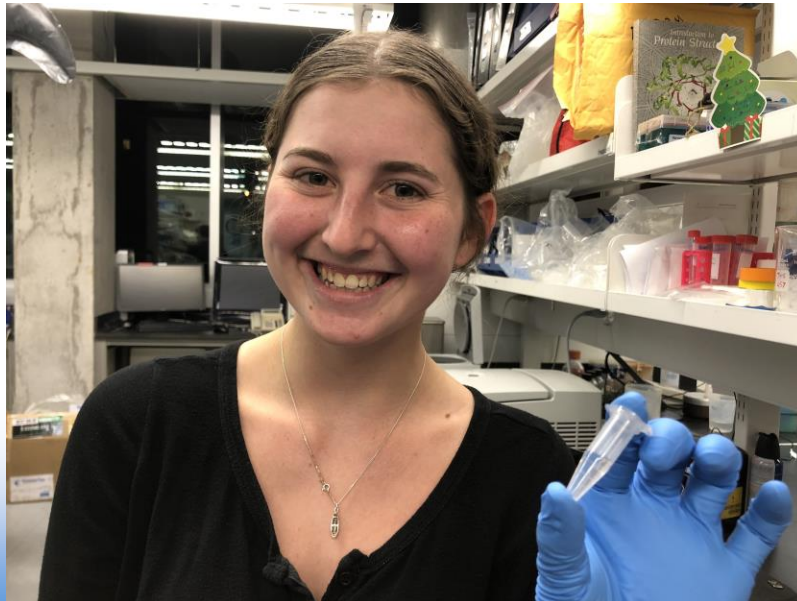
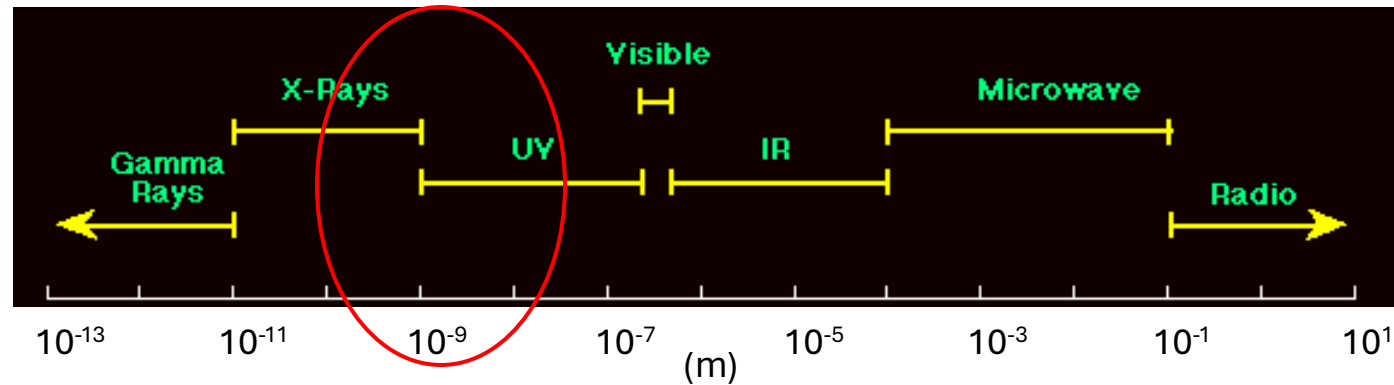
- (i) determining the **molecular basis** of disease and treatments
- (ii) leveraging **biophysical, chemical and synthetic biology** for health

To train a new generation of structural biologists and biophysicists with outstanding expertise in using cross-disciplinary approaches for biomedical research

To make structural biology and its many strengths accessible to the broader biomedical research and health community.

## The Fundamental Problem:

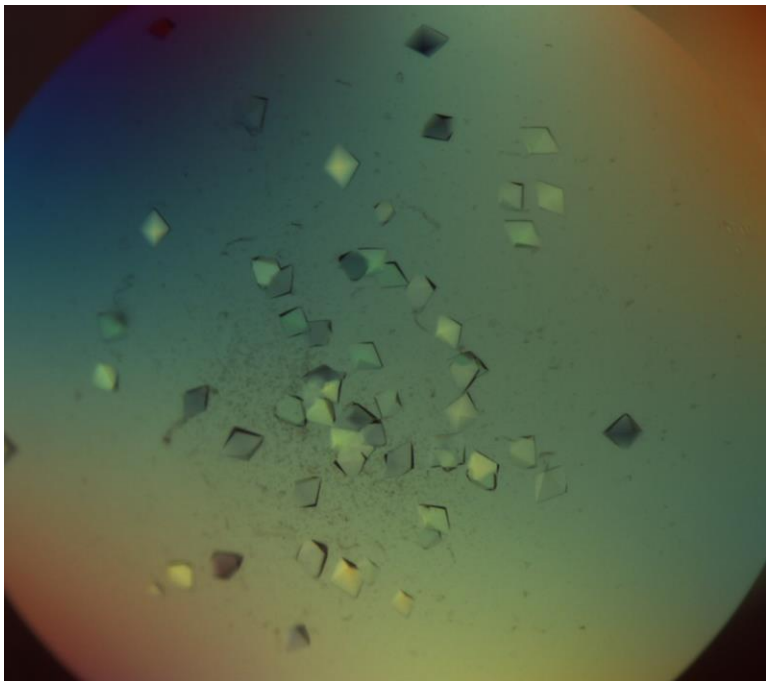
-the molecules that underpin all health and disease are very small and thus not visible even with a very, very powerful light microscope



## Biophysics & Structural Biology to the Rescue!

- What protein/RNA/DNA/complex looks like
- What it binds, how it binds (no biomolecule is an island, entire of itself)

## Equipment to address this Fundamental Problem:

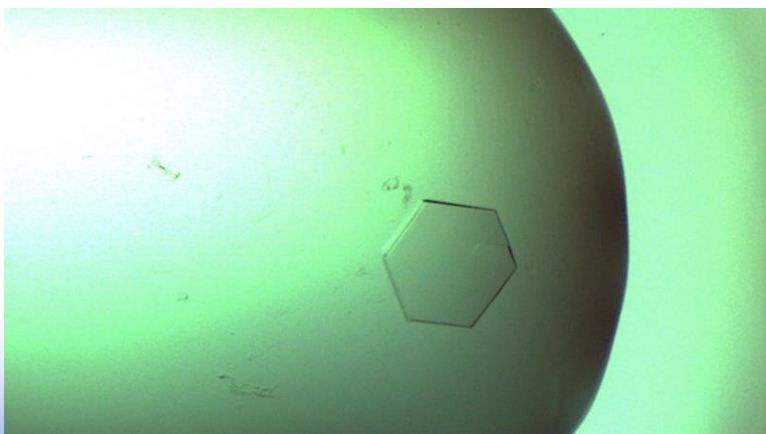


Crystallography 1: High-throughput crystallization hub

Robots

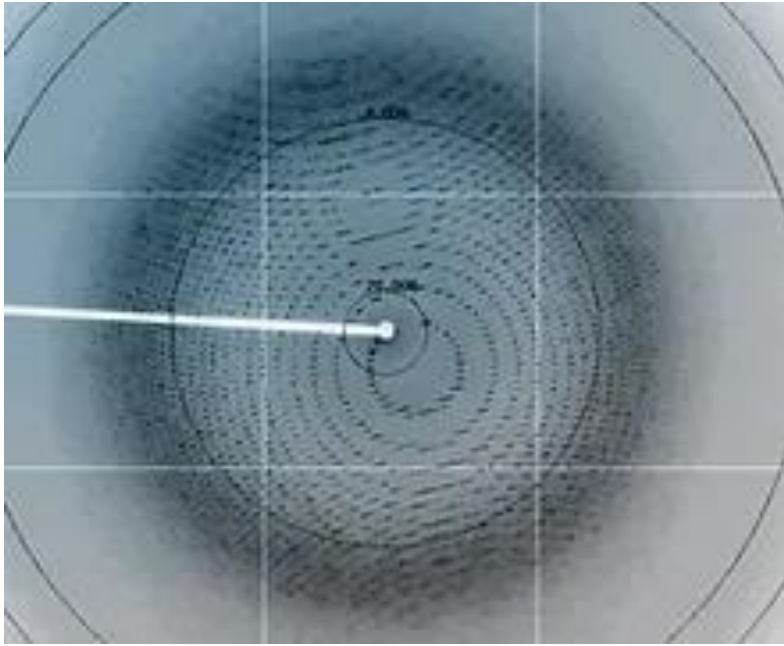


Controlled incubation & UV/vis camera

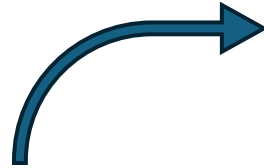
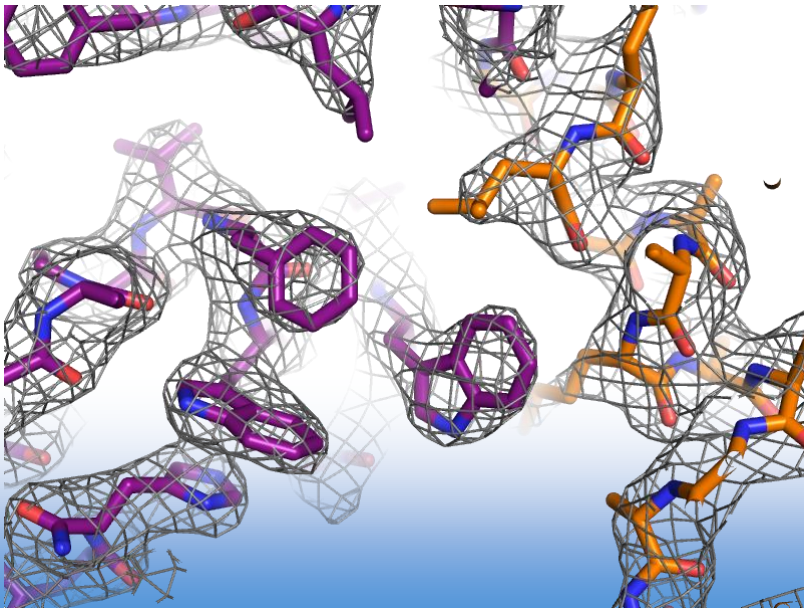




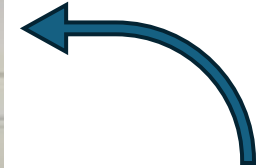
## Equipment to address this Fundamental Problem:



### Crystallography 2: Macromolecular X-ray analysis hub



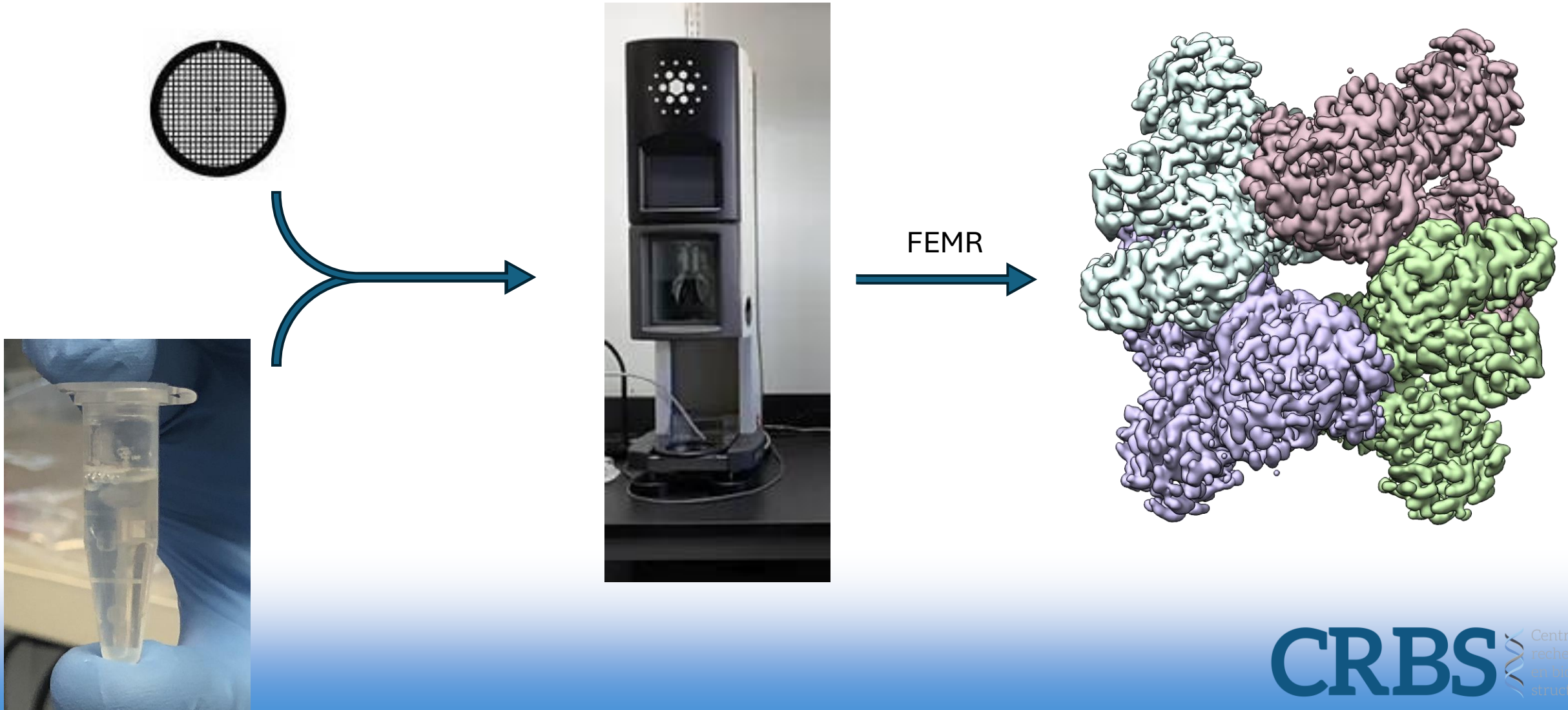
Single crystal  
diffraction  
(CRBS's D6 & RI-  
MUHC's D6):  
High res structure  
& crystal screening



Small angle X-ray  
scattering (SAXS):  
Shape info

## Equipment to address this Fundamental Problem:

EM prep: Vitrobot, carbon coater and glow discharger



## Equipment to address this Fundamental Problem:

NMR: 600 MHz spectrometer with cold probe. Small molecule structure determination and protein dynamics, ligand binding and structure.

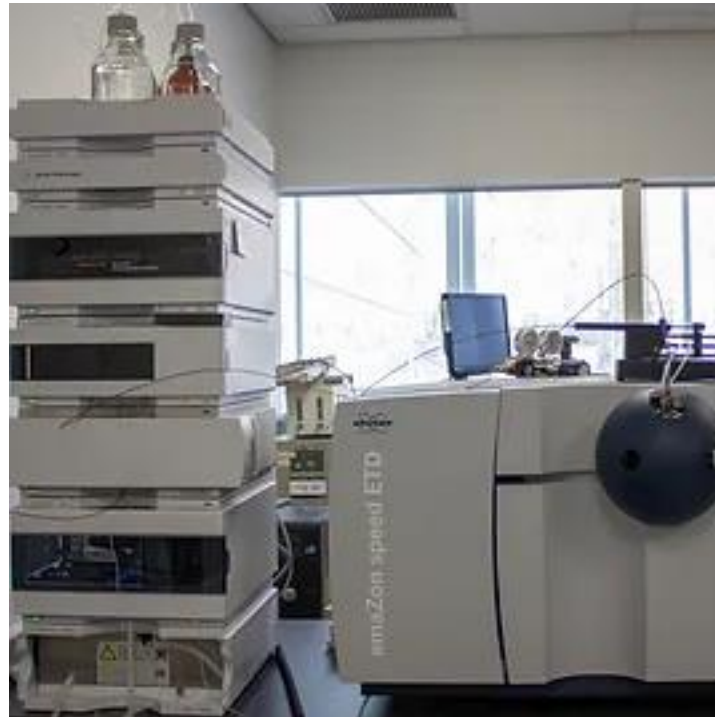


## Equipment to address this Fundamental Problem:

Mass Spec:

Microflex MALDI –TOF: Metabolite, small molecule and small protein detection and analysis.

Amazon Speed ETD ion trap mass – HPLC: intact protein mass and small molecule mass.



These MSs has MS friends in SPR-MS platform



## Equipment to address this Fundamental Problem:

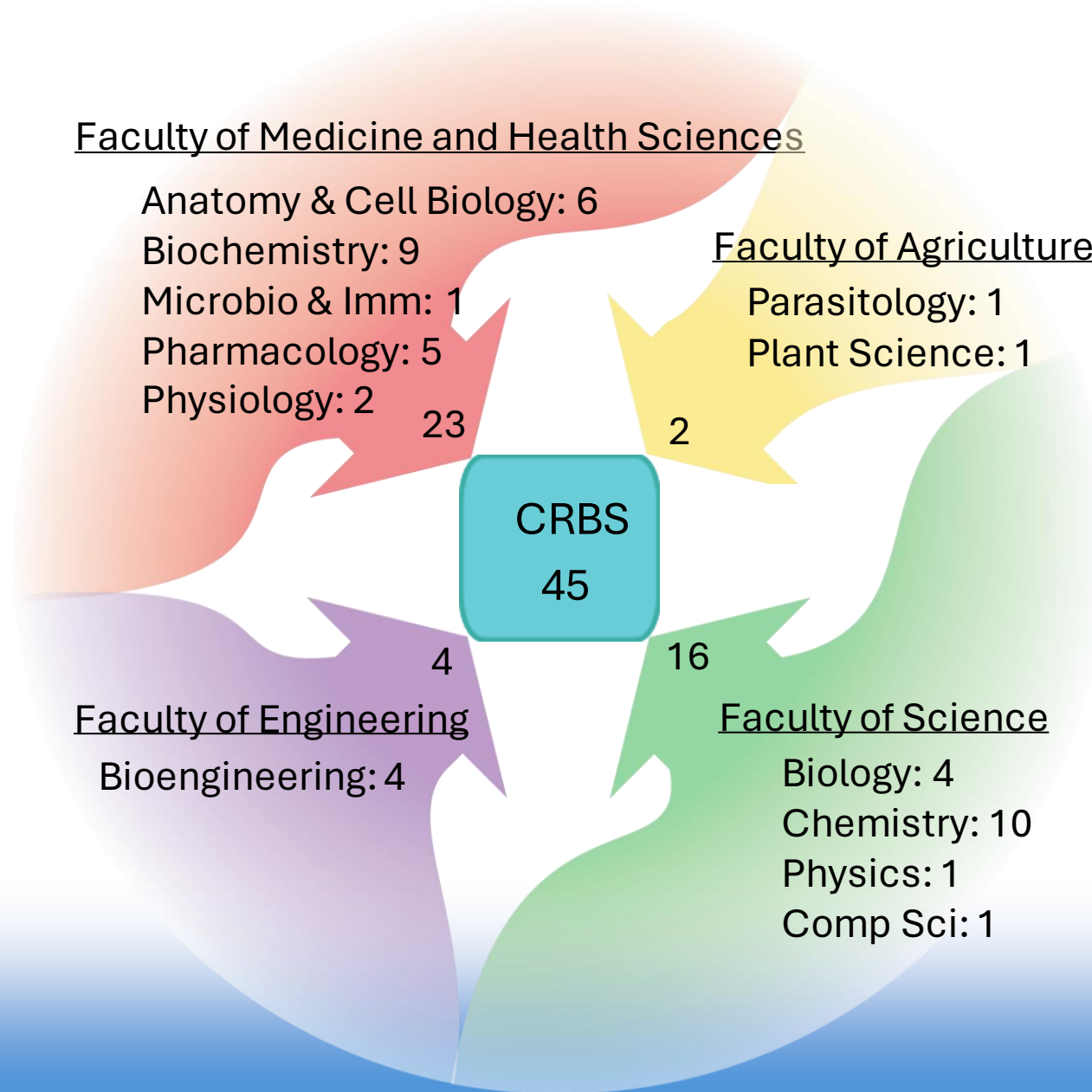
All the Biophysics: Protein-protein interactions platform

- three isothermal titration calorimeters (ITC) for measuring binding affinities,
- multiangle and dynamic light scattering instruments and analytical ultracentrifuges for characterizing shape and oligomeric mass
- a circular dichroism (CD) spectrometer for determining protein secondary structure
- a Biocomp Gradient Station for purification, crosslinking and molecular density analysis
- several fluorometers for measuring binding affinities and protein stability.



These MSs has MS friends in SPR-MS platform

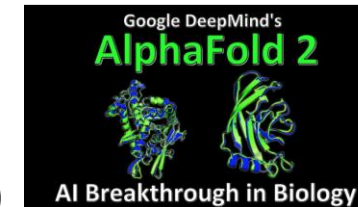
# CRBS PIs / Core users of Biophysics Platforms



# Linking to others

We can:

- Facilitate your research if you would like to get into biophysics
- Collaborate on a PI-PI level (funding available! RI-MUHC, MI4, GCI)
- Put on joint events with your Centre / Platform
- Access the power of AlphaFold (one-on-one, or group)
- (Place your idea here!)



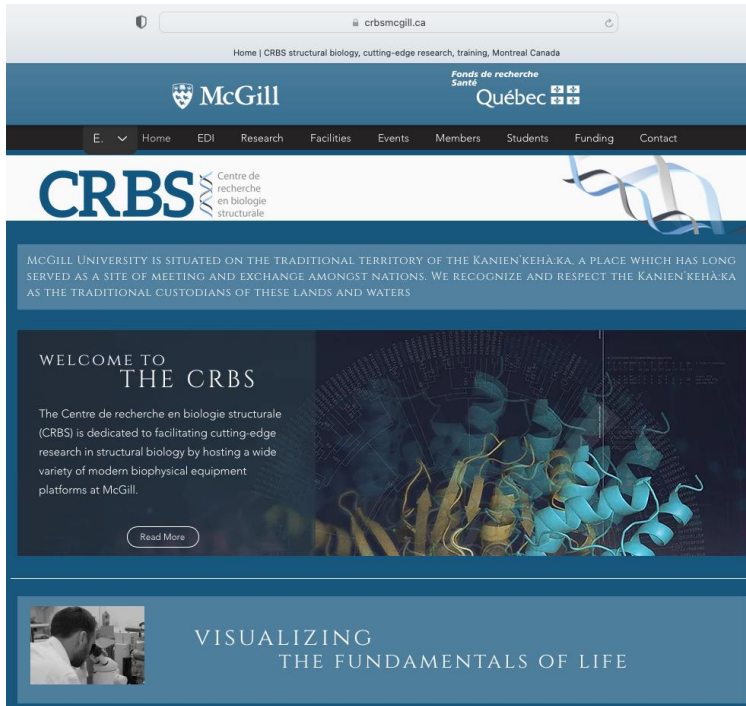
**CRBS** Centre de  
Recherche  
en Biologie  
Structurale

**Atelier du CRBS**  
Exploration des structures de protéines avec PyMOL et AlphaFold2

Instructeur: Jean-François Trempe  
Date: vendredi, 17 juin 2022, 13-16h  
Session virtuelle sur Zoom

Le Centre de Recherche en Biologie Structurale (CRBS; <https://fr.crbsmcgill.ca/>) de l'Université McGill a le plaisir d'annoncer la tenue d'un atelier d'exploration des structures biomoléculaires, qui aura lieu le **17 juin 2022 à 13h**. Ce cours intensif de 3h sur Zoom, en français, est gratuit et

www.CRBSMcGill.ca



## How to find us



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### CRBS Directorship:

Martin Schmeing, Director (me!)  
Christopher Thibodeaux, Associate Director  
Natalie Zeytuni, Associate Director

Plus an awesome exec council and student body



Home CRBS Floor: Fourth Floor Bellini