

# Introduction to The Rockefeller University Resource Centers



Overview for New Graduate Fellows  
September 11, 2019

# The Role of Core Facilities

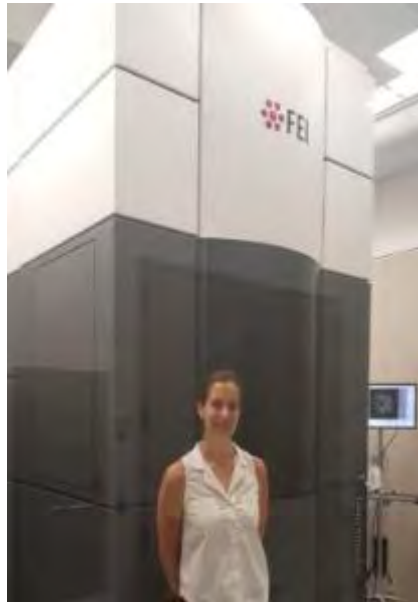
- Ensuring and encouraging access to expertise, equipment, services and products that are beyond the financial or technical means of most individual laboratories
- Supporting research in a cost effective and scientifically effective manner
- Improving competitiveness for researchers and the University
- Providing continuity of staffed expertise
- Training and education in specialized methods and technologies
- Serving as scientific brokers both internally and externally

# Cryo-Electron Microscopy Resource Center

Microscopy Suite: CRC B13  
Phone: 212-327-7282



Mark Ebrahim  
Senior Staff Scientist  
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Johanna Sotiris  
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Honkit Ng  
Research Support Specialist  
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# Cryo-Electron Microscopy Resource Center

200 kV

**Talos Arctica**



300 kV

**Titan Krios G2 – “Krios 1”**



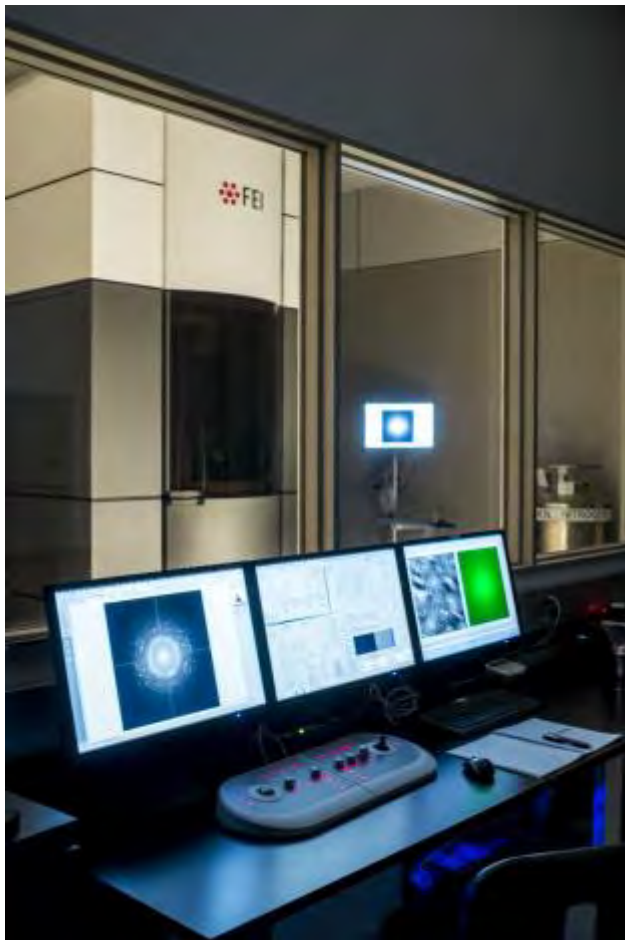
**Titan Krios G3i – “Krios 2”**





# Cryo-Electron Microscopy Resource Center

## FEI Titan Krios G2 (“Krios 1”)



300kV Transmission  
Electron Microscope

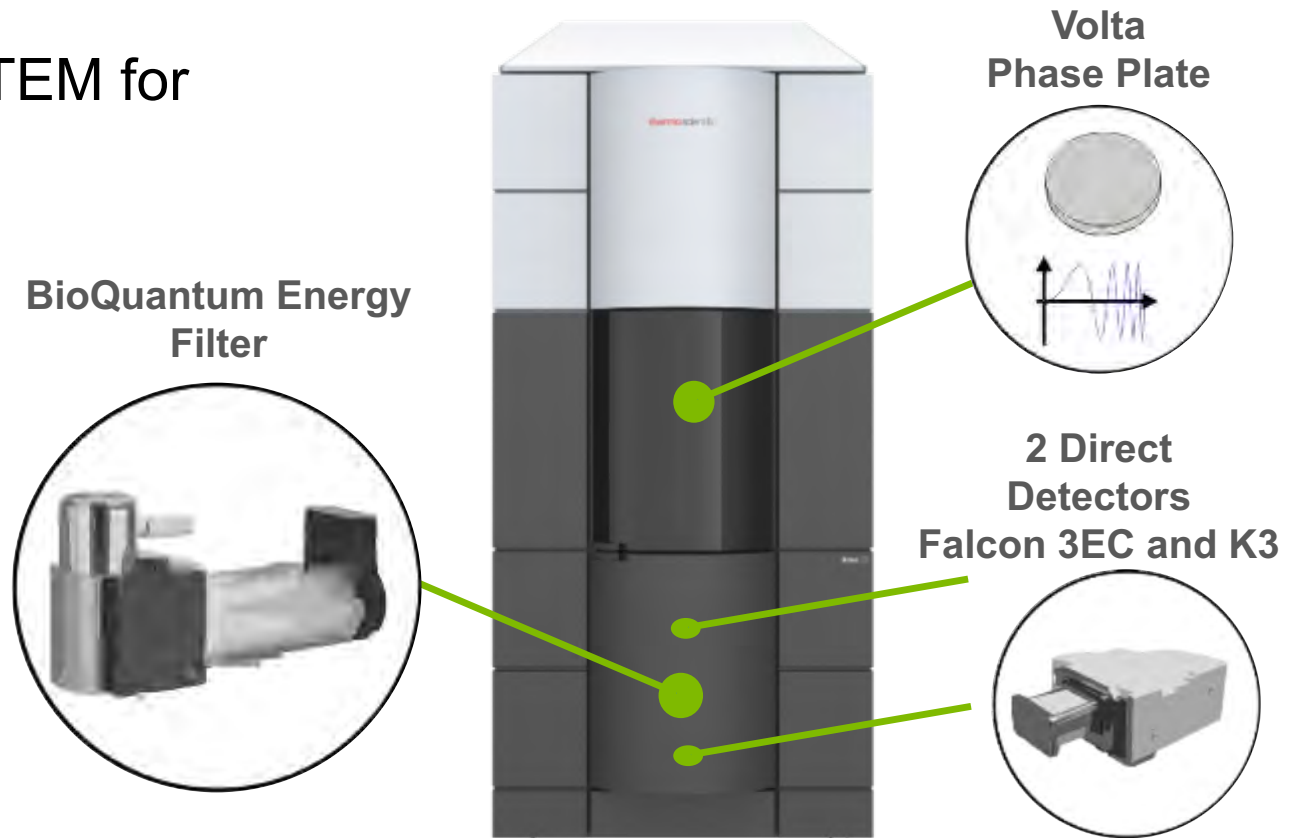
1.0 Angstrom Resolution

Equipped with Gatan K2  
Summit direct electron  
detector



# Krios 2 has a configuration suitable for Cryo Electron Tomography of Eukaryotic Cells

- Krios 300 kV Cryo-TEM for
- Structural Biology
- Cell Biology



# Aquilos

## Cryo Focused Ion Beam Scanning Electron Microscope (FIB/SEM)

- For optimal sample preparation for high-end cryo-electron tomography
- Cryo-FIB milling of vitreous specimens
- Produces cryo-lamellae for the cryo-tomography workflow

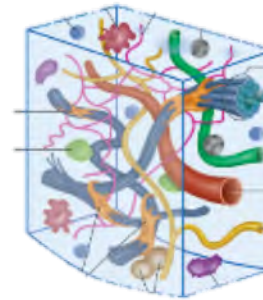
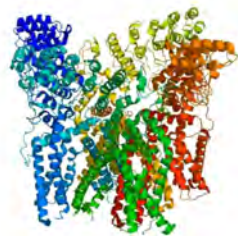


Protein coding Gene

Structure of  
folded Protein

*Function of  
Protein in cells*

*Cellular organisation  
in tissue*



NextGen Sequencing

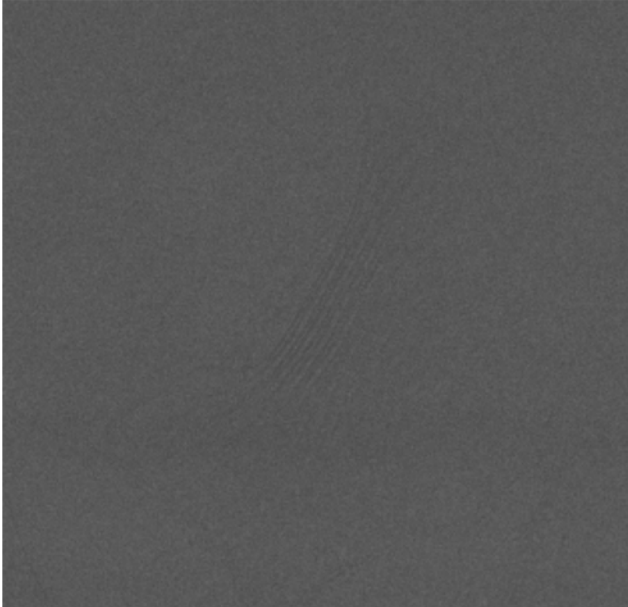
SPA

*Cryo-Tomography*

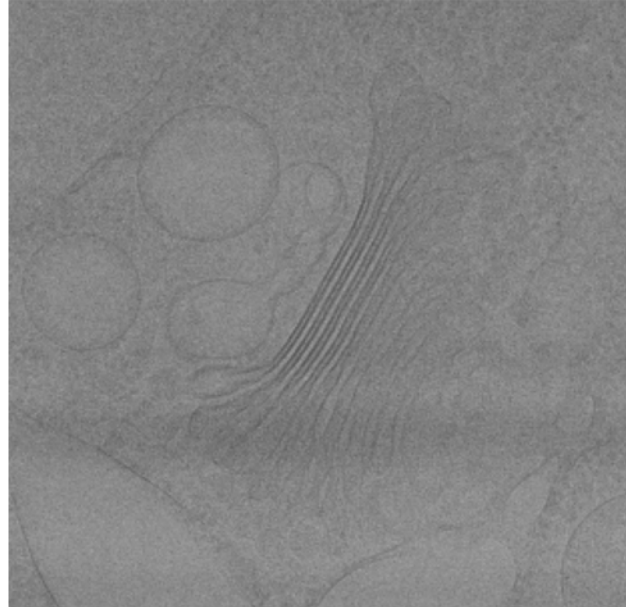
*Large Volume Analysis*

Illustration: Chuck Twiss (ESD)

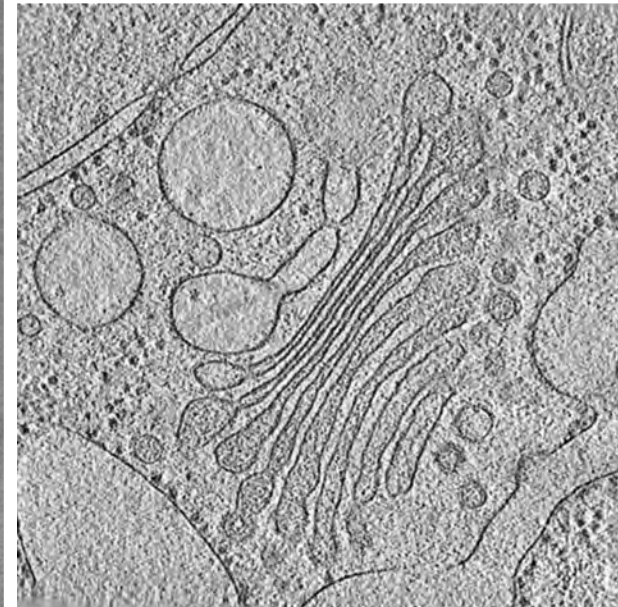
Single frame



Tilt image projection



Reconstructed tomogram



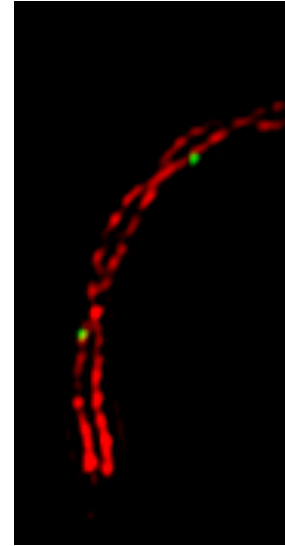
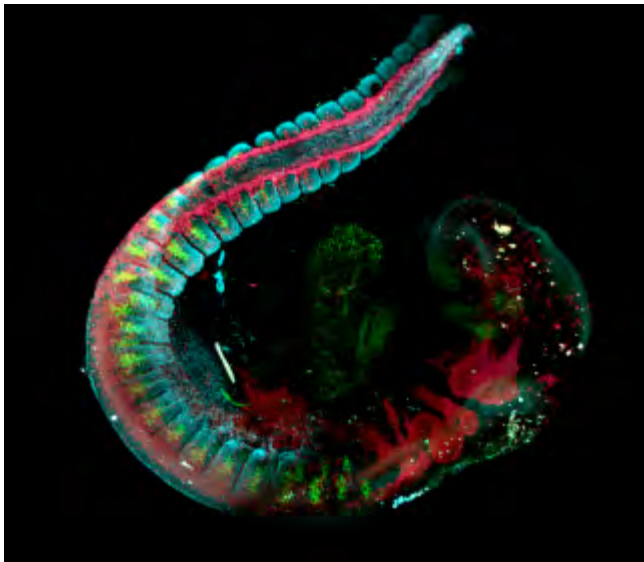
**Thermo Scientific Titan Krios**  
+ Volta phase plate, K2 detector  
Spot size 7, 13 frame/s, 42kMag, total dose:  $\sim 60 \text{ e}/\text{\AA}^2$ , Defocus =  $-0.5 \mu\text{m}$   
 $\sim 180\text{-nm}$  thin cryo-FIB lamella

Bykov et al., eLife 2017



# Bio-Imaging Resource Center

Advice on microscopy and specimen preparation  
Training on the microscopes  
Training in image analysis  
Initial consultations and tours  
Possible collaborations



Widefield

Light Sheet

TIRF

Super-resolution

Confocal

Spinning disk confocal

Deconvolution

Multiphoton

FRET

FLIM

FCS

Laser microdissection

You can even climb inside your own 3D data sets with our Virtual Reality system...





# Staff of the BIRC

(DWB 201-203)

<http://inside.rockefeller.edu/bioimaging/>



**Alison**

➤ **Alison North (Senior Director) - Ph.D. in Cell Biology** - any microscope I can get my hands on in between doing the less exciting administrative stuff, plus OMX specialist;

➤ **Christina Pyrgaki (Senior Research Support Specialist) - Ph.D. in Molecular Biology** – all types of microscopy, multiphoton and light sheet specialist;

➤ **Carlos Rico (Research Support Specialist) - Ph.D. in Chemical Biology** – all types of microscopy, FCS specialist;

➤ **Tao Tong (Research Support Specialist) - Masters in Computing and in Biochemistry/Molecular Biology** – systems administration, image analysis, microscopy;

➤ **Kate Cialowicz (Research Support Specialist) – Ph.D. in Biology**, with an emphasis in super-resolution microscopy;



**Christina**



**Carlos**



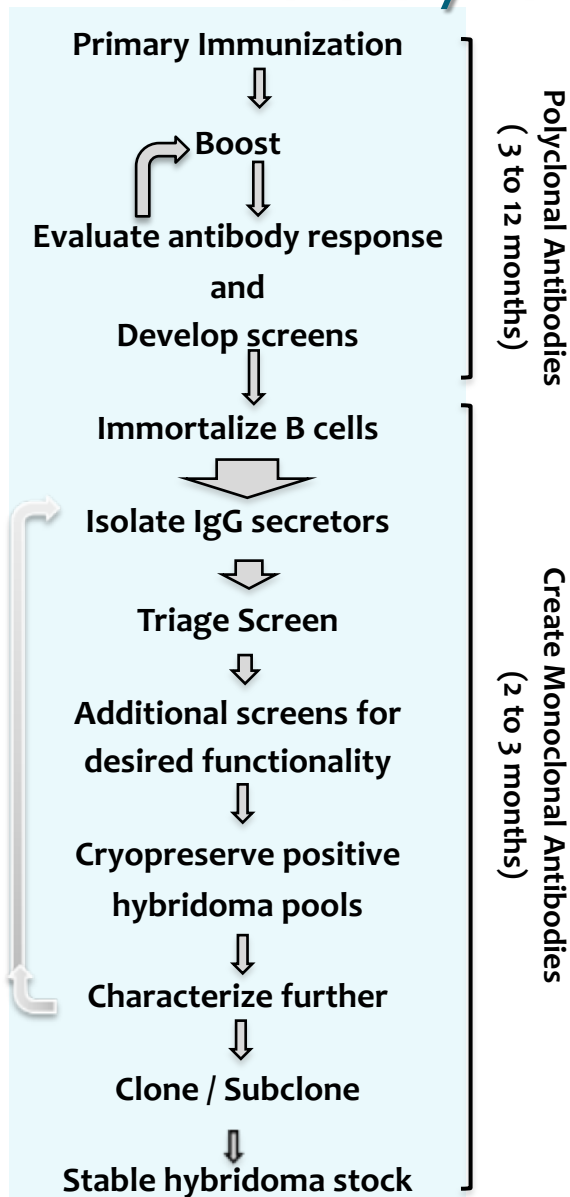
**Kate**



**Tao**

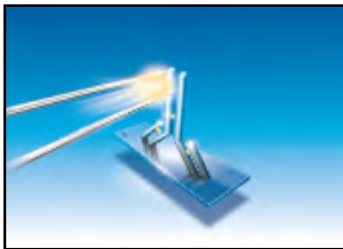
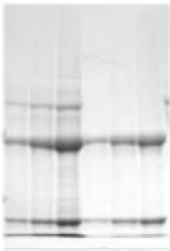
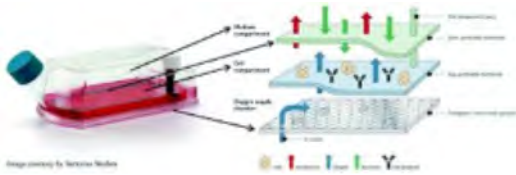
# Antibody and Bioresource Core Facility

## Custom MAb Development



- Comprehensive project design and management
- Generate robust humoral response
  - Immunogens: Cells, protein, peptides, and other haptens
  - Hosts: Mice (WT/KO), hamsters, and rats
- B cell immortalization (10's of thousands of hybridomas)
- Isolate IgG secreting hybridomas (100's pf hybridomas)
- Screen hybridomas by ELISA
- Cryopreserve antigen specific hybridomas
- Clone hybridoma cultures to establish stable lines
- Advise on further characterization and validation

# Antibody and Bioresource Core Facility



## Monoclonal Antibodies

### ○ *In vitro* production

- Conditioned media
- Large scale productions in bioreactors

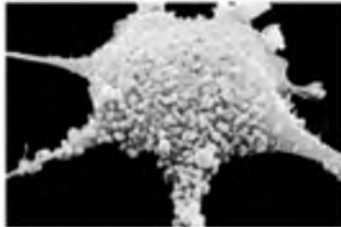
### ○ Purification

- milligrams to grams at > 95% purity

### ○ Modification

- Conjugation to fluorophores, HRP & biotin
- Fragmentation into Fab & F(Ab')<sub>2</sub>

# Antibody and Bioresource Core Facility



## Mycoplasma Testing



## Cell Line Distribution Service

- Alleviates the research lab the work associated with distributing cell lines (e.g. MTA, validation, shipping logistics)
- Informational and physical repository for published cell lines
- Hybridomas & cell lines (e.g. tumor and stem cell)

# Antibody and Bioresource Core Facility



Anne Mui

Lou Mattera

Greg Davis

Syeda Rizvi

Michelle Riley

Frances Weis-Garcia

Bronk 415  
x 7030

ZRC 1553  
646-888-2331

[skiabcf@mskcc.org](mailto:skiabcf@mskcc.org)  
[macfwebext.mskcc.org](mailto:macfwebext.mskcc.org)

# Electron Microscopy Resource Center

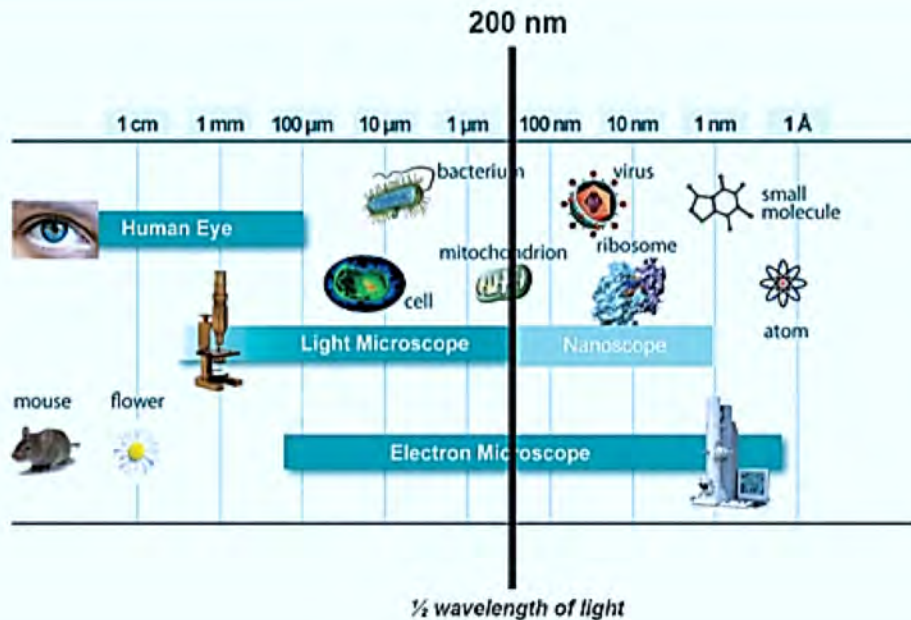


Hilda Amalia Pasolli, Ph.D.  
Director



Nadine Soplop, Ph.D.  
Senior Research Support  
Specialist





- Sample preparation for transmission and scanning Electron Microscopy
- Development of EM protocols according to the scientists needs
- Immuno-labeling (colloidal gold, HRP, APEX)
- Correlative light-electron microscopy (CLEM)
- Training in the use of equipment and techniques

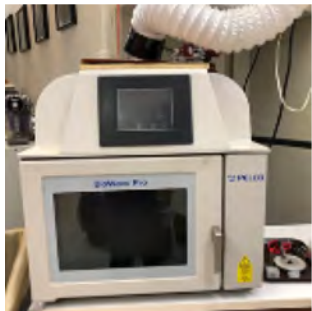
**We can do EM for a variety of experimental models!**





# How do we do EM?

- Chemical fixation
- High pressure freezing followed by freeze-substitution
- Microwave fixation and embedding
- Semi-thin and ultrathin sectioning
- Critical point drying
- Metal coating
- TEM and SEM imaging



Microwave

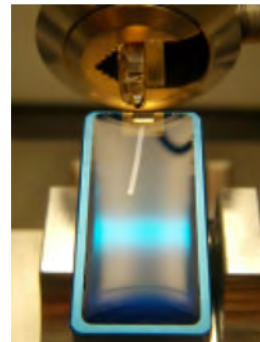


High Pressure  
freezer



Freeze-  
substitution

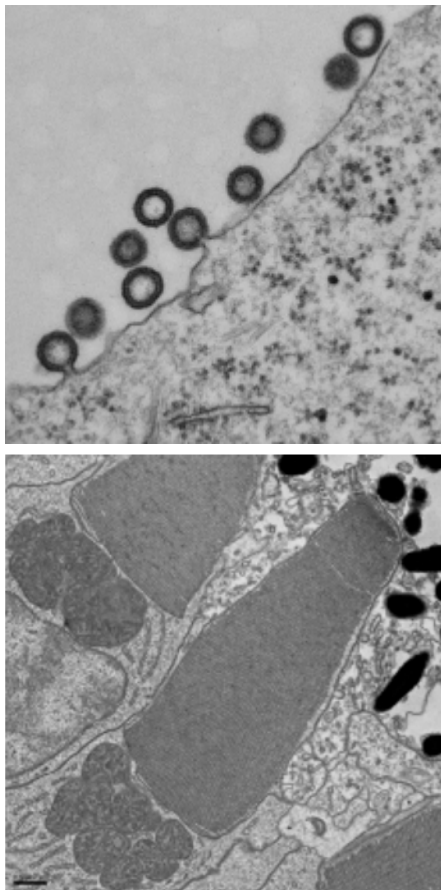
Ultramicrotome



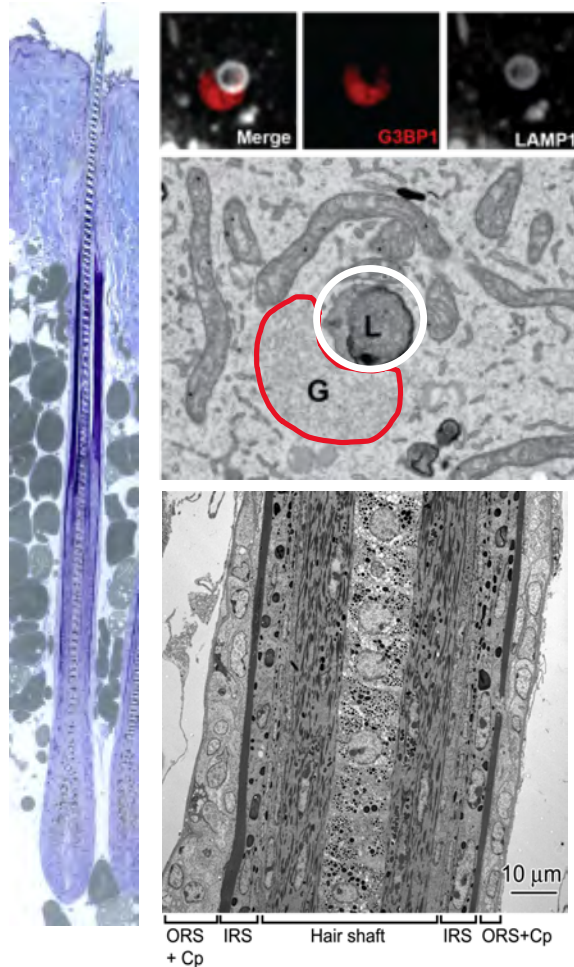
Diamond knife

# Electron Microscopy Resource Center

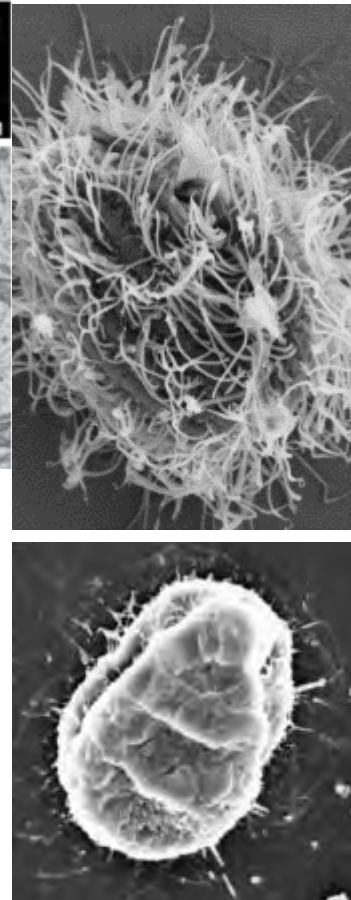
TEM



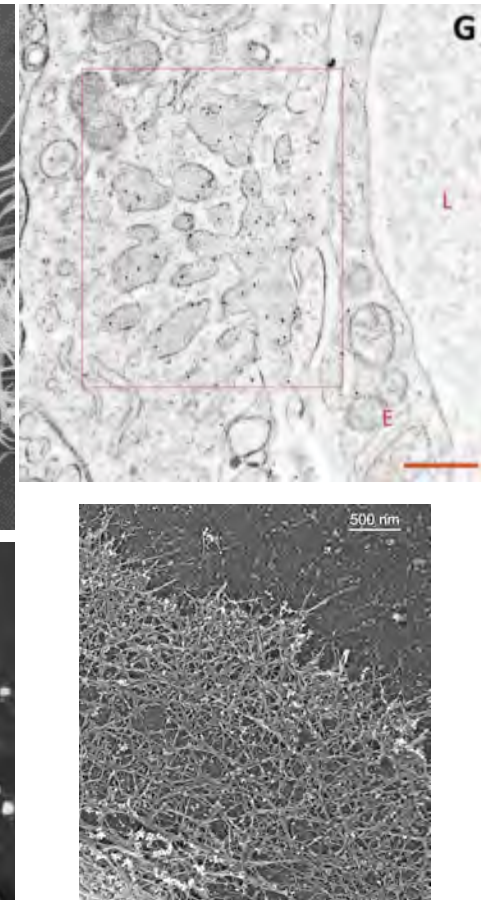
Correlative LM-TEM



SEM



Immunogold



# Flow Cytometry Resource Center (FCRC)

<http://www.rockefeller.edu/fcrc/>

DWB 205 - DWB 211



## ➤ Svetlana Mazel

- Director
- [mazels@rockefeller.edu](mailto:mazels@rockefeller.edu)
- #7656



## ➤ Songyan Han

- Research Support Specialist
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## ➤ Stanka Semova

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- #7657



## ➤ Alena Keprova

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- #7657



## ➤ Samer Shalaby

- Research Support Associate
- [sshalaby@rockefeller.edu](mailto:sshalaby@rockefeller.edu)
- #7657

## ➤ Brandon Yoo

- Information Technology
- Senior Computer Support Specialist





# FCRC Services

## On Equipment

- **Staff-operated equipment at FCRC**
  - Cell sorting on three BD FACSria (cell sorters)
  - Data acquisition on the ImageStream-X (imaging flow cytometer)
- **All the Flow Cytometry equipment at FCRC**
  - Maintenance
  - Quality control testing
  - Troubleshooting and minor repairs
  - Communication with the vendors for appropriate service
    - Service request
    - Post-service follow-up
  - Data management
- **Equipment and computer/programs upgrades and acquisitions**
  - Proposals
  - Installations with sufficient quality controls



## Education and Training

- **“Beyond the Basics” Flow Cytometry Class**
  - FCRC Classes in groups of 5-15 people
  - Pre-scheduled approximately once a two months



- **Instrumental Training “Hands-on”**
  - Personal sessions sc based on re
  - Hands-on trainings
  - Help with instrument setup and troubleshooting



- **Software Training (Introduction or Troubleshooting)**



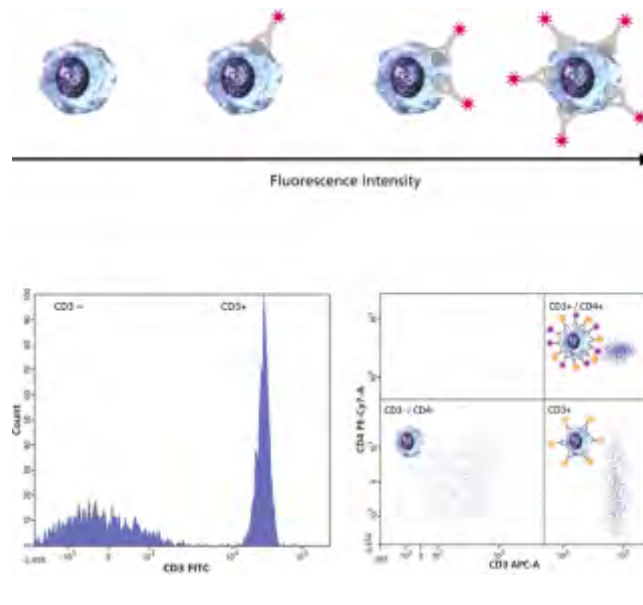
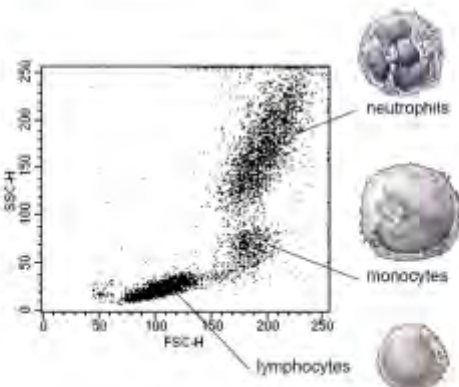
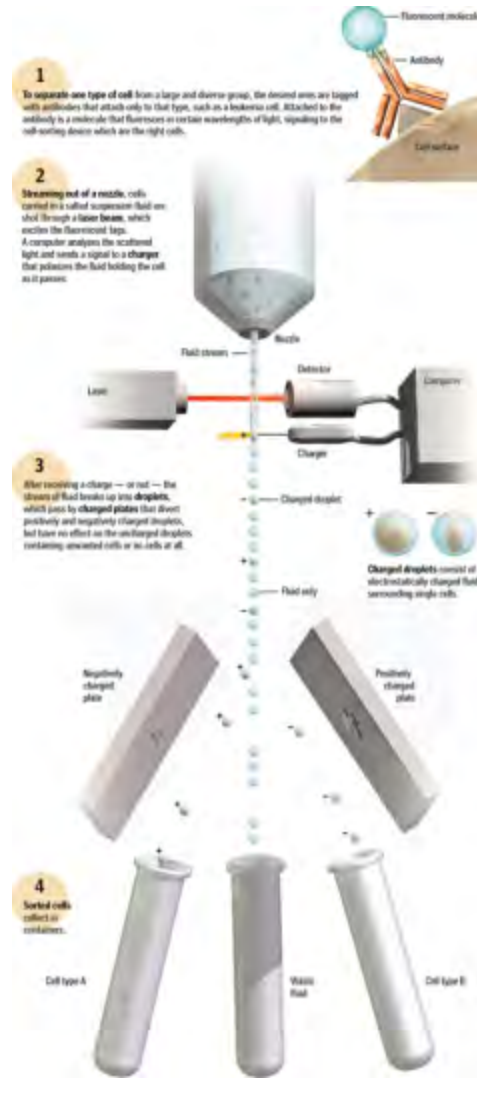
## Consultation and Help

- **Consultations**
  - Experimental Design
  - Pre-Sort
  - Pre-ImageStream
  - Pre-Hands-On
- **Troubleshooting**
  - On experimental design
  - On the instruments
- **Help and Assistance with:**
  - Data analysis
  - Data preparation for the publication and scientific presentations
  - Experiments to produce preliminary data for proposals and grant applications
  - Letters of support for proposals and grant applications



# What Could Be Done at FCRC, the Flow Cytometry Resource Center?

Operations	Extended Business Hours, by FCRC Staff				24/7/365, Self-Operated					
Lasers (Wavelength) and # of Fluorescent Detectors	Sorters			Image Analyzer	Spectral Analyzer	Advanced Analyzers			Basic Analyzers	
	BD FACS Aria-II-1	BD FACS Aria-II-2	BD FACS Aria-II-3	Image Stream-X	Cytek Aurora	BD LSRII-1	BD LSRII-2	BD LSR- Fortessa	ThermoFisher Attune NxT	BD Accuri C6
Blue (488nm)	3	3	3	5	14	3	3	3	3	4/3/2
Red (633-658 nm)	3	3	3	2	8	3	3	3		0/1/2
Violet (405-407 nm)	3	4	4	5	16	4	4	5		
Yellow/Green (561nm)	4	4	4	4	10	4	4	4	4	
UV (355 nm)		2	2			2		3		
Blue/Violet (445 nm)			2				2			
Total # of Detectors	13	16	18	10	48	16	16	18	7	4



# What Could Be Done at FCRC, the Flow Cytometry Resource Center?

## Graphical Population Definitions

Define populations using familiar graphical tools and combine them with logical functions.

## Comprehensive Population Statistics

Characterize your cell populations with a wide range of statistical metrics to reveal differences in cell morphology, phenotype, and function.

## Operational Parameters

Lasers (Wavelength and # of Fluorochromes Detected)

Blue (488)

Red (633)

Violet (405)

Yellow/Green (568)

UV (351)

Blue/Violet (405)

Total # of Cells Detected

## Inspect Your Populations

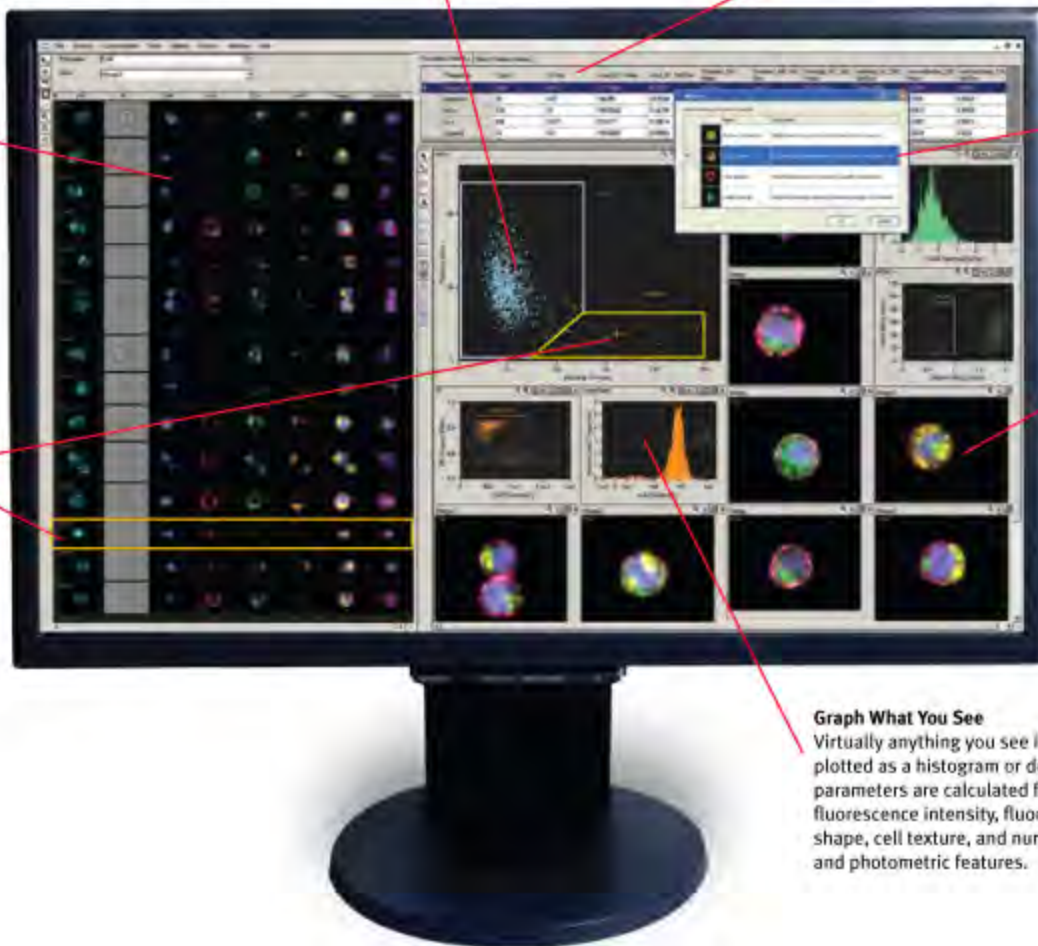
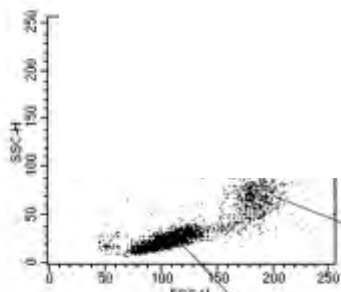
The Image Gallery allows you to see every image of every cell or perform a "virtual cell sort" to inspect and validate the cells within a specific population.

## Images for Every Dot

Every dot in every scatter plot is linked to the corresponding cell imagery. Simply click on a dot to see the associated cell images or vice-versa.

Bacteria: 0.5 µm  
Phytoplankton: 2 µm

Smaller



## Wizards Simplify Analysis

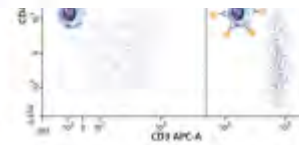
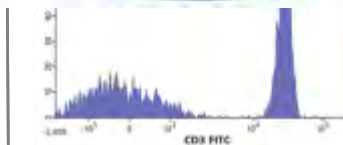
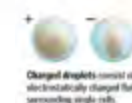
Pre-configured and optimized analysis wizards are provided for many common applications.

## Flexible Image Display Tools

Create composite images, pseudo-color representations and a host of other image transformations for reporting and publication.

## Graph What You See

Virtually anything you see in the imagery can be plotted as a histogram or dot plot. Hundreds of parameters are calculated for every cell, including fluorescence intensity, fluorescence location, cell shape, cell texture, and numerous other morphologic and photometric features.

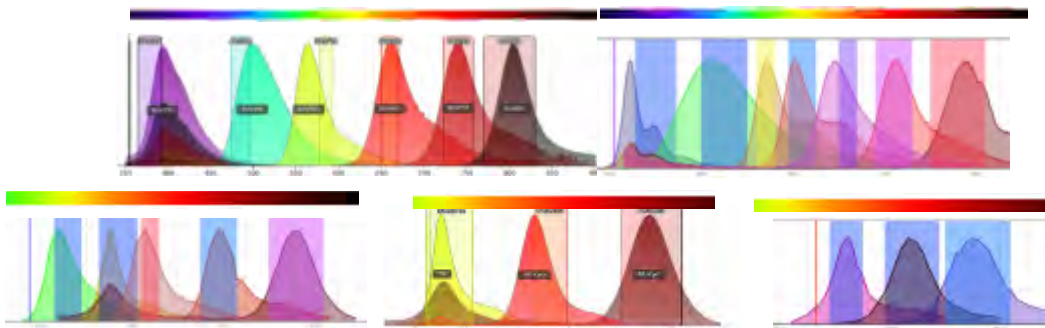




# Revolution in Flow Cytometry - Spectral Flow Cytometer, Cytek



## Cytek Aurora Spectral Analyzer

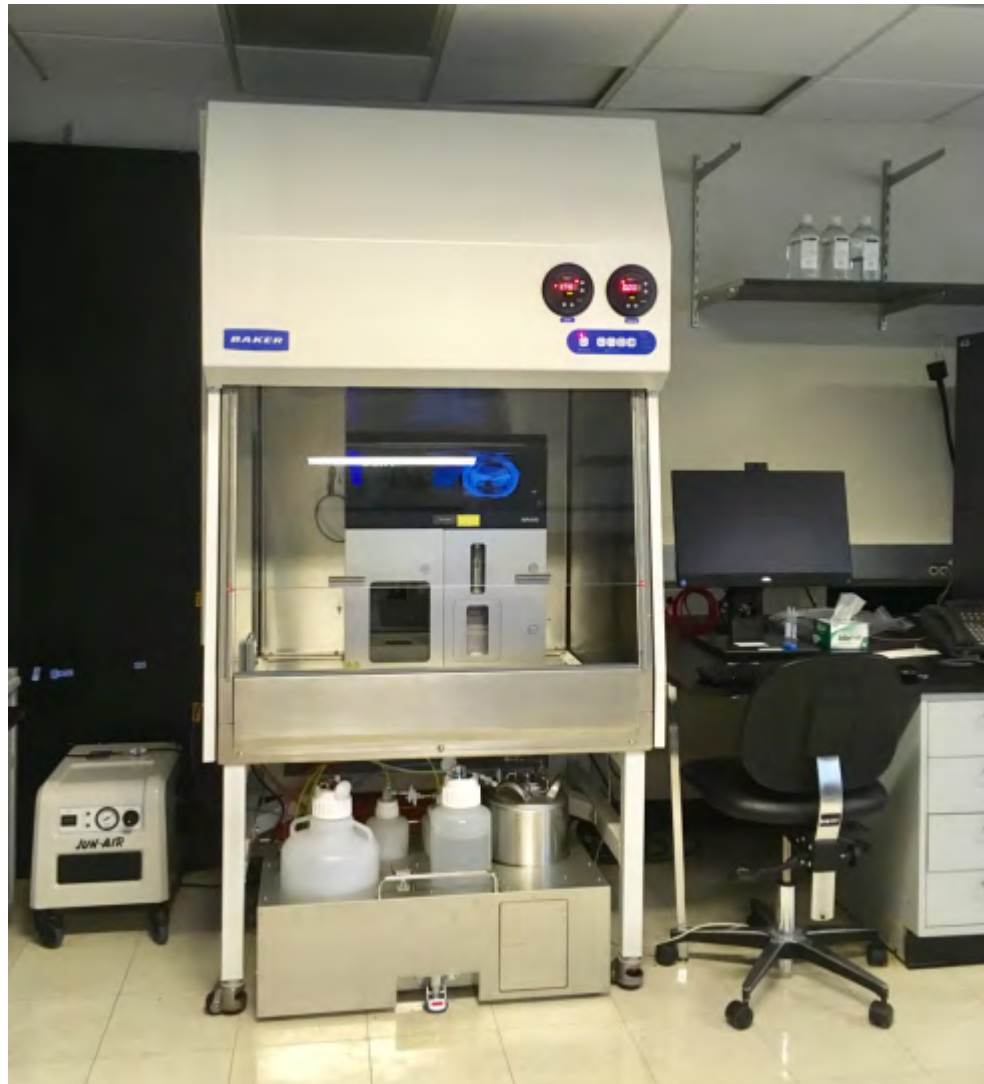


## Aurora™

1980	1 Laser, 2 Colors
1990	2 Lasers, 4 Colors
2000	3 Lasers, 8 Colors
2010	3 Lasers, 13 Colors
2017	3 Lasers, 20 Colors
2018	4 Lasers, 24 Colors
2019	5 Lasers, 32 Colors



# Sony MA900 Cell Sorter



**Self Sorting???**

# Services at the Genomics Resource Center

- **Next-generation sequencing**

- Whole genome and whole exome sequencing
- Targeted sequencing
- RNA-Seq: Transcriptome analysis
- Small RNA seq: small RNA discovery and quantitation
- ChIP-seq: Protein-DNA binding
- Methyl-Seq: epigenetic analysis of DNA methylation
- Ribo-Seq: Ribosome profiling for active translation
- TRAP-Seq: cell type specific mRNA profiling
- ATAC-Seq: chromosomal accessibility
- CROP-Seq: pooled CRISPR screening with single-cell transcriptome readout



- **10 x Genomics Chromium Single Cell System**

- Single cell gene expression
- Single cell full-length V(D)J profiling
- Single cell ATAC-Seq
- Single cell RNA Seq with CRISPR Screening



- **Realtime PCR systems**

- Individual gene expression analysis
- Pathway analysis
- Individual SNP genotyping

- **Others**

- Covaris Ultrasonicator
- Agilent Bioanalyzer
- Agilent TapeStation
- Qubit fluorometr
- NanoDrops
- PCR machines 96/384-well



# Sequencer Selection



NovaSeq  
800 million to 10 Billion reads

- Whole genome and Whole exome sequencing
- Paired-end RNA-Seq
- ATAC-Seq
- Large scale RNA-Seq
- Single cell RNA-Seq



NextSeq  
400 million reads

- Small scale RNA-Seq up to 16 samples
- ChIP-Seq
- Small RNA Seq



MiSeq  
1 – 20 million reads

- PCR Amplicon Sequencing
- 16S rRNA Metagenomic Sequencing as 300bp x 2
- Library prep method development
- Library QC

# Transcriptome Sequencing (RNA-Seq)

- **mRNA sequencing**  
Start with 100 ng total RNA  
Gene expression and splicing analysis of coding RNA
- **Total RNA sequencing (with rRNA depletion)**  
Start with 100 ng Total RNA  
Gene expression analysis of coding and non-coding RNA
- **Low-input mRNA sequencing**  
Start with as low as 100 pg – 1 ng total RNA
- **Ribo-profiling**  
Which mRNA are actively translated
- **TRAP-Seq**  
Cell type specific mRNA expression
- **Single Cell RNA sequencing**  
mRNA profiling at single cell level

# Genomics Resource Center

WRB 725 24/7 access

[www.rockefeller.edu/genomics](http://www.rockefeller.edu/genomics)

genomics@rockefeller.edu



Connie Zhao, Ph.D.  
Director



Hong Duan, PhD  
Research Support Specialist



Bin Zhang  
Research Support Specialist



Christine Lai  
Research Support Specialist



Xiaoyun Qiu  
Research Support Assistant



Sophie Huang  
Bioinformatics Specialist

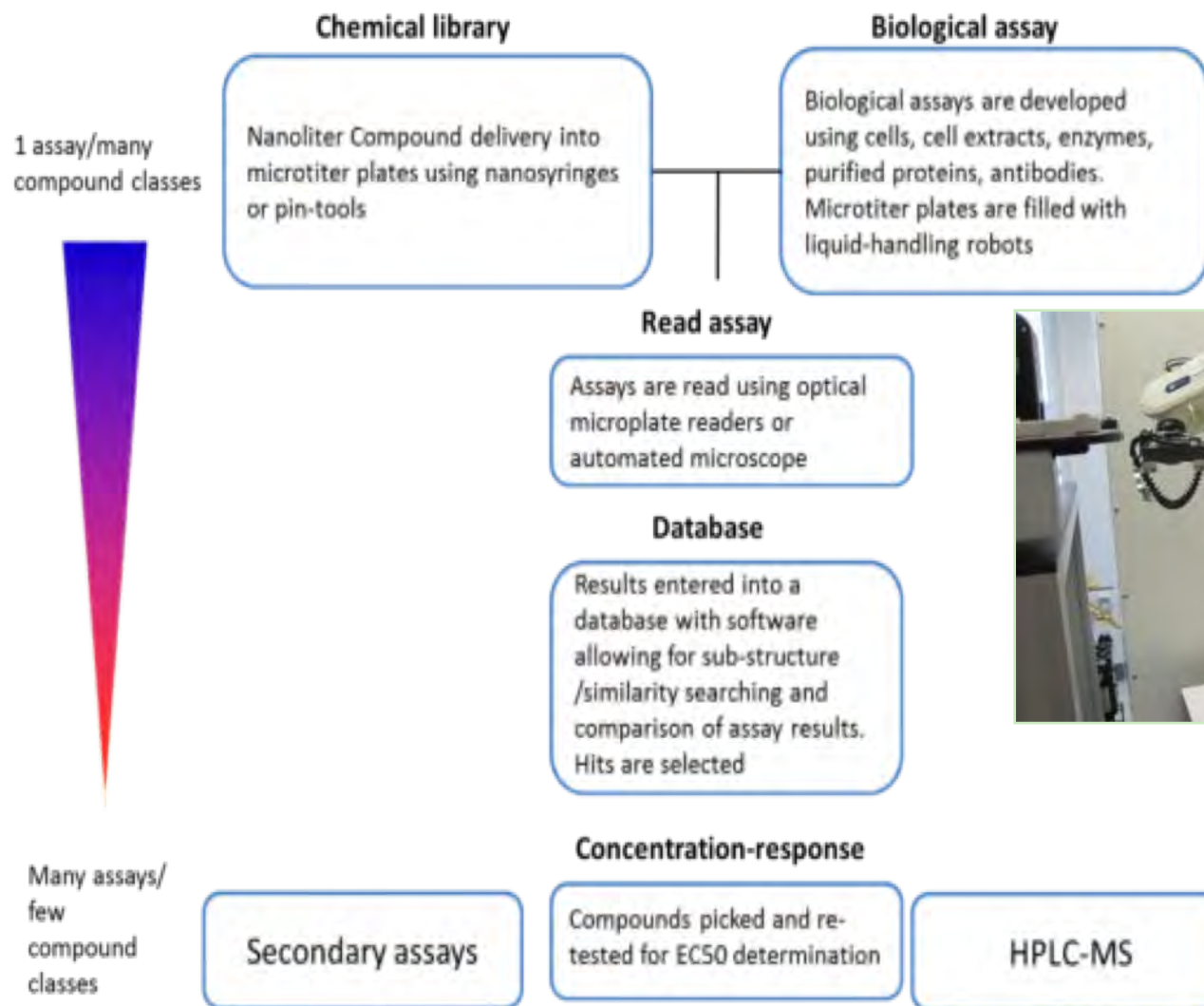
# High Throughput and Spectroscopy Resource Center, DWB 216

<http://inside.rockefeller.edu/htsrc>

**The HTSRC provides scientific and technical support and access to instrumentation in the following areas:**

- assay cascades used in drug discovery projects
- miniaturized assay development
- high throughput liquid handling
- high-throughput screening
- identifying tool compounds or lead compounds for drug discovery
- determination of structure activity relationships/medicinal chemistry efforts
- measuring the kinetics and thermodynamics of biomolecular interactions
- analysis of small molecular weight compound structure and purity

# How Is An HTS Project Accomplished?



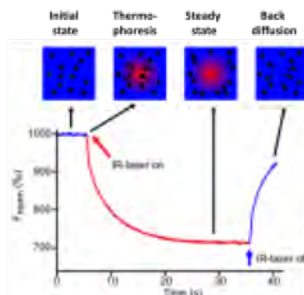


# Biophysical Techniques for Binding Measurements

## Microscale Thermophoresis: Nanotemper MST

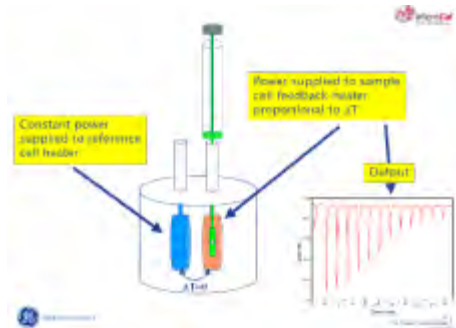
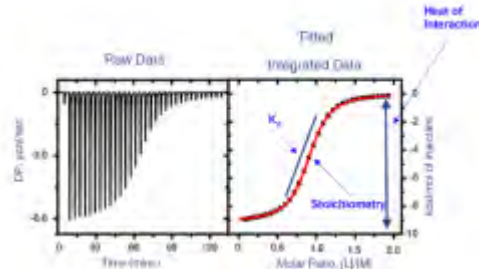
### Principle

Molecules migrate along a temperature gradient at different rates depending upon size, shape (hydration shell) and charge thus allowing differentiation between unbound and bound state



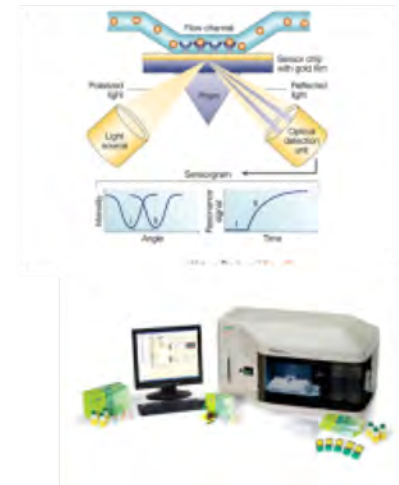
## Isothermal Calorimetry

- Label-free
- Autosampler
- 400uL/1 mg protein per experiment



## Surface Plasmon Resonance : Proteon XPR

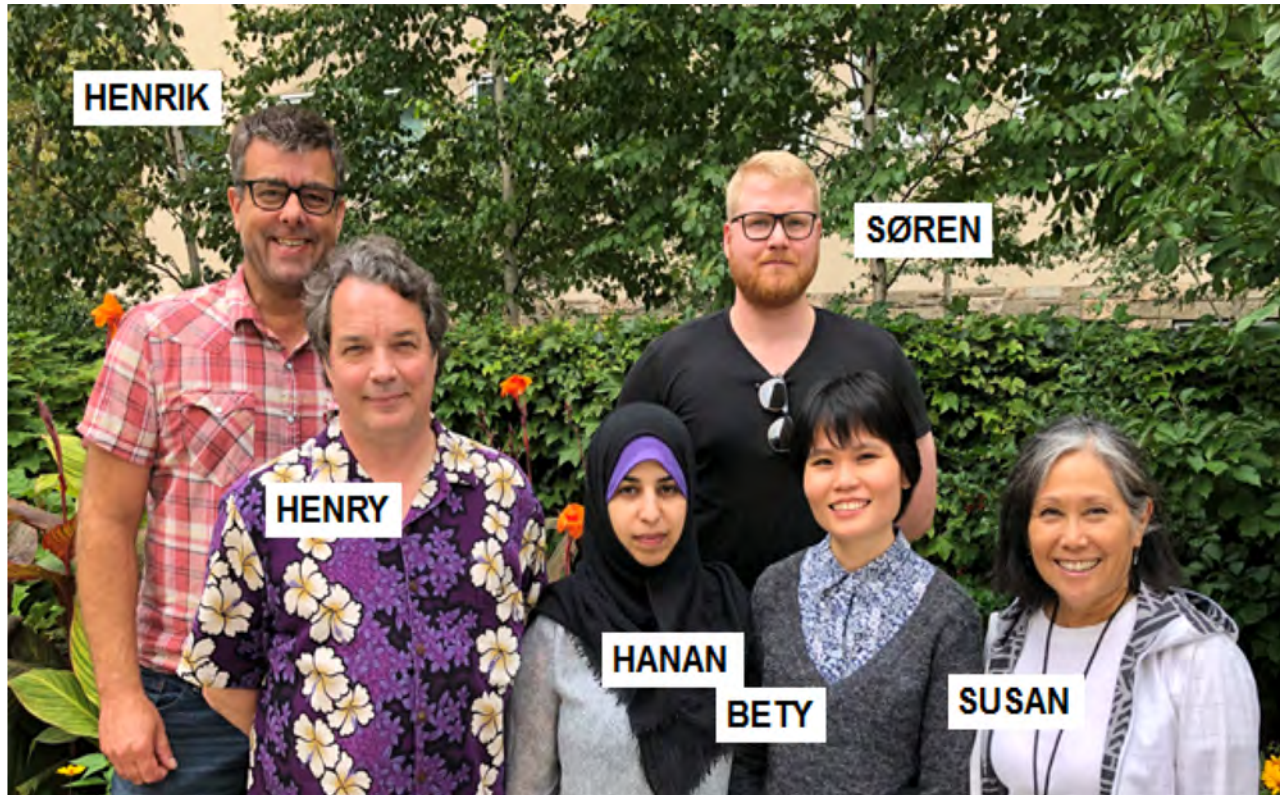
20-50uM Protein Immobilized to Gold Sensor Chip Measures on-rate and off-rate Immobilization through free-amines, antibody, biotin



### HTSRC Staff

- Fraser Glickman, Ph.D., *Director*
- Carolina Adura, Ph.D., *Manager of Spectroscopy*
- Jeanne Chiaravalli, M.S., *Research Support Specialist*
- Lavoisier Ramos-Espiritu, Ph.D., *Research Support Specialist*

# Proteomics Resource Center



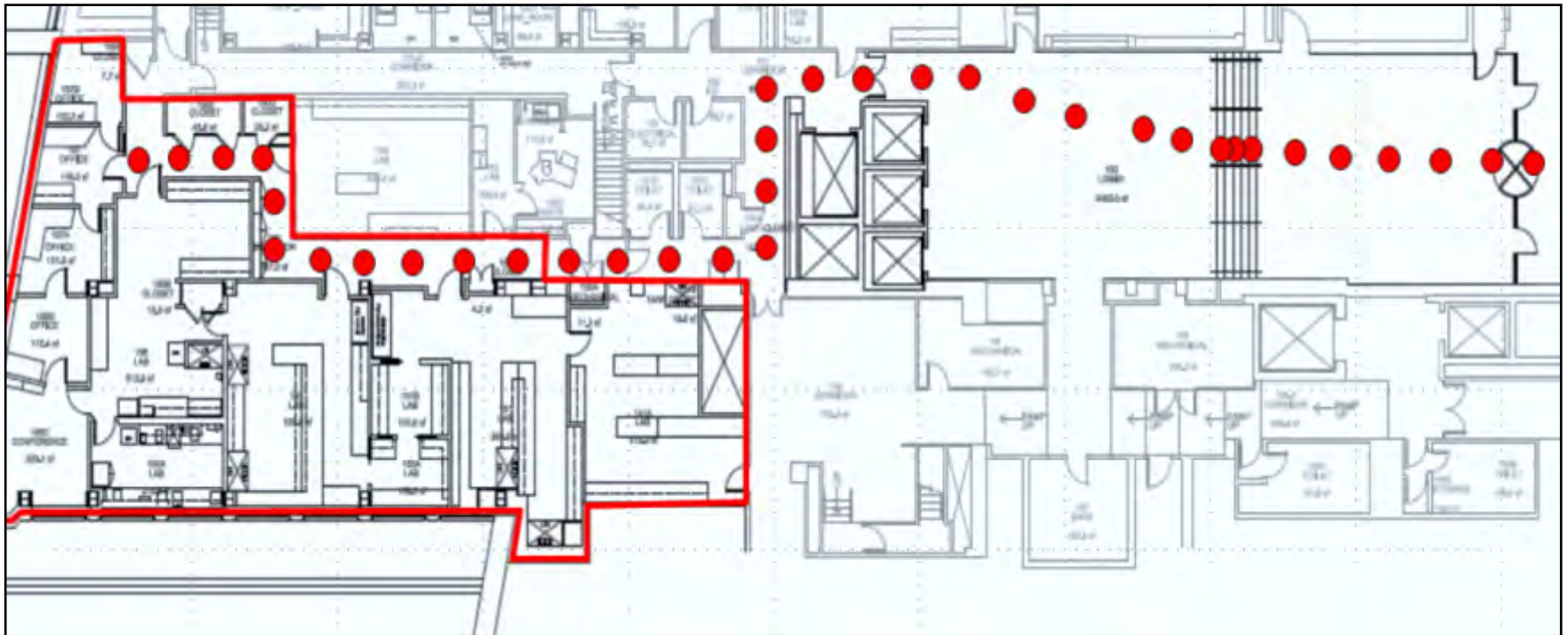
# Proteomics Resource Center

## What we do...

- Proteomics – identification and quantitation of proteins, peptides and small molecules by mass spectrometry
- Peptide synthesis
- Analytical chemistry
- User instrumentation
- Help planning experiments...

# Proteomics Resource Center

Where to find us...



Rockefeller Research Bldg. Room 157

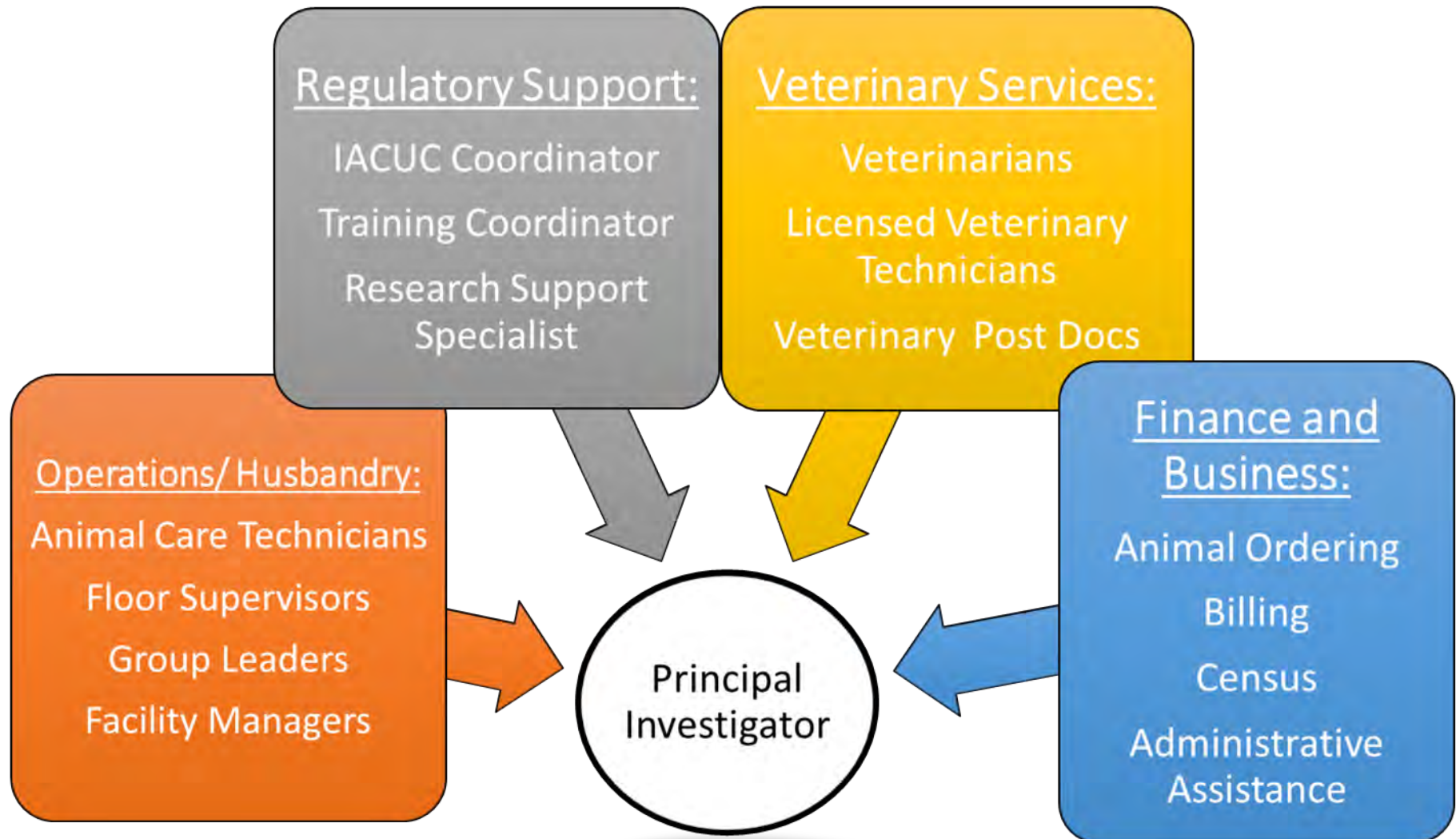
<http://inside.rockefeller.edu/proteomics/>



# Comparative Bioscience Center (CBC)



# Structure of the CBC





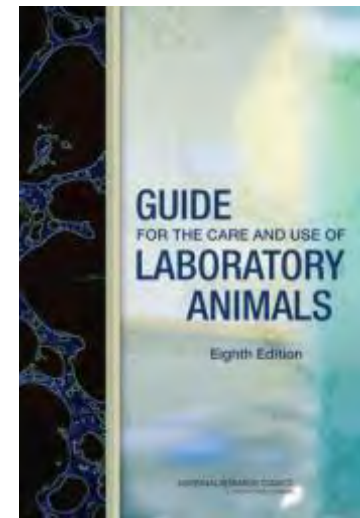
# The CBC Facilitates Quality Biomedical Research

## IACUC

- Institutional Animal Care and Use Committee
  - Ethics review
  - Scientific review

## High Standards of Animal Care

- Optimal environment
- Optimal health
  - Maintained disease free
  - Extensive health surveillance programs
- Minimize research variables



# Educational/ Research Support

1. CBC Orientation
  - Lecture & Lab (Restraint, Anesthesia, Dosing)
2. Aseptic Techniques for Surgery
3. Micro-Ultrasound/IVIS Imaging
4. Safe use of Hazards
5. Dosing: SC, IM, IV, PO
6. Blood Collections
7. Tissue Collections
8. Tri-Institutional Seminars
  - Rodent Breeding, Phenotypic Characterization of Mice



# Vevo 2100 Ultrasound

- High anatomical resolution
  - Color Doppler for blood flow identification
- Research areas applicable:
  - Cardiovascular studies
  - Urology
  - Contrast imaging
  - Cancer
  - Developmental biology
- Useful with mice, rats, fish



# IVIS Spectrum

- High Sensitivity in vivo fluorescence and bioluminescence imaging
- Monitor disease progression
- Cell trafficking
- Gene expression
- Measure up to 5 mice at a time

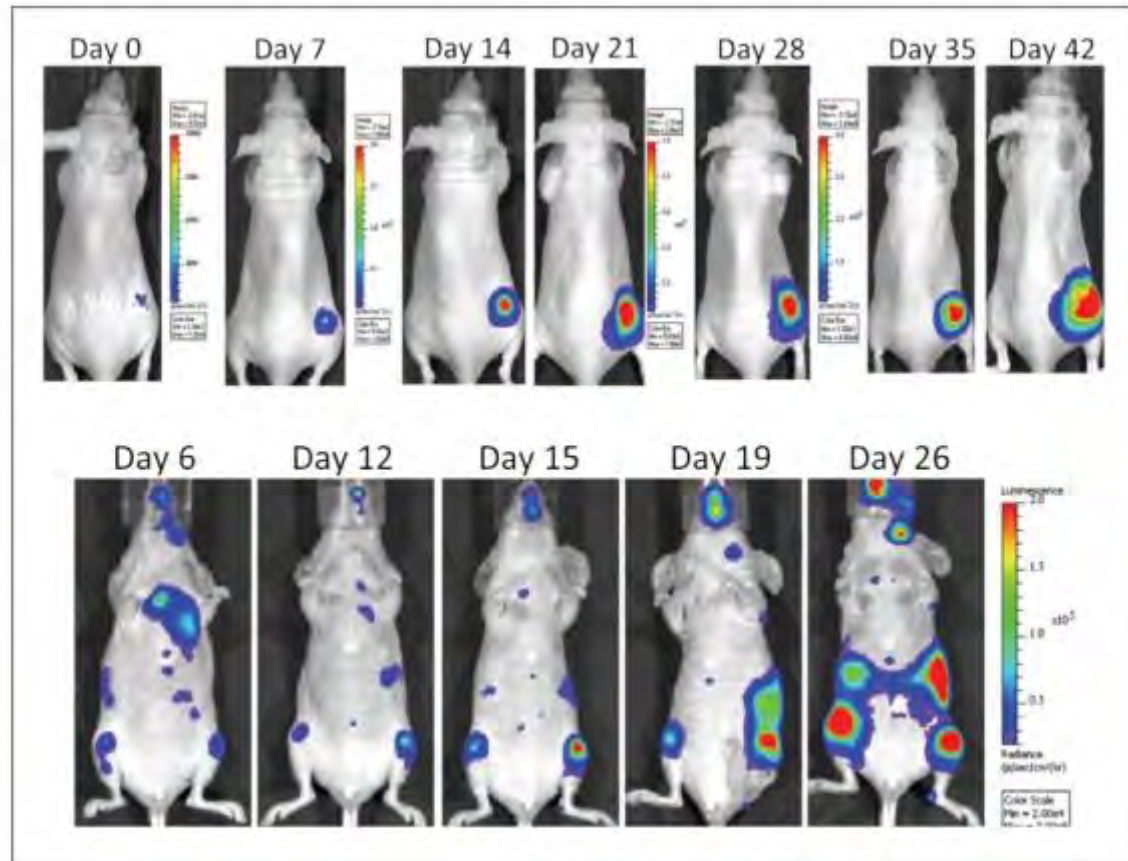
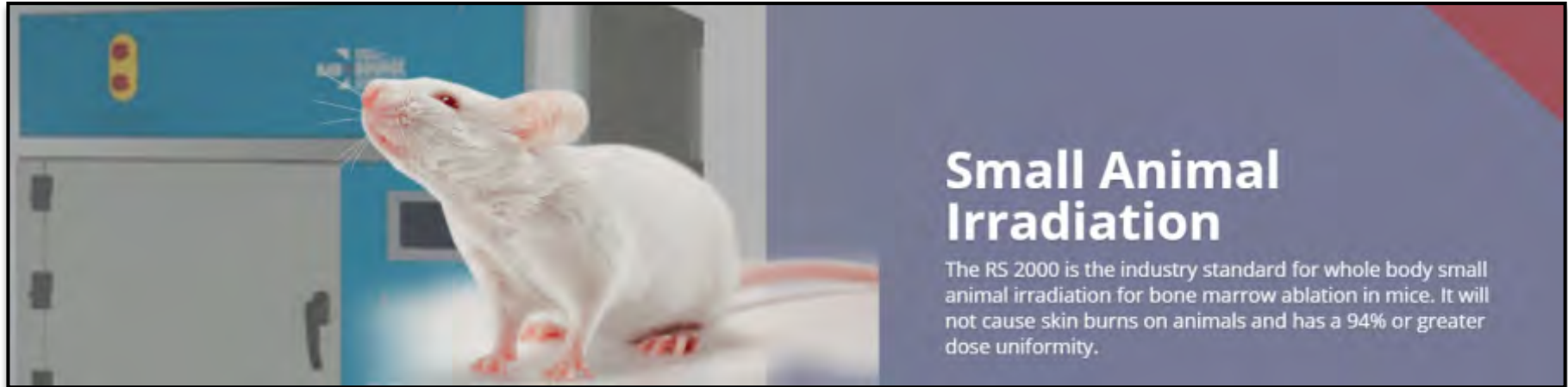


Figure 4. Detection of five 4T1-luc2 cells injected subcutaneously in nude mice (top) and monitoring metastasis post intracardiac injection of MDA MA-231-luc2 cells (bottom) longitudinally.

# Rad Source 2000 X-Ray Biologic Irradiator

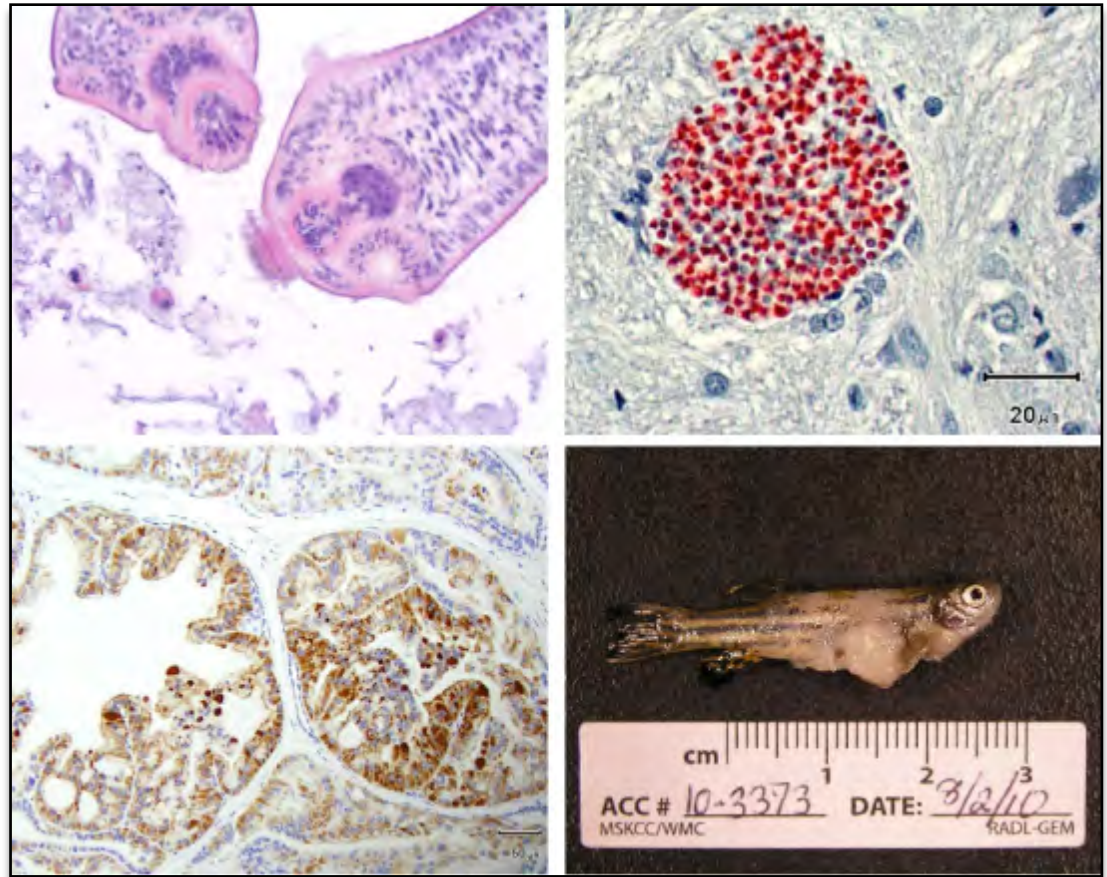


- Generate bone marrow ablation in mice
- Useful for
  - Cancer studies
  - Targeted Tumor irradiation
  - Host - Viral Infection



# Laboratory of Comparative Pathology

- Genetically Modified Animal Phenotyping
- Hematology and Clinical chemistry
- Complete Necropsy
- Histology
- Radiology
- Bone marrow evaluation
- Organ-specific research studies: mammary gland, prostate, heart





*Leslie Diaz, Associate Director, CBC*

[ldiaz@rockefeller.edu](mailto:ldiaz@rockefeller.edu)





# CRISPR & Genome Editing Center

Chingwen Yang, Ph.D.  
Director  
DWB 703

Tel: 212-327-8649  
yangc@rockefeller.edu

## Gene editing Service

mouse embryos

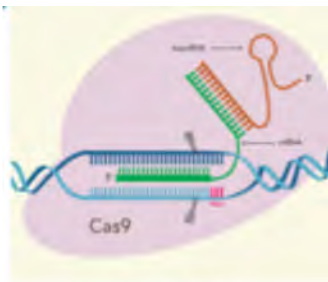
adult mice

mES cells

mouse lymphoma cell lines

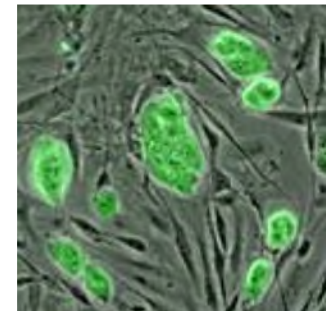
human iPSC

primary human cancer cells



**RNP**

**ssDNA**



# Genome Editing in mouse zygote

## **KO: ctRNP**

frameshift mutation

genomic deletion & Inversion

*high efficiency, easy to achieve homozygous mutants*

*microinjection, embryo electroporation, iGONAD*

## **KI of short modification: ctRNP + ss oligo (<200 bases)**

missense mutation

insertion of loxP site, small tag

*modification up to 60nt*

*high efficiency, sometimes homozygous*

*microinjection, embryo electroporation, iGONAD*

## **EasiCRISPR: ctRNP + ssDNA (500 -2500 bases)**

KI of reporter

conditional KO allele

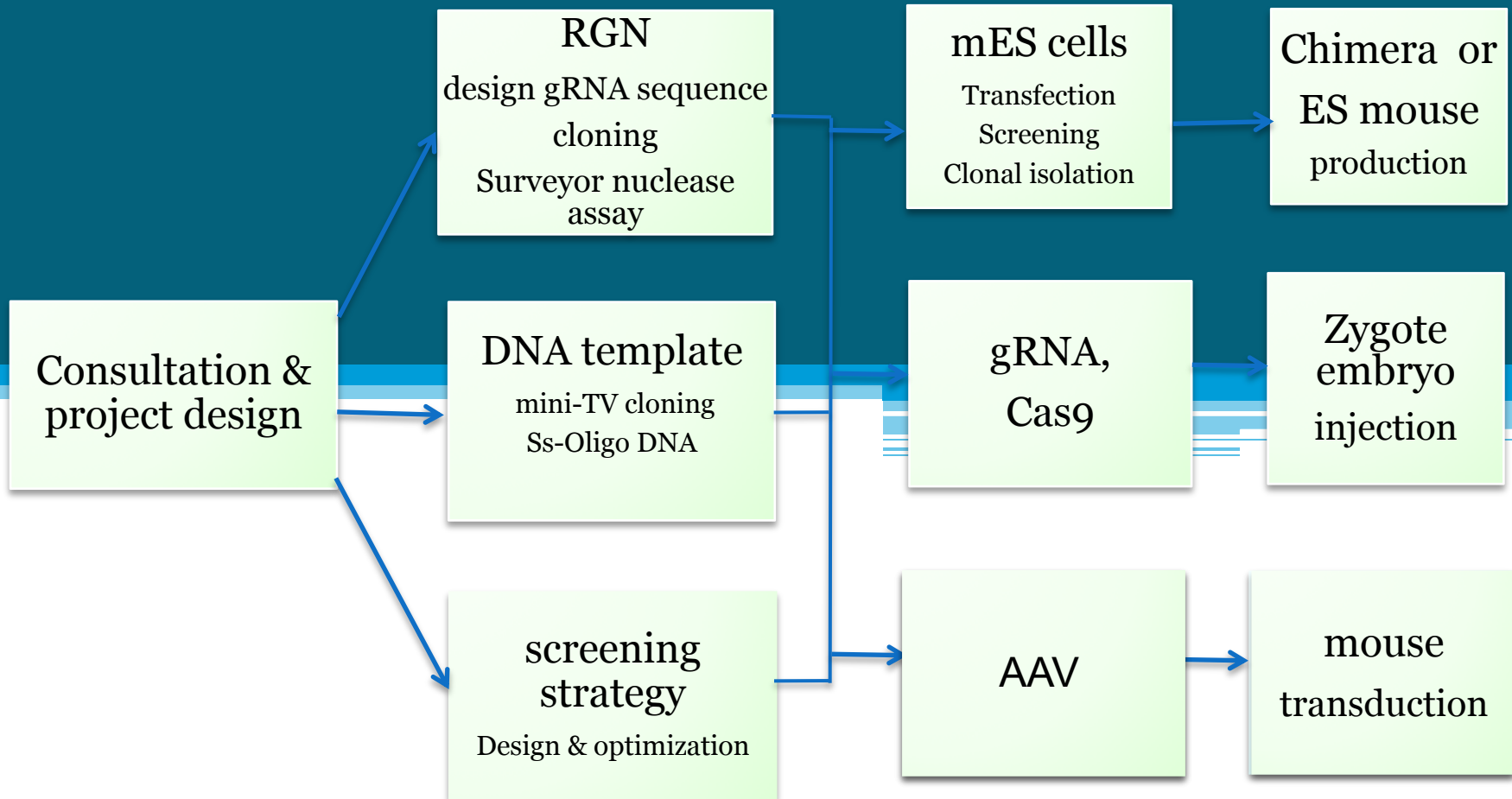
exon replacement

*modification up to 1500bp*

*medium efficient, less likely for complete homozygous alleles*

*microinjection, ~~embryo electroporation~~, iGONAD*

# CRISPR & Genomic Editing Center





# ***CRISPR & Genome Editing Center***

*DWB 702-703*

*327-7307, -8650, -8649*



**Chingwen Yang**, Ph.D., Director

**Jing Gao**, M.D., Manager in cell biology

**Dorjee Shola**, Ph. D., Manager in molecular biology

**Chia-Yun Han**, M.S, Research support specialist

**Pradip Kar**, M.S., Research support specialist

**Qilie Luo**, Ph. D. Research support specialist

**Vhy-Shelta Kewalder**, B.S., Research support specialist



## Transgenic and Reproductive Technology Center

CBC 542-546  
x7783, x7738

Director:

Rada Norinsky

[Rada.Norinsky@rockefeller.edu](mailto:Rada.Norinsky@rockefeller.edu)

Research Support  
Specialists:

Jahnney Torres

Eunyong Kim

William Ramirez

Roxana Cubias

# Mutant Mouse Models Production Services

## *Targeted Mutations (KI, KO, conditional KO)*

- Microinjection of CRISPR/Cas9 complex directly into zygotes

## *Transgenic Mice*

- Microinjection of plasmid DNA and BAC DNA into zygotes

## *Chimeric Mice*

- Microinjection of ES cells into blastocysts





# Assisted Reproduction/Rederivation Services

- Accelerated colony expansion  
(*IVF with fresh and frozen sperm*)
- Accelerated embryos cryopreservation
- Assisted reproduction for difficult lines
- Resuscitation of cryopreserved mouse embryos
- IVF recovery of lines from frozen sperm
- Rederivation from frozen embryos/sperm for generation pathogen free mouse lines



# GLASSWASHING SERVICES



**CRC Room CO2E (Greenberg Building)  
7AM-3PM**

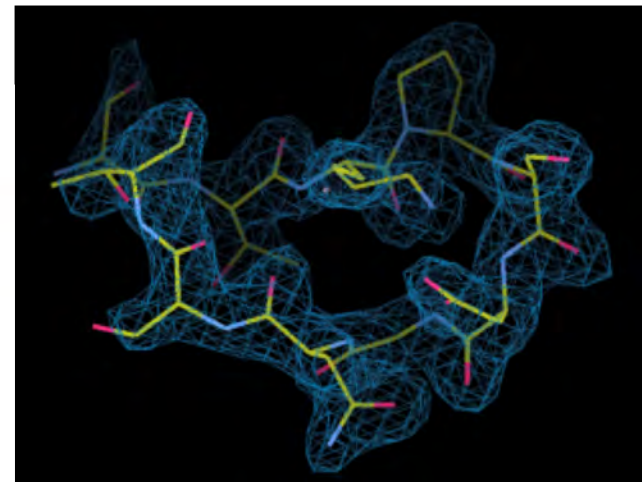
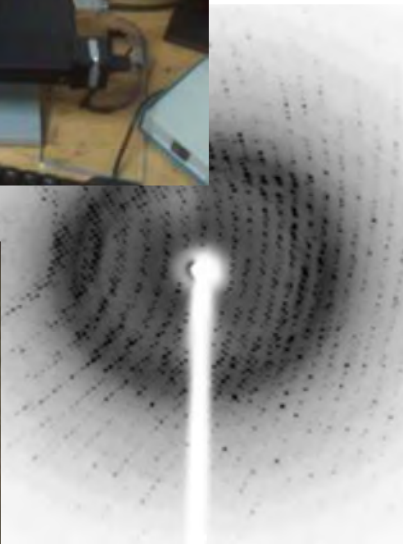
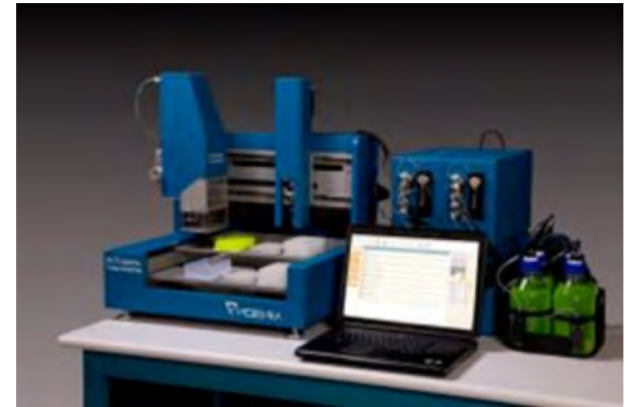
<http://www.rockefeller.edu/glasswashing/>



# Structural Biology Resource Center

Do X-ray  
crystallography  
with us

From crystal growth to  
structure determination



# Structural Biology Resource Center

Or.... make and purify proteins with us



## NEW SEC-MALS

What is SEC-MALS?

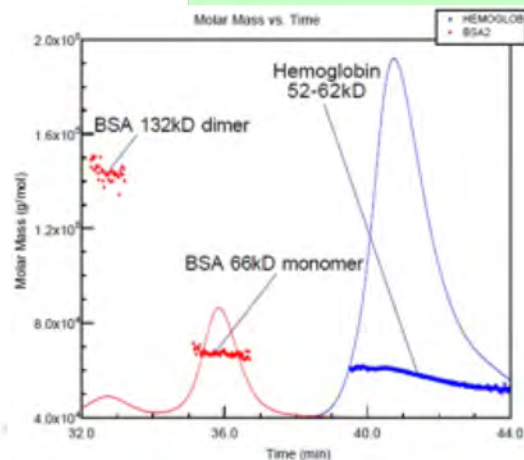
Size Exclusion Chromatography coupled to Multi-Angle Light Scattering



## DO YOU WANT PURIFIED PROTEIN?

The Structural Biology Resource Center  
has a new protein expression and  
purification training center.

- We will provide the tools and knowledge for satisfactory results.
- You will develop the expertise to continue working independently.



Deena Oren, Ph.D.  
Manager  
RRB Suite 140  
[inside.rockefeller.edu/sbrc](http://inside.rockefeller.edu/sbrc)

# Precision Instrumentation & Technology Makerspace

## How Can You Improve Your Science?

- Training to use various design and fabrication tools
- Stock materials (plastic and metal)
- Workspace, CAD computers
- Equipment
  - Sketch, Surface, Solid Body Modeling
  - 3D Printers (FDM, DLP, MJP)
  - Laser Cutter (CO<sub>2</sub> Emission)
  - CNC Mills (5-Axis, 4-Axis)
  - Electronics and Microcontrollers
  - Lathe
  - Micro Welder
  - Knee Mill



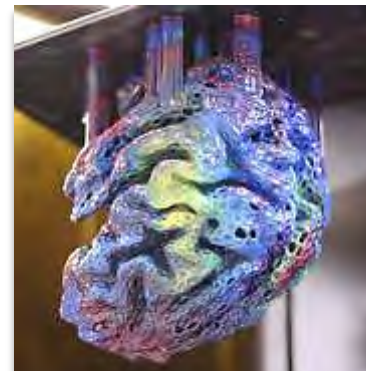




# Services

## How Can We Improve Your Science?

- Free Expert Consultations
- Custom Scientific Equipment Design & Fabrication
  - Precision Machining
  - Prototyping
  - Parametric Modelling
  - Metrology
  - Integrated Systems
- Procurement & Referrals
- Process Improvement
- System Analysis, Diagnosis & Repair
- Training







# The PIT



**Dan Gross**  
Scientific Engineer

**Jim Petrillo**  
Instrumentation Engineer

**Peer Strogies**  
Scientific Machinist

Location: Plaza Building, A level  
[inside.rockefeller.edu/fabrication/](https://inside.rockefeller.edu/fabrication/)

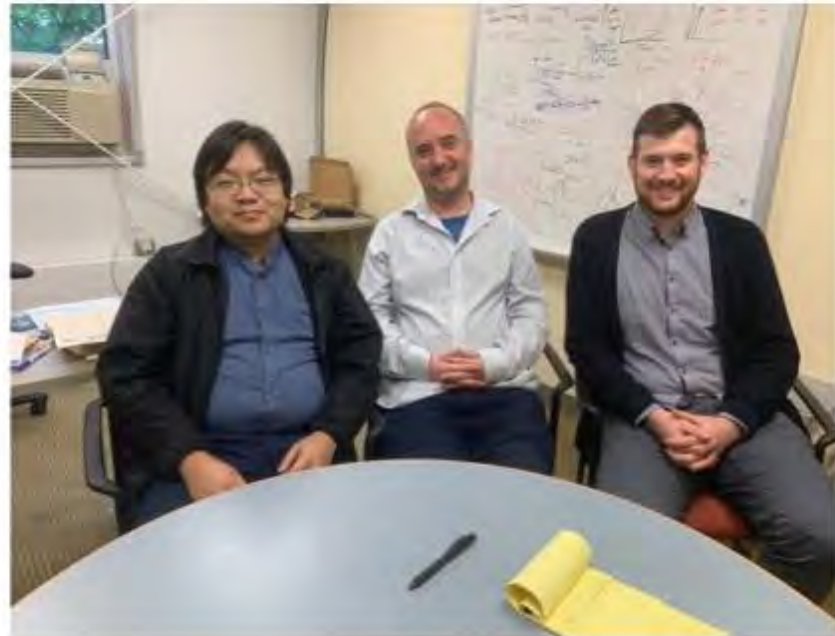
# Bioinformatics Resource Center

- Provide bioinformatics software infrastructure.
- Conduct training courses and community events.
- Work with wet and dry biologists on both a consultative and collaborative basis.
- Experimental design.



# Bioinformatics Resource Center

- Who we are.
  - Tom Carroll
  - Ji-Dung Luo
  - Matt Paul
- What we do.
  - Training
  - Infrastructure
  - Analysis on demand.





# BRC- Training

- Analysis of high-throughput-data in R and Bioconductor.
- New York R and Bioconductor meetups
- Bioconductor Conference 2019 at RU -> 2020 organizing committee.
- 1 to 1 training through Github.

## Upcoming training

In 2018 we will be running training on R, reproducible research, visualising data in genome browsers and analysis of RNA-seq/ChIP-seq using R and Bioconductor

To book a place on one of these courses please contact Thomas Carroll ([thomas.carroll@rockefeller.edu](mailto:thomas.carroll@rockefeller.edu))

Material linked below is for reference and example only.

Material to be used for course will be made available to attendees only and weblinks distributed on the day of the course.

**NOTE** All materials are under strict licensing. Material is free for review but may not be redistributed 'as is' or modified without explicit consent. For more details contact the Head of Bioinformatics ([Thomas.Carroll@rockefeller.edu](mailto:Thomas.Carroll@rockefeller.edu))

Show  entries

Search:

Course	Material	Dates	Availability
Introduction to Bioinformatics Resource Centre	<a href="#">Link</a>	TBD	TBD
ATACseq Workshop	<a href="#">Link</a>	TBD	TBD
Introduction to Spring 2018 Bioinformatics Course	<a href="#">Link</a>	TBD	TBD
Intro To R	<a href="#">Link</a>	TBD	TBD
Plotting In R	<a href="#">Link</a>	TBD	TBD
Genomic Data	<a href="#">Link</a>	TBD	TBD
IGV	<a href="#">Link</a>	TBD	TBD
Introduction to Bioconductor	<a href="#">Link</a>	TBD	TBD
ChIPseq in Bioconductor	<a href="#">Link</a>	TBD	TBD
ATACseq in Bioconductor	<a href="#">Link</a>	TBD	TBD
RNAseq in Bioconductor	<a href="#">Link</a>	TBD	TBD
Visualising genomics data in R and Bioconductor	<a href="#">Link</a>	TBD	TBD

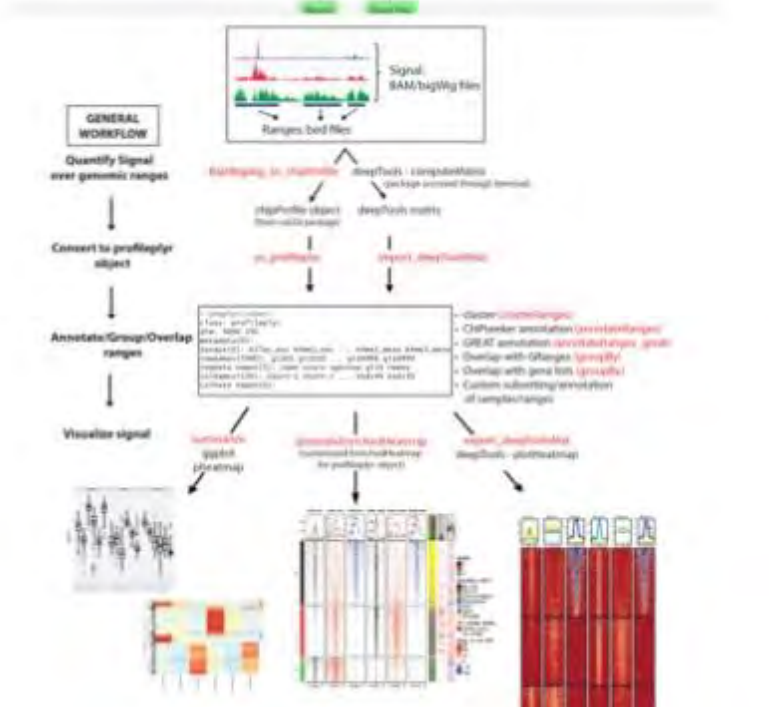
Showing 1 to 12 of 12 entries

Previous  Next

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- 
- ```

graph TD
    SampleCollection[Sample Collection] --> SamplePreparation[Sample Preparation]
    SamplePreparation --> LibraryConstruction[Library Construction]
    LibraryConstruction --> Sequencing[Sequencing]
    Sequencing --> DataGeneration[Data Generation]
    DataGeneration --> DataAnalysis[Data Analysis]
    DataAnalysis --> Interpretation[Interpretation]
    Interpretation --> Reporting[Reporting]
  
```



# Analysis on demand

## Analysis as requested

- ~30 groups.
- Custom analysis fitted to project.
- Regularly assess requests and adapt pipelines and/or training dependent on demands for analysis types.



The screenshot shows a web-based interface for managing analysis requests. It features a table with columns for various attributes of the requests, organized into sections.

| Section                  | ID  | Name         | Status      | Priority | Owner    | Created          | Updated          |
|--------------------------|-----|--------------|-------------|----------|----------|------------------|------------------|
| Analysis 1 (Completed)   | 101 | Genetic Data | Completed   | High     | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 102 | Genetic Data | Completed   | Medium   | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 103 | Genetic Data | Completed   | Low      | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 104 | Genetic Data | Completed   | High     | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 105 | Genetic Data | Completed   | Medium   | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 106 | Genetic Data | Completed   | Low      | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
| Analysis 2 (In Progress) | 201 | Genetic Data | In Progress | High     | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 202 | Genetic Data | In Progress | Medium   | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 203 | Genetic Data | In Progress | Low      | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 204 | Genetic Data | In Progress | High     | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 205 | Genetic Data | In Progress | Medium   | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 206 | Genetic Data | In Progress | Low      | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
| Analysis 3 (Pending)     | 301 | Genetic Data | Pending     | High     | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 302 | Genetic Data | Pending     | Medium   | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 303 | Genetic Data | Pending     | Low      | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 304 | Genetic Data | Pending     | High     | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 305 | Genetic Data | Pending     | Medium   | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |
|                          | 306 | Genetic Data | Pending     | Low      | John Doe | 2023-10-26 10:00 | 2023-10-26 10:00 |





Jason Banfelder  
Director, HPC Systems and Applications



Rebecca Bennett  
Senior Systems Engineer

# High Performance Computing



Bala Jayaraman  
Senior HPC Systems Administrator



Logan Sweezy  
Scientific Systems Administrator



# High Performance Computing



runs Linux

- On campus support for **data intensive science**
- Processing Power (257 double-precision TFlops)
  - More cores to devote to a problem (4,808 cores under management)
  - Specialized architectures: 54 GPUs; large memory (3 TB RAM) nodes
  - Batch vs. interactive
- Data Storage (2.8 PB)
  - Hundreds of disks for more IOPS
- Networking (56 Gbps/100 Gpbs)
  - Low latency InfiniBand



# Other Data Storage Options

- Box.com
  - No cost to users (IT provided)
  - File size limit
  - Extramural bandwidth limits
- Data Park
  - “cheap, deep, and simple”
    - 1/3 the cost of other options
  - Available for
    - SMB (Mac/Windows on campus)
    - sftp (Linux and cyberduck)
- Other Cloud (e.g. Amazon glacier)
- Protected or regulated data (e.g. HIPAA)?



$$a + \frac{1}{4} \cdot \sum_{n=1}^{\infty} a \cdot r^n = 2 \cdot a$$

when  $r = 0.8$

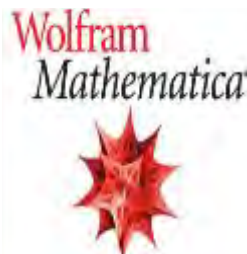


# HPC Software and Applications

- Image Processing (e.g. CryoEM)
- Genomics
- Neural Simulations
- Molecular Dynamics
- Artificial Intelligence (Deep Neural Networks)
- $\vdots$
- Campus licenses for Mathematica and Schrödinger



<http://hpc.rockefeller.edu>



# The Vertebrate Genome Laboratory (VGL)

Olivier Fedrigo, Director



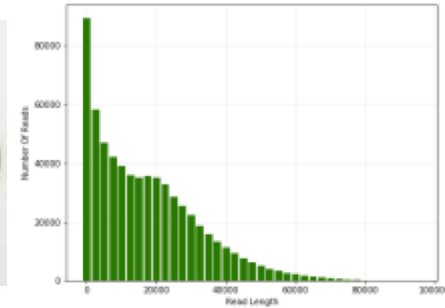
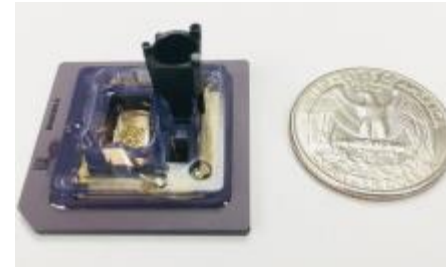
Weiss 7<sup>th</sup> floor

Email: [VGL@rockefeller.edu](mailto:VGL@rockefeller.edu)



## PacBio Sequels

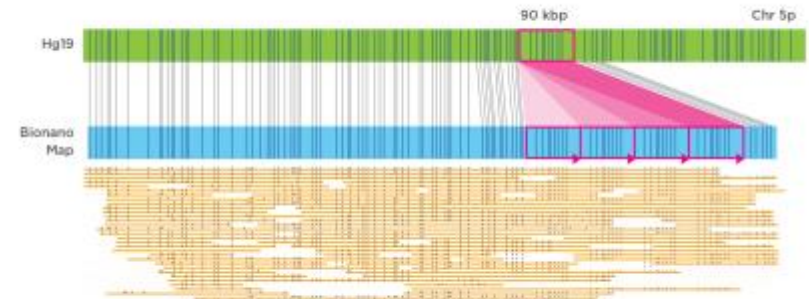
- Single molecule sequencing
- 3 Sequel and 2 Sequel II instruments
- 15-25 kilobases sequences (average)
- Up to ~120 gigabases per run



## Bionano Saphyr

- Optical mapping
- >150kb fragments (N50 >220kb)
- 24+ hours run; >300Gb per run

TANDEM REPEAT



- Whole genome sequencing
- Targeted sequencing (e.g. HLA)
- Complex populations (e.g. microbial communities)
- RNA sequencing (full length transcriptomics)
- Epigenetic
- Structural variants
- Genome scaffolding

# The Vertebrate Genome Laboratory



(212) 327-8216



VGL@rockefeller.edu

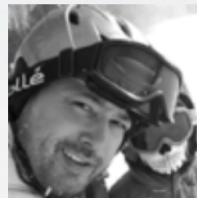


<https://vertebrategenomelaboratory.youcanbook.me>

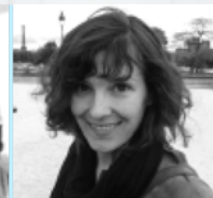


<http://inside.rockefeller.edu/vgl/>

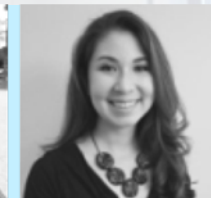
## Our team:



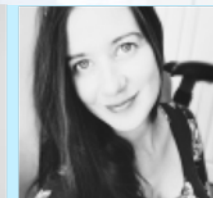
Olivier Fedrigo



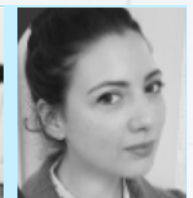
Bettina Haase



Jackie Mountcastle



Melanie Couture



Jennifer Balacco

Weiss 7<sup>th</sup> floor  
Room 735

If you are interested in long reads, come visit us at the VGL to discuss your ideas and projects

# Resource Center Locations

- **Antibody and Bioresource Core Facility** (DWB 415)
- **Bio-Imaging** (DWB 201-203)
- **Bioinformatics Resource Center** (IT Pavillion, 1st floor)
- **Comparative Bioscience Center** (CBC)
- **CRISPR & Genome Editing Center** (DWB 703)
- **Cryo-Electron Microscopy** (CRC B13)
- **Electron Microscopy** (RRB 120)
- **Flow Cytometry** (DWB 205-211)
- **Genomics** (WRB 725)
- **Glasswashing** (CRC-C level)
- **High Throughput and Spectroscopy** (DWB 219)
- **High Performance Computing** (IT Pavillion, 1<sup>st</sup> floor)
- **Instrument Design and Fabrication** (Plaza Building–A level)
- **Precision Fabrication Facility** (Plaza Building–A level)
- **Proteomics** (RRB 157)
- **Structural Biology** (RRB 1<sup>st</sup> floor)
- **Transgenics (CBC 542-546)**
- **Vertebrate Genome Laboratory** (WRB, 7<sup>th</sup> Floor)