#### Let Our State-of-the-Art Rangos Cores Ratchet Up Your Research

**August 11, 2020** 

12:00 PM - 1:00 PM

**Animal Imaging Core** 



Dr. Yijen Wu

**Histology Core** 



Dr. Sunder Sims-Lucas

**Cell Imaging Core** 



Dr. Krishna Prasadan

**Flow Cytometry Core** 



Dr. Abbe de Vallejo

**Metabolic Core** 



Dr. Clinton Van't Land

**Genetic Sequencing Core** 



Dr. Amanda Poholek

**Gnotobiotic Core** 



Dr. Tim Hand

**Bioinformatics Core** 



Dr. Dhivyaa Rajasundaram

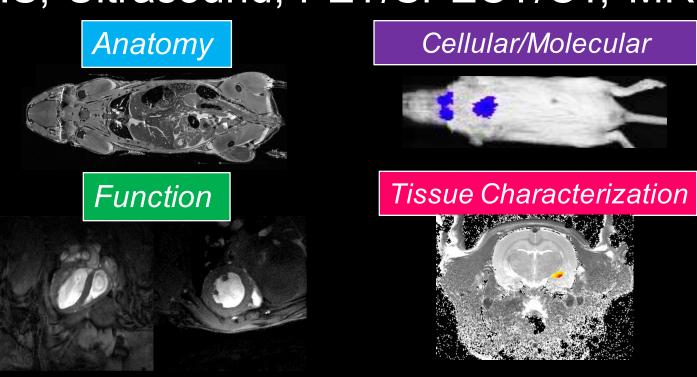
**AAV Production** 



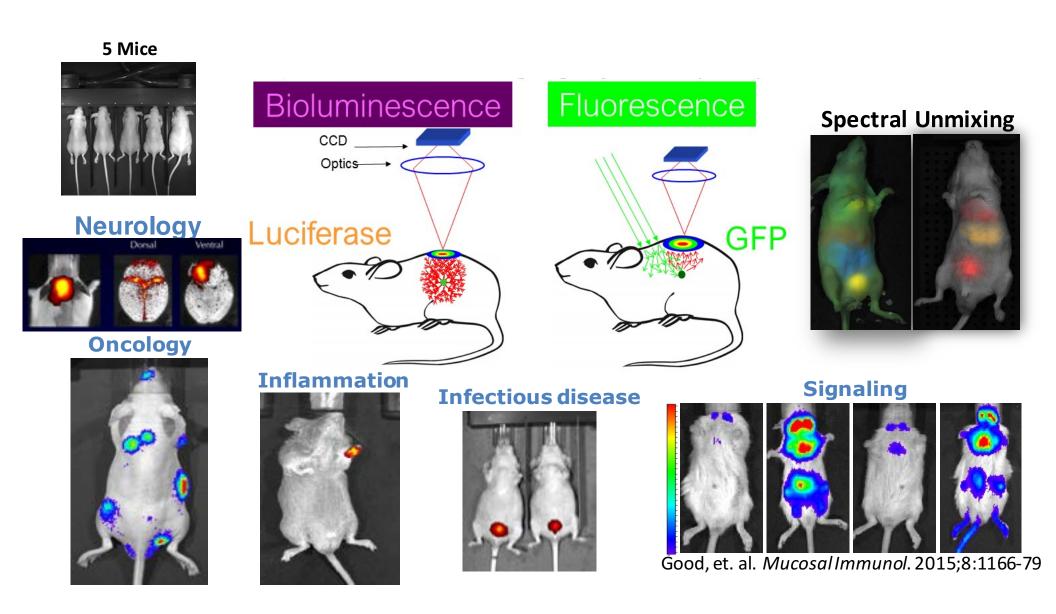
**Dr. George Gittes** 

# **Animal Imaging Core**

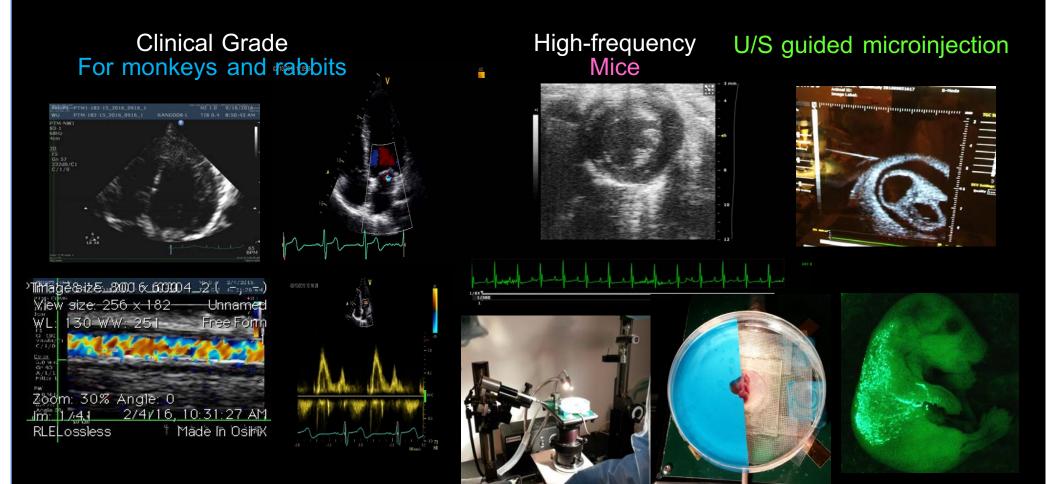
Systems, Tissue
in vivo, ex vivo
IVIS, Ultrasound, PET/SPECT/CT, MRI



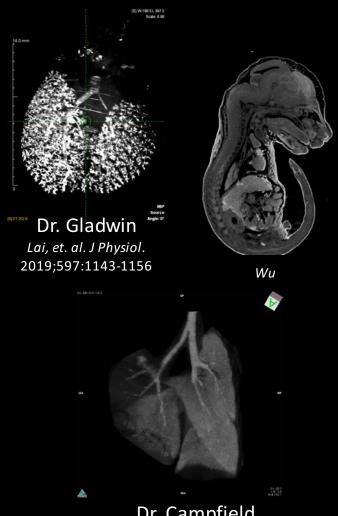
## Optical In Vivo Imaging System (IVIS)



## Ultrasound & Ultrasound-guided Microinjection



# PET/SPECT/CT

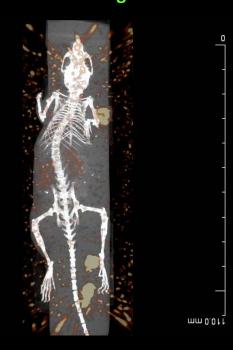


Dr. Campfield

Henkel, et. al. Am J Respir Crit Care Med. 2020;201:934-945.

## Molecular Imaging

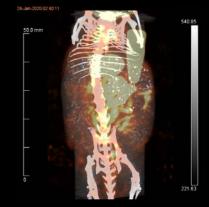
## Theranostics SPECT/CT Lu-177-PSMA-targeted CTT1403



Dr. Anderson Ling, et. al. *Mol Imaging Biol*. 2020;22:274-284

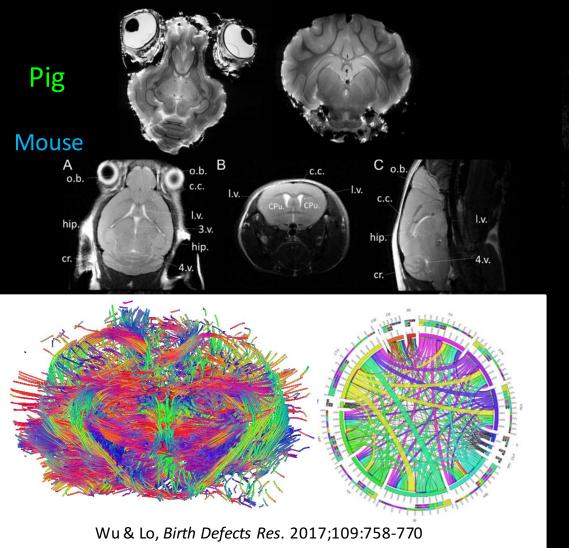


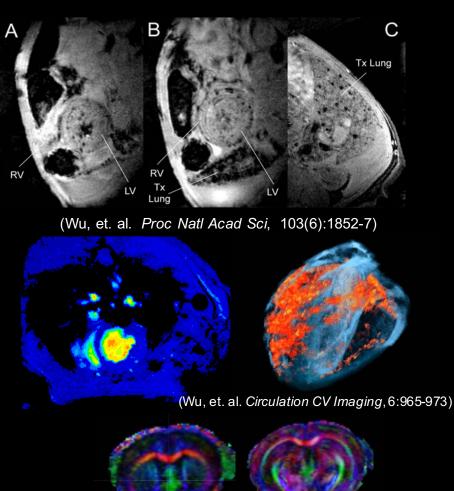
SPECT/CT mTc99-Sestamibi

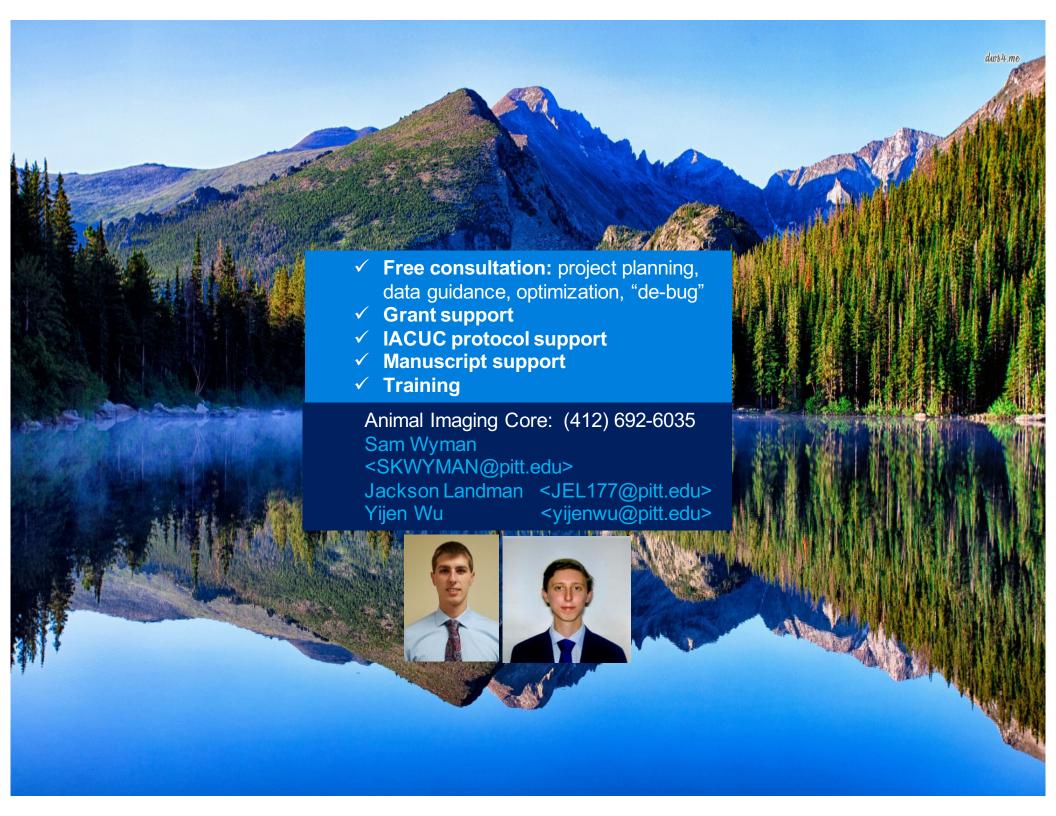


Wu

# MRI







# Histology Core Rooms 3537

**Mission:** To provide high quality individualized histological services in order to facilitate productivity and success of investigators based in the Rangos Research Center and researchers in other units of Pitt / UPMC, as well as clients in other academic institutions in the Pittsburgh area.

#### Website:

https://www.pediatrics.pitt.edu/research/cores-and-research-support/histology-core

#### **Scientific Director:**

Sunder Sims-Lucas, PhD (sunder.sims-lucas@chp.edu; sus58@pitt.edu) Ph: 412-692-9440

#### Manager:

Dan Bushnell (daniel.bushnell@chp.edu)

#### **Histologist:**

Michele Mulkeen (mlm159@pitt.edu)

#### Student worker

Sarah Su

## Services offered

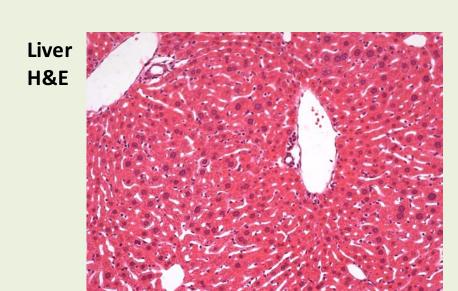
#### **Routine Histological Services:**

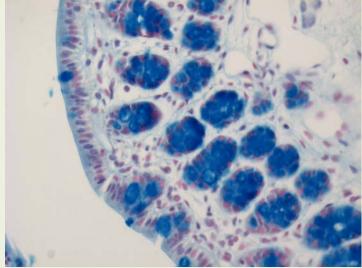
```
Routine Histology including:
Tissue processing (frozen and paraffin)
Tissue embedding
Sectioning (blank slides and staining)
```

Routine histological stains Hematoxylin and Eosin

Special stains
Oil red O
Trichrome
PAS
Toludine blue
Alcian blue

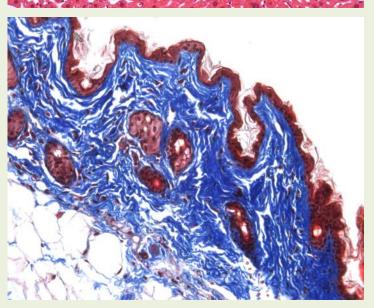
# Routine/Special Histological stains

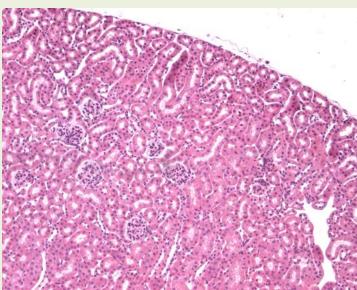




Small Intestine Alcian Blue

Skin Trichrome





Kidney PAS

## Services offered

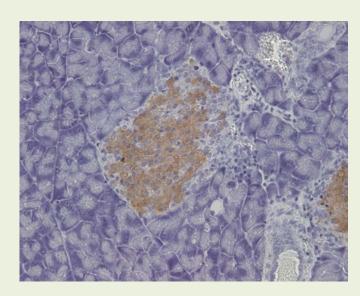
- Immunoperoxidase staining:

   IHC and IF staining with working protocols for Kim1, cd31, endomucin, f4/80, MHCII, and caspase-3, and quite a few more
- Antibody work up:

   Antibody optimization for those
   antibodies without a current working
   protocol
- Other services upon request:

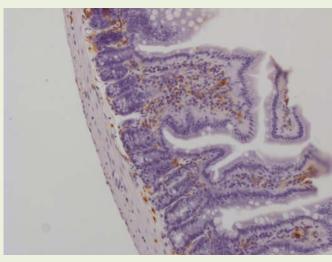
# Immunoperoxidase staining Antibody work up

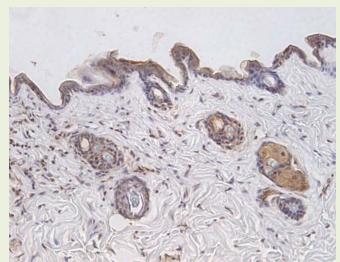
Pancreas Insulin



Kidney PMP70

Small intestine MHCII





Skin Smad2

## **Cell Imaging Core**

Zeiss LSM 710



Olympus Vivaview



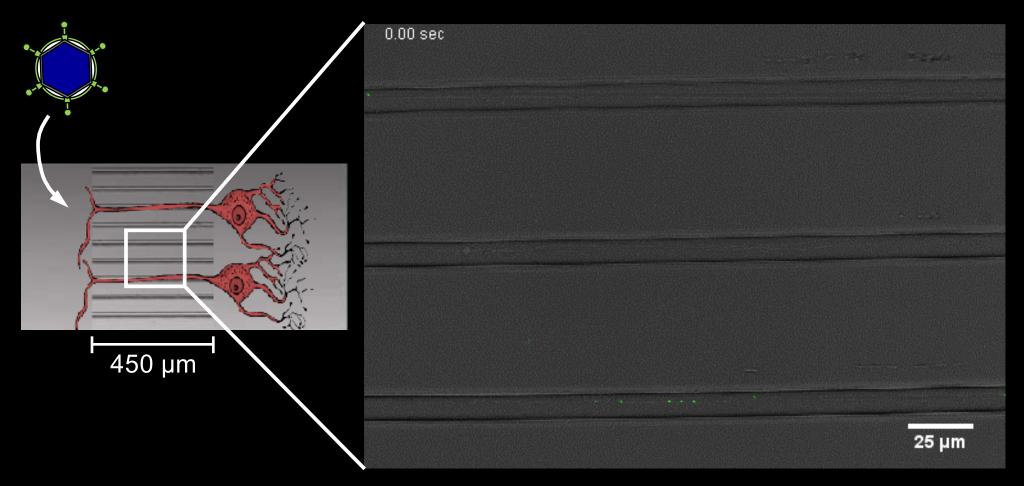
Leica TCS SP8



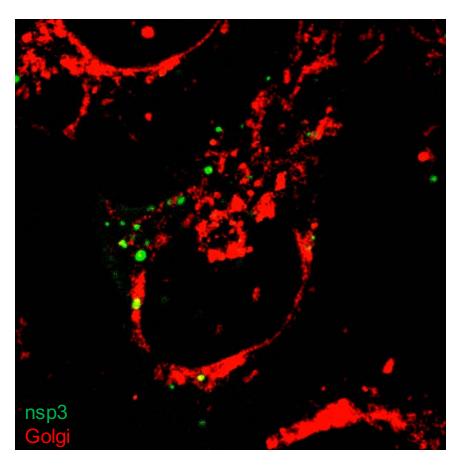
Olympus MMI CellCut laser microdissection (LMD)

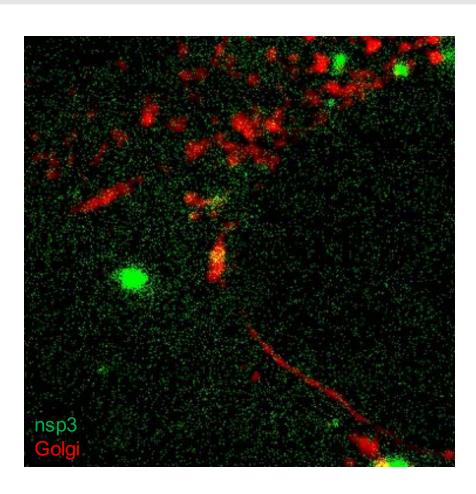


# Live Imaging Shows Single Reovirus Particles Transported in Neurons Cultured in a Microfluidic Device

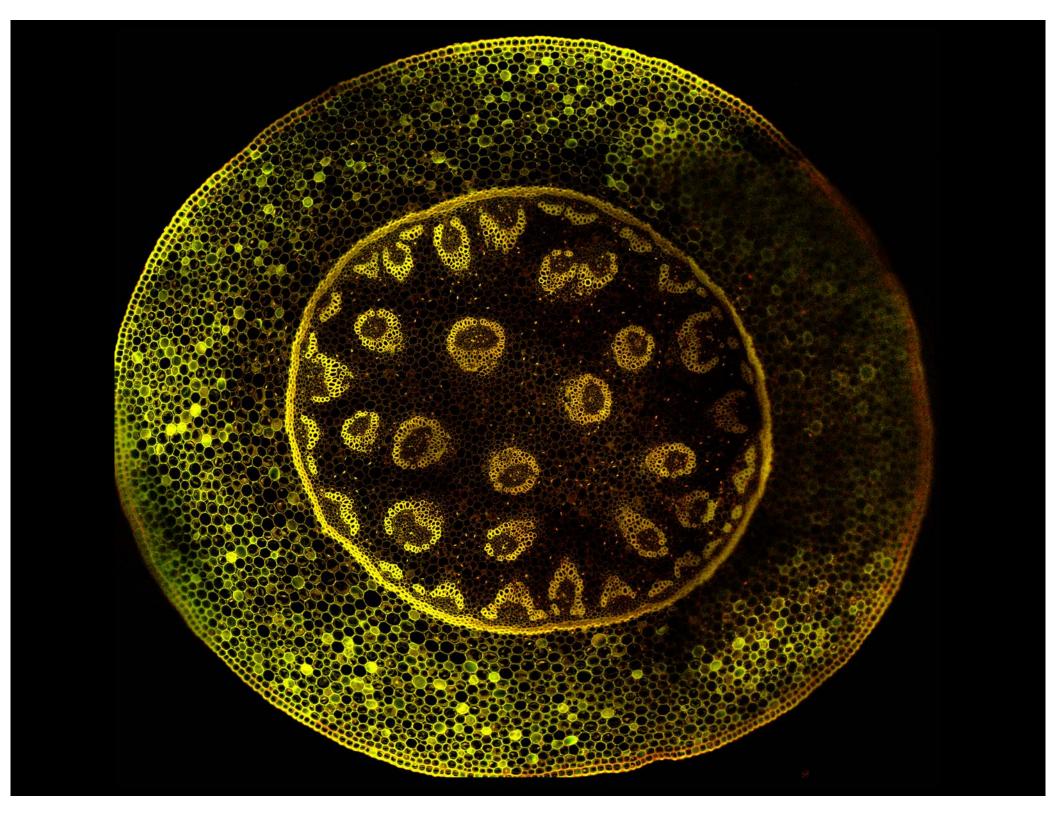


## Rotavirus protein NSP3 nsp3 co-traffics with golgi marker





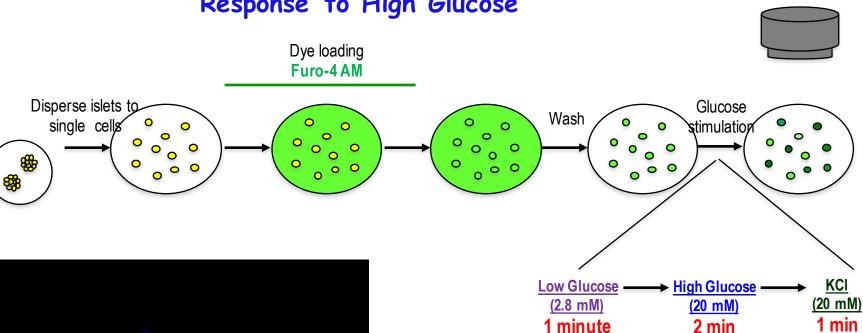
181-GFP, 5MOI U2-OS, 7.5 8.5 hpi



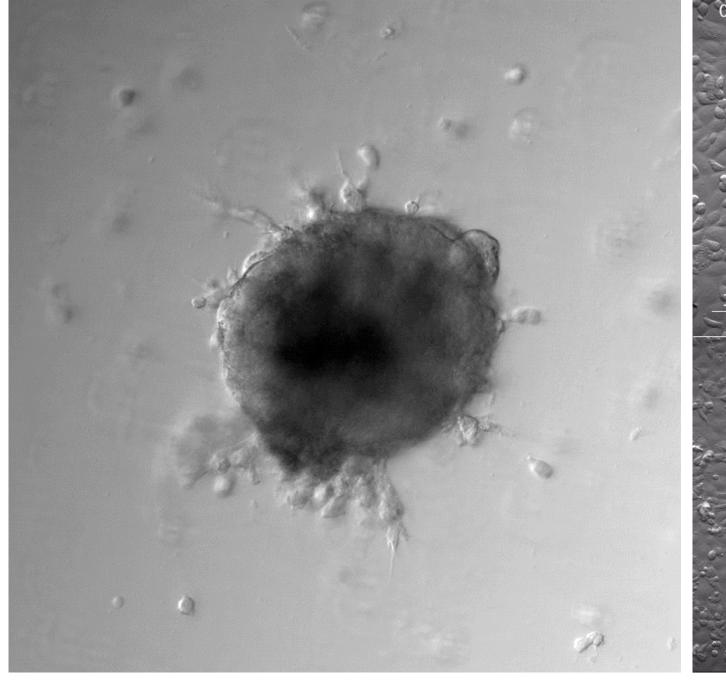
## Calcium Imaging

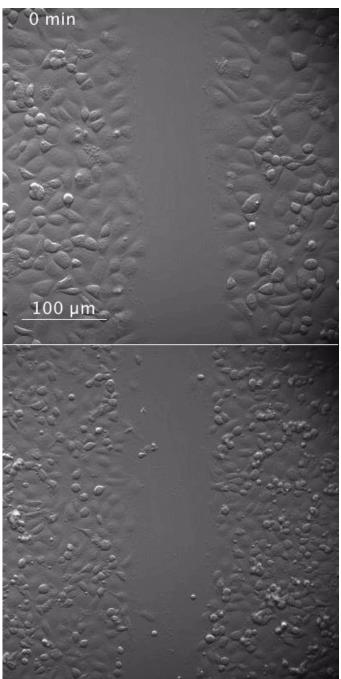
Zeiss LSM 710 confocal

# FoxO1 Deficient Islets Have Reduced Calcium Release Response to High Glucose



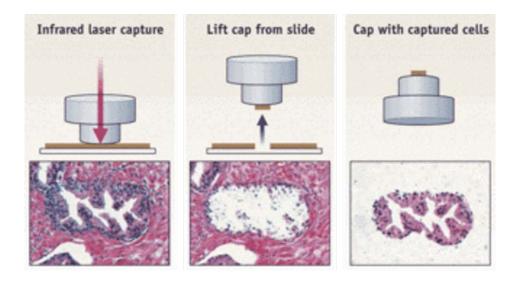
# Live cell imaging using Olympus Vivaview





# Laser capture imaging





# Cell Imaging Core 5<sup>th</sup> floor room 5148

Contact: Krishna Prasadan

Krishna.prasadan@chp.edu 412-692-9211

- Training is required to get access to the imaging core.
   First training is free. Re-training is charged \$60/- per hour.
- The core microscopes can be reserved for up to three hours at a time during regular working hours 8 am-5 pm, and longer duration after regular hours. The facility can be accessed 24/7 throughout the year.
- The flat rate is \$30 per hour, and the minimum time that can be reserved is one hour.
- Consultation is available for experiment setup or for grant submission.

## Flow Cytometry Core

John G. Rangos Sr. Research Center Rooms 8148 and 6148

Mission:

To provide congenial and individualized cytometry services in order to facilitate productivity and success of investigators that are based in the Rangos Research Center and researchers in other units of Pitt/UPMC, as well as clients in other academic institutions in the Pittsburgh area

Email contact: flow1core@chp.edu

Online scheduling: <a href="https://rangosflowcytometrycorechp.setmore.com">https://rangosflowcytometrycorechp.setmore.com</a>

#### Cytometrists:

Josh Michel (Room 8148, Tel: 692-6968)

Alex Styche (Room 6148, Tel: 692-3025)

Scientific Director: Abbe N. de Vallejo PhD (andv26@pitt.edu; vallaj@upmc.edu)

#### Flow cytometry

Technology platform employed to study cells, organelles, or molecules by taking advantage of cell-intrinsic refractive property and/or fluorochrome signals of molecular probes when they are exposed to light.

#### **Uses**

- Cell phenotypes or changes thereof (surface, cytoplasmic, nuclear)
- Cellular activities/responses (e.g. cell cycle, mitochondrial function, DNA damage)
- Cell sorting (isolate live cells for further experimentation)
- Bacteria, mycoplasma, viral particles
- Identify/quantify molecules in solution
- (Pharmaco)Kinetics of receptor-ligand interactions
- Dying (apoptotic) cells
- Membrane bound vesicles, cell organelles (mitochondria), isolated chromosomes
- Flow FISH

#### Flow cytometry

#### **SERVICES**

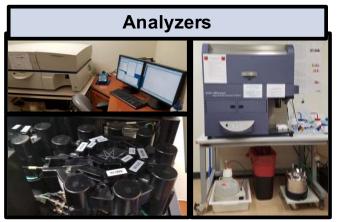
Electronic transfer of cytometry data to PI through the PI-only access subdirectory (\*courtesy service requiring IMS authorization)

Services offered (BSL1, BSL2, BSL2+ specimens; NO BSL3 specimens)

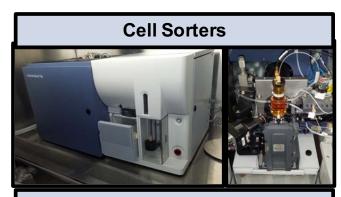
- Staff-assisted analysis and cell sorting
- Training for self-users on the analyzers, and "super-users" on the cell sorters
- Training on the use of Flow-Jo software
   (access to Flow-Jo require annual user-license fee; contact Dewayne Falkner
   falkner@pitt.edu in the Dept Immunology)
- Off-line data analysis
- Design / troubleshooting of cytometry protocols

BSL2+ cytometry work requires PI-authorized research personnel to submit IRB(PittPRO) [or IACUC] plus IBC approval information

Flow Cytometry Core
John G. Rangos Sr. Research Center
Rooms 8148 and 6148



Cytometer	Configuration	n set up	OS
Fortessa	5 lasers	18 PMTs	Windows 7
	355nm (UV)	2	
	405nm (violet)	6	
	488nm (green)	2	
	561nm (yellow)	5	
	640nm (red)	3	
LSRII	3 lasers	•	DiVa 8.0.1
	405nm (violet)	2	Windows 7
	488nm (green)	4	
	640nm (red)	2	



8th Floor and 6th Floor					
Cytometer	Configuratio	n set up	os		
Aria IIu	3 lasers	16 PMTs	Windows 7		
	405nm (violet)	8			
	488nm (green)	5			
	640nm (red)	3			
Aria II SORP	5 lasers	16 PMTs	Windows 7		
	355nm (UV)	2			
	405nm (violet)	5			
	488nm (green)	5			
	561nm (yellow)	2			
	640nm (red)	2			

## **Flow Cytometry Core**

John G. Rangos Sr. Research Center Rooms 8148 and 6148

#### **GENERAL GUIDELINES**

- a) User-training is required regardless of background
- b) Access to cytometry rooms by badge swipes (sharing of badges is prohibited)
- c) Enforcement of Biosafety as required by EH&S and by the COVID-19 committee
   \*PPE: Face mask, gloves, lab coats
  - \*Physical distancing; Disinfectant wiping of work areas including keyboards/mic
  - \*Entry logbook: Sign in/out
- c) Prior on-line scheduling required
- d) Submission of job request and safety disclosure form
- e) Penalty charges for no-shows, overtime, or undertime Schedule appropriately; call the core staff for schedule changes 24 hours BEFORE anticipated change (\*penalty fee CANNOT be charged to federal grants)
- f) Machine clogs, software failures, and other emergent problems must be reported to the core staff immediately

#### Metabolic Core

https://www.pediatrics.pitt.edu/research/cores-and-research-support/metabolic-core

Rangos Rooms 5157, 5156 & 5151

Clint Van't Land 412-692-7652

Kaitlyn Bloom 412-692-9133

clv19@pitt.edu / clinton.vantland@chp.edu

knb52@pitt.edu / kaitlyn.bloom@chp.edu

The Metabolic Core supports investigators involved in the study of energy metabolism, metabolic pathways, and cellular metabolic function in cells from biological fluids, cell cultures and tissues.

#### Staff-assisted Service(s)

Seahorse extracellular flux analyzer

#### Staff-conducted Services

- LC/MS/MS (tandem mass spectrometry)
- Amino Acid analyzer
- HPLC with UV(PDA)/fluorescence/electrochemical detectors

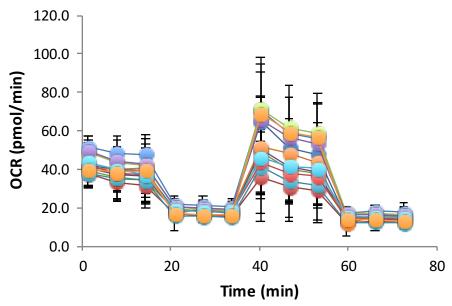
#### Self-conducted Services

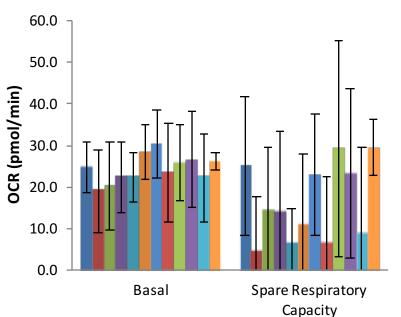
- Spectrofluorometer
- Spectrophotometer

#### **Project Consultation**

Technical consultation will be required prior to the initial (first) sample submission.

#### **Mitochondrial Respiration**

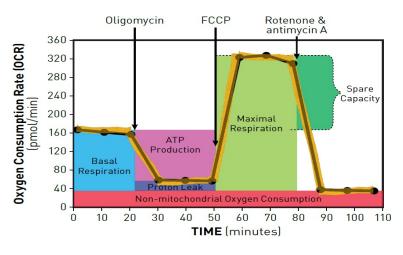


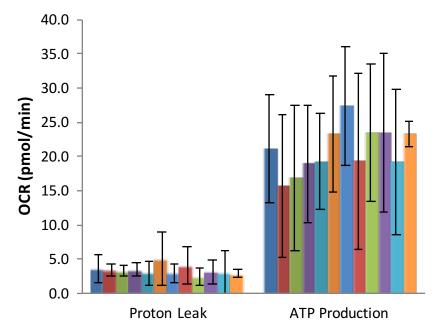


#### This slide contains unpublished data

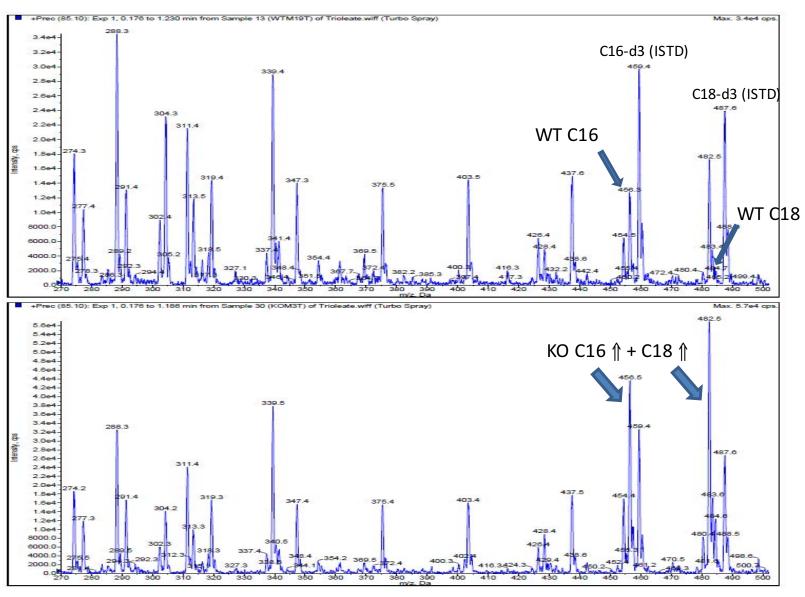
#### **Seahorse XF Cell Mito Stress Test Profile**

Mitochondrial Respiration



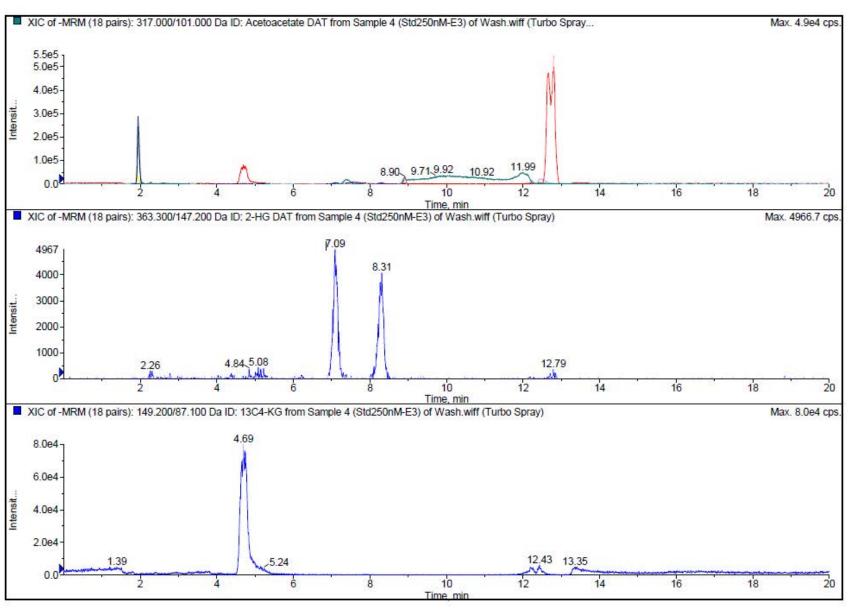


# LC/MS/MS Assay: Acylcarnitine Profiling WT & KO Mouse serum after feeding fatty acid oil for {N}-days



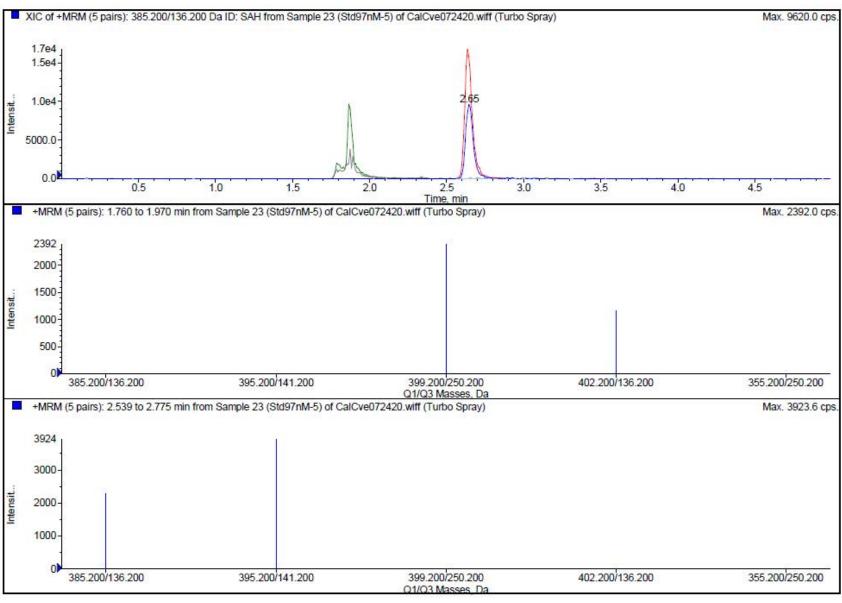
This slide contains unpublished data

# LC/MS/MS Assay: (D & L)-2-hydroxyglutarate "oncometabolite" D- & L-2-Hydroxyglutarate (253 nM) / [13C4]-Ketoglutarate (505 nM) standard solution



This slide contains unpublished data

# LC/MS/MS Assay: S-Adenosylmethionine (SAM → SAH) SAM and SAH (≈ 97 nM) / SAM-d3 (112 nM) & SAH-13C10 (218 nM) standard solution



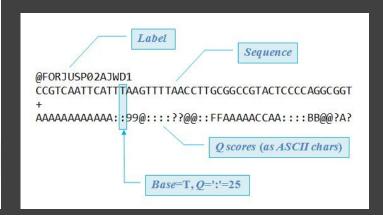
# Health Sciences sequencing core

@ Children's Hospital of Pittsburgh Rangos 8<sup>th</sup> floor

Website: http://nextgen.pitt.edu/







## HSSC @ CHP

- STAFF BASED AT RANGOS AND THE GENOMICS RESEARCH CORE
- PERFORM THE SAME SERVICES.
- CONSULT WITH THE MOST CONVENIENT FACILITY

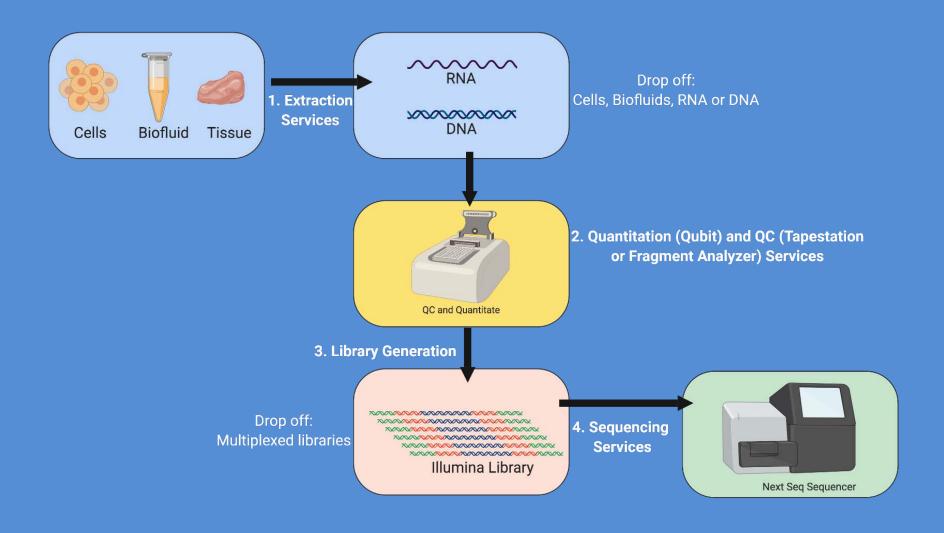
Children's Team: 8th floor Rangos, Rm 8145

- Dr. Amanda Poholek, PhD Director <u>poholeka@pitt.edu</u>
- Dr. Will MacDonald, PhD Assistant Director w.a.macdonald@pitt.edu
- Rania Elbakri Technician Rania. Elbakri@chp.edu

Oakland Team (Genomics Research Core - Forbes ave.):

- Janette Lamb Director, GRC
- Debby Hollingshead Technical Director, GRC
- Bryan Thompson Technician, GRC
- Heidi Monroe Technician, GRC

## **SEQUENCING WORKFLOW**



### LIST OF SERVICES OFFERED

#### 1. RNA/DNA EXTRACTION

TISSUE, CELLS, WHOLE BLOOD, SALIVA, BIOFLUID

#### 2. QUALITY CONTROL

- ADVANCED ANALYTICS FRAGMENT ANALYZER
- AGILENT TAPESTATION 2200
- QUBIT FLUOROMETER

#### LIBRARY PREPARATIONS

- мRNA
- TOTAL RNA
- RNA Access (FFPE) TARGETED SEQUENCING
- SMARTER-SEQ v4 ULTRA-LOW INPUT RNA- CLONTECH
- SMALL RNA
- NEXTERA/NEXTERA XT DNA
- CHIP-SEQ
- ATAC-SEQ

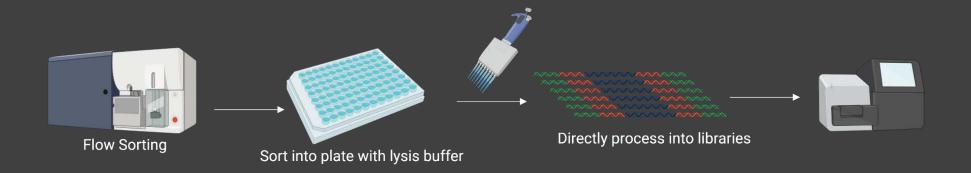
#### 4. SEQUENCING

- NEXTSEQ 500
  - HIGH OUTPUT (400M READS)
  - MID OUTPUT (130M READS)
- MISEQ (OAKLAND)
  - 16S SEQUENCING
  - TCR SEQUENCING

#### Other Projects:

- Amplicon Sequencing
- ATAC-seq
- ChIP-seq
- Cell free DNA
- Cell free RNA
- Bulk mouse TCR sequencing
- CUT&RUN (low cell number ChIP-seq assay)
- Crispr library screens
- If you have a sequencing project, please contact us. We want to help you go from sample to sequencer

## **SMART-SEQ LOW-INPUT RNASEQ**



#### Starting material:

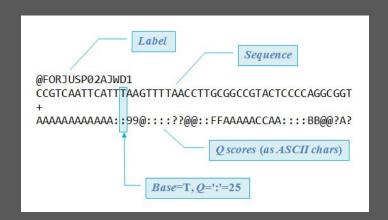
- 1-1000 cells
- 10pg-10ng RNA
- CHP Flow Core has experience working with us on this, we can also prepare plates of lysis buffer for other sort facilities

### WHAT DO WE DELIVER TO YOU?

## **Data!!** (Raw.....)

- Fastq files main file output from the sequencer
  - Direct to CRC cluster path of your choice
  - Pitt Box folder
  - Hard Drive
  - Etc ..... we will deliver to you in whatever is best option for you

Additional resources:
Next Generation Sequencing
@ The University of Pittsburgh
<a href="http://www.hscrf.pitt.edu/ngs">http://www.hscrf.pitt.edu/ngs</a>

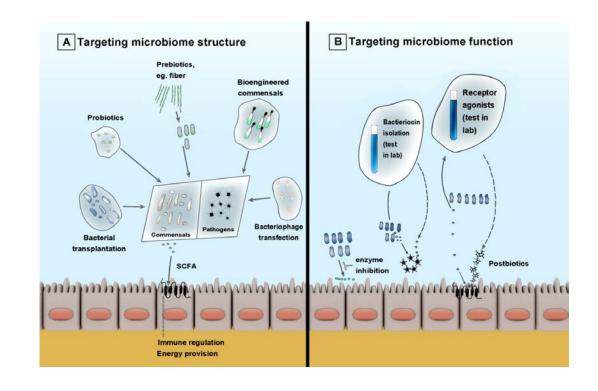


#### **Gnotobiotic Core**

# Microbiome – new paradigm for understanding biology and medicine

#### **Therapeutics**

- -Pro-biotics
- -Fecal transplant
- -Pre-biotics
- -Engineered bacteria
- -Bacteriophages
- -Metabolite/drug discovery



**Director:** <u>timothy.hand@chp.edu</u>

Manager: javonn.musgrove2@chp.edu

## **Gnotobiotic mice**

- Latin for 'known life'
- What does that mean in practice?
  - Germ-free (GF) mice completely lack all culturable micro-organisms, no bacteria, archaebacteria or fungi.
  - GF mice can be colonized with micro-organisms (single, multi, communities) to test whether a particular organism or microbiome is necessary and sufficient to produce a phenotype.
  - Germ-free or gnotobiotic mice are kept in isolators or bubbles so as to maintain their microbiological status



#### What can you do with gnotobiotic animals? Koch's postulates for the microbiome



I think gut microbes are associated with fast running?

Associate mice with intestinal microbes

Measure response

### What is the 'University of Pittsburgh Gnotobiotic facility'?

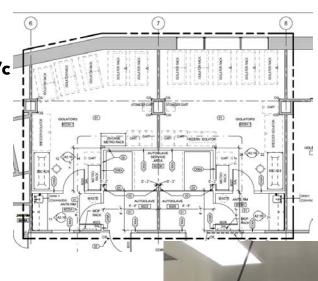
#### In BST3

- Currently have established germ-free C57BL/6 and Balb/c breeding stock.
- Serviced by three highly trained staff:
  - Facility Manager Javonn Musgrove
  - Technicians Eugenio Alvarez and Abby Mulhorn

#### Two Mirrored Facilities

- Limits the effect of autoclave breakdowns that cause contamination
- We have 10 isolator carts (each with 1-4 bubbles)
  - 2 breeder isolators (30+ cages)
  - 1 're-derivation' isolator
  - 2 'double' carts (isolators hold 15+ cages)
  - 5 'quad' carts (isolators hold 5 cages)

Total = 24 experimental isolators (capacity for 20 individual conditions)



### What can the gnotobiotic facility do for your research?

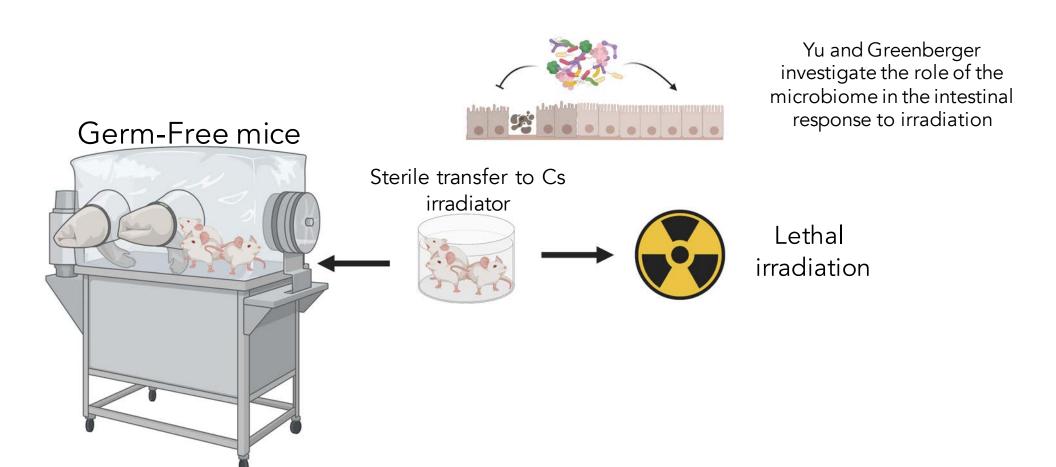
#### Strains in the facility:

- Initially, we will breed and provide C57BL/6 and Balb/c.
- If there is interest we could also acquire Rag1-/- IL-10-/- and Swiss Webster.

#### • Services that we will be able to provide:

- Javonn Musgrove is highly skilled and has 9+ years experience as technician at NIH, 5+ years in GF facility. He is happy to learn your favorite technique. **Let us say yes to your research.**
- Facility will be certified as BSL2, so human samples can be used
- We will be able to associate with any mouse microbiome you're interested in (no BSL3/4)
- Can associate microbiome from intestine, skin, mouth, lung, urogenital tract, eye
- Will be able to use BSL2 pathogens
- We can treat with drugs, biologics
- Can autoclave and administer custom diets.
- We can adoptively transfer cells. We can irradiate mice for bone marrow chimeras.
- Can inject **tumors** for cancer treatment studies
- Our staff can carry out all procedures (we can draw blood/feces etc.).
- At conclusion of experiments mice can be euthanized within biosafety cabinet and tissues removed to your own facility
- BONE MARROW CHIMERAS

# Technical breakthroughs in the Pitt Gnotobiotic facility – Gnotobiotic bone marrow chimeras



**Director:** <u>timothy.hand@chp.edu</u>

Manager: javonn.musgrove2@chp.edu

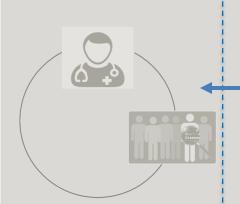
### **Bioinformatics Core**

Director: Dhivyaa Rajasundaram, PhD

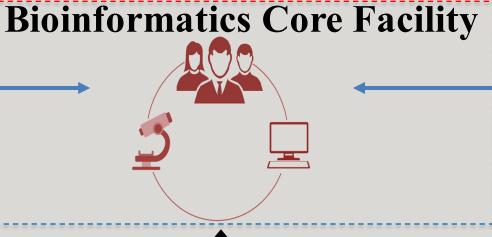
# Cross-facility workflow

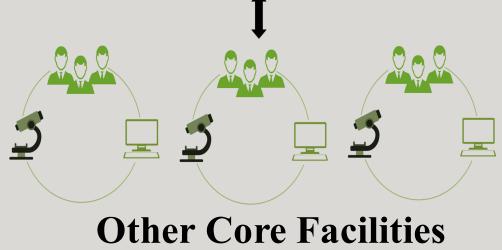


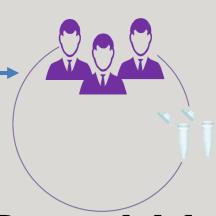




**Physicians** 







Research labs

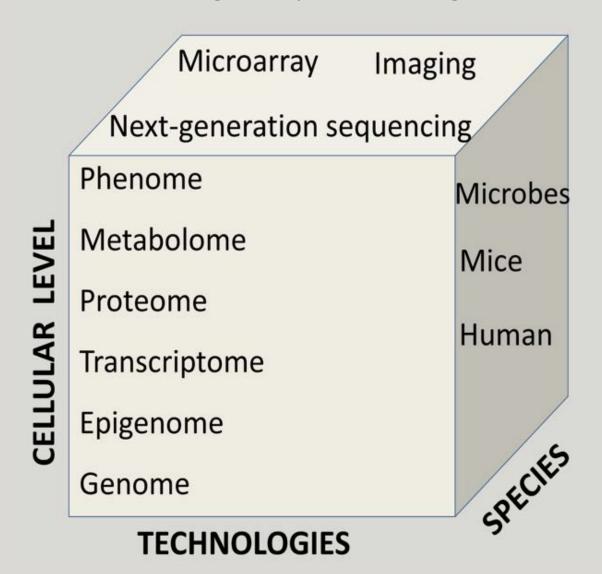
### Services offered



- □ Study design implementation
- High-throughput data analysis
- ☐Statistical support
- Proposal writing
- ☐ Faculty collaborations

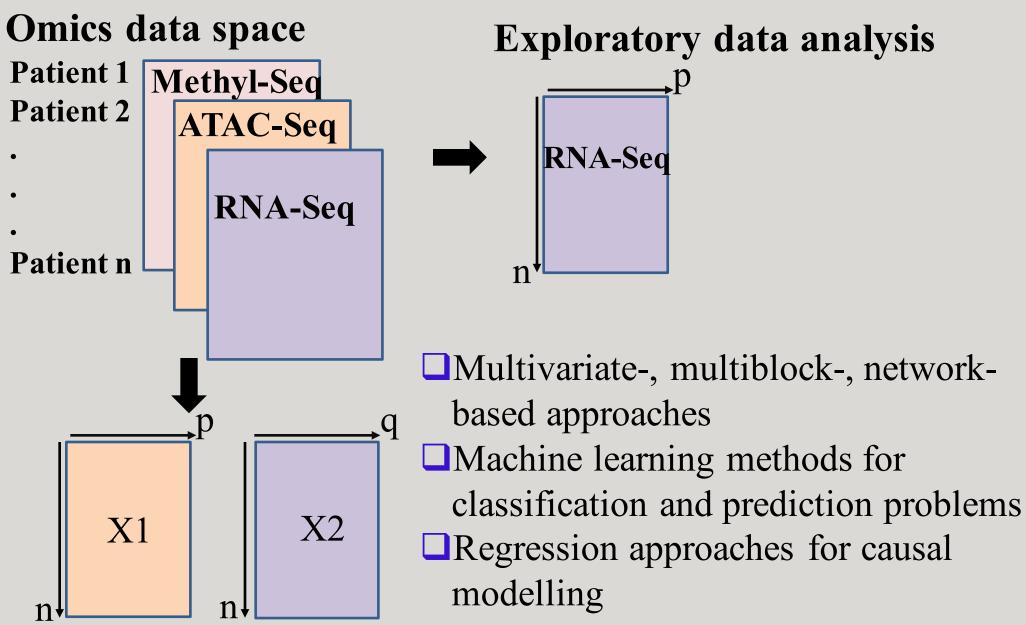
## High-throughput data analysis

Transform high-throughput data from clinical, translational, and basic research into biologically meaningful information.



## Statistical support





Integrative omics analysis

### Core users



- □ 15 unique users
- □ 27 projects completed in fiscal year 2019 2020
- □ Data submission forms
  <a href="https://www.pediatrics.pitt.edu/research/cores-and-research-support/bioinformatics-core">https://www.pediatrics.pitt.edu/research/cores-and-research-support/bioinformatics-core</a>
- □ Data transfer Pittbox/Teams or hard disk

## Leveraging publicly available data



□ Landscape of tumor infiltrating T-cell repertoire of pediatric brain tumors (using bulk RNA-Seq) - **Dr. Gary Kohanbash** 

#### **CBTTC PEDIATRIC BRAIN TUMOR ATLAS**

1ST RELEASE - LARGEST COLLECTION OF PEDIATRIC BRAIN TUMOR DATA



**30** brain tumor types



Data from over **1,000** patients



Releasing data in **REAL-TIME** 



More than **16** partner institutions



Over **50** foundation sponsors



Includes **WGS**, RNAseq, Proteomics, Clinical, Imaging & Histology Data











## Training



- □Pitt CRC Fall 2020 workshops
- Differential expression analysis Dhivyaa September 24, 2020 (tentative) (1-4p)
- □ Single cell RNA-Seq analysis (beginner) Dhivyaa October 20, 2020 (1-4p)
- □ Single cell RNA-Seq analysis (Advanced) Dhivyaa October 22, 2020 (1-4p)

### Cost model



□ Project based pricing, and percent effort on grants

Project	Examples	СНР	Non-CHP
size		users	users
Small	RNA-Seq analysis	\$1500	\$2000
Medium	Exome-Seq data analysis, CHIP-Seq analysis, ATAC-Seq analysis, Time-course RNA-Seq analysis	\$2500	\$3000
Large	Single-cell sequencing data analysis, T cell repertoire analysis, integrative omics using multivariate statistics	\$4500	\$5000

## Practical aspects



- □Do we offer software based (Partek, CLC,...) services?
- Do we mentor students who are interested to learn their own analysis?
- Do we have to pay for the analysis even if we do not get the expected result?
- Could two projects be charged for the price of one because the analysis pipeline is the same?
- Why do we require authorship?

# Rangos Virus Production

#### **Dr. George Gittes**

- AAV serotypes 1-10 and DJ. Custom serotypes available as well.
- Adenovirus also available
- Expected titer 10<sup>13</sup>-10<sup>16</sup>vg/ml, volume of 0.5-1.0ml
- Turnaround time 2-3 weeks
- Investigator must have IBC approval prior to virus production
- Fees: Reagents billed to the investigator (~\$1000/virus). Brief subcontract will be arranged to pay salary of the virus producer (~\$700/virus). Commercial virus typically ~\$3000.