

Sixth Annual Interdisciplinary Quantitative Biology Winter Boot Camp

# SINGLE-PARTICLE CRYO-EM

ORGANIZED BY





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## **PROGRAM**

**January 7 – 11, 2019** 

Institute for Quantitative Biomedicine (IQB)

Rutgers University | Busch Campus Proteomics Building 174 Frelinghuysen Rd GPS Address: 679 Hoes Lane West Piscataway, NJ 08854-8076

igb.rutgers.edu/bootcamp

#### Introduction

### 6<sup>th</sup> Annual Interdisciplinary Quantitative Biology Winter Boot Camp

## SINGLE-PARTICLE CRYO-EM

The Institute for Quantitative Biomedicine, is delighted to be hosting the Sixth Annual Rutgers Interdisciplinary Quantitative Biology Boot Camp. This year's workshop is focused on Single-Particle Cryo-Electron Microscopy (Cryo-EM).

The 2019 immersive one-week Boot Camp has been designed to provide participants with: (1) a fundamental understanding of the cryo-electron microscope and its capabilities and limitations; (2) an overview of sample preparation techniques for single-particle Cryo-EM applications; (3) an introduction to high-throughput data collection workflows; (4) familiarity with the computational approaches and tools used for processing Cryo-EM image data and determining and interpreting three-dimensional structures derived therefrom.

Instruction will take the form of morning lectures on the theory and applications of Cryo-EM, addressing the advantages of this modern form of microscopy, and methods for analyzing and interpreting the data obtained. In addition to the lectures and invited presentations, a broad range of collaborative hands-on afternoon workshops will equip attendees with the skills needed to uncover the three-dimensional structures of complex biological machines.

The Boot Camp would not have been possible without tireless efforts of the many contributors listed pages 8-10 of this Program.

Financial contributions from Angstrom Scientific, Leica, NanoSoft, OHAUS, Thermo Fisher Scientific and Waters Corporation are much appreciated.

Cover images sources: D. Sirohi *et al.* (2016) A. Merk *et al.* (2016)

#### Locaction:

All sessions will be held or initiated in Proteomics, Room 120

#### Laptops:

Laptop computers are required. See page 11 for software specifications.

#### **Activity Legend**







Lab Demo



9:00 AM		Welcome S. David Kimball
9:10 AM		Welcome & Overview Stephen K. Burley and Arek Kulczyk
9:20 AM	İ	Cryo-EM Resolution Revolution from the Perspective of the Protein Data Bank Stephen K. Burley
10:15 AM		Coffee Break
10:30 AM	İ	Structures of Biological Macro-molecules by Single-Particle Cryo-EM: The roots of the "resolution revolution" Marin van Heel
11:30 AM	İ	Cryo-Electron Microscopy 101: Why, What and How Arek Kulczyk
12:30 PM		Lunch & Coffee/Tea
1:30 PM	5	Practical Session (1 hour)
2:45 PM		Image Processing (Coffee/Tea provided)
4:15- 5:00 PM	1	Poster Session

## Tuesday, January 8<sup>th</sup>

9:00 AM	İ	A Service Vision on Microscope Setup Gijs Janssen
10:00 AM		Coffee Break
10:15 AM	İ	Strategies for Large Scale Data Collection and On-the-fly Processing at SEMC William Rice
11:00 AM	İ	Recent Xmipp/Scipion advances in Image Processing for Single Particle Analysis. Carlos Sorzano
11:45 AM		Panel Discussion Led by Stephen K. Burley and Arek Kulczyk Panel members: Marin van Heel, Gijs Janssen, William Rice, and Carlos Sorzano
12:30 PM		Lunch & Coffee/Tea
1:30 PM	\$	Practical Session (1 hour)
2:45 PM		Image Processing (Coffee/Tea provided)
4:15- 5:00 PM	The state of	Poster Session

9:00 AM	İ	Cryo-Electron Microscopy and Tomography of the E.coli DnaB - phage Lambda P Complex David Jeruzalmi
10:00 AM		Coffee Break
10:15 AM	İ	<b>Sample Preparation for Cryo-EM and CLEM</b> Wei Dai
11:15 AM	İ	Microscope Tuning For High Resolution Data Collection Sean Mulligan
12:15 AM		Lunch & Coffee/Tea
1:30 PM	\$	Practical Session (1 hour)
2:45 PM		Image Processing (Coffee/Tea provided)
4:15 - 5:00 PM	T. II	Poster Session

## Thursday, January 10th

9:00 AM	Ť	<b>Challenges and Opportunities for EM Automation</b> Bridget Carragher
10:00 AM		Coffee Break
10:15 AM	İ	Integrated Approach to Membrane Surface Complexes: from NMR to EM Andrew Byrd
11:15 AM	İ	Preserving the Native State Nancy Rizzo
12:15 PM		Lunch & Coffee/Tea
1:30 PM	5	Practical Session (1 hour)
2:45 PM		Image Processing (Coffee/Tea provided)
4:15- 5:00 PM		Poster Session
5:15 PM	That I	Networking at Harvest Moon (Hors d'oeuvres provided) 392 George St, New Brunswick

9:00 AM	İ	Cryo-EM: An Imaging Tool Beyond Crystallography Wah Chiu
10:00 AM		Coffee Break
10:15 AM	İ	Cryo-EM Data Resources, Validation and Challenges Cathy Lawson
11:15 AM	İ	Emerging Cryo-EM methodologies: FIB-SEM Jason Kaelber
12:15 PM		Lunch & Coffee/Tea
1:30 PM	\$	Practical Session (1 hour)
2:45 PM		Image Processing (Coffee/Tea provided)
4:15- 5:00 PM		Closing remarks Stephen K. Burley and Arek Kulczyk

#### Contributors

## The organizers would like to extend a special thank you to the Boot Camp contributors:

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#### Additional Information

#### Software

Boot Camp Participants will be using their own laptops. Calculations will be performed on Linux clusters at Rutgers.

Please make sure you have the following software installed PRIOR to the start of Boot Camp:

- Web Browser: current generation of Safari or Google Chrome
- UCSF Chimera: free for academic use; follow download and installation instructions at www.cgl.ucsf.edu/chimera/ download.html

#### Recommended items:

- 1. Power supply cords
- 2. External mouse (very useful for structural visualization)

#### Wi-Fi

#### Rutgers affiliates:

Use RUWireless with NetID

#### **Guests:**

Network:

**Proteomics Guest** 

Password:

1766Rutgers1997Proteomics

#### Sakai

Sakai access will be set up for all full-time registered participants.

#### **Parking**

**Rutgers students, staff, and faculty:** Park according to your permit.

#### Visitors:

Park in Busch **Lots A & B** located at 679 Hoes Lane West (see map on the back cover).

Visitors must register their vehicles to park on campus at bit.ly/2LdTDGA

#### Questions?

Registration and logistics: bootcamp@iqb.rutgers.edu

The program: arek.kulczyk@rutgers.edu

