

Network Biology Collaborative Centre Fundamentals and Applications Workshop

Tuesday, November 22nd, 2016 60 Murray Street, L3-201, Mount Sinai Hospital

Workshop #1: High-Throughput Screening & Imaging

Fundamentals:	
9:00 am	Towards challenges and rewards of high-throughput screening: current and future amenities, Alessandro Datti, High-throughput screening manager
9:30 am	Microscopy-based screening: from high content to high resolution and beyond,
	Mikhail Bashkurov and Monica Hasegan, High-content screening & high-resolution imaging managers
Application	ons:
10:00 am	May the Light be with you: using LUMIER to unveil novel signaling pathways from protein interaction networks, Miriam Barrios-Rodilles, Wrana lab
10:25 am	Morning Break
10:45 am	Going beyond the diffraction limit of light; Using super resolution microscopy to study centrosomes, Bahareh Adhami, Pelletier Lab
11:10 am	Identifying activators of the neuroprotective E3 ligase Parkin, Natalia Moskal, McQuibban lab
11:35 am	The beauty of High-throughput screening in drug discovery: Hippo pathway, Dohee Lee, Wrana Lab

12:15 pm Complimentary Lunch

Workshop #2: Proteomics

Fundamentals:		
1:00 pm	Success in proteomics: an overview of LC-MS, Brett Larsen, Proteomics manager	
1:30 pm	Finding the diamonds in the rough: from raw data to compelling and high-	
	confidence hits, James Knight, Software engineer	
Application	ons:	
2:00 pm	Expanded proteomic toolbox, Payman Samavarchi-Tehrani, Gingras lab	
2:25 pm	Afternoon Break	
2:40 pm	Mapping endosomes with BioID, Geoffrey Hesketh, Gingras lab	
3:05 pm	Using data-independent acquisition mass spectrometry in interactome mapping experiments, Jean-Phillipe Lambert, Gingras lab	
3:30 pm	Using Kinobead Technology to Assess Kinome Reprogramming in Ovarian Cancer Cell Lines in Response to Cisplatin Treatment, Kevin Brown, Rottapel lab	

4:00 pm Our Facilities in Action – join a tour!