







# Genomics - One Cell at A Time Opening of the Single Cell Analysis Core Sunday, October 22, 2017 Yaglom Senate Hall



#### 10:00 - 11:00

Special seminar **Oliver Hobert**, Columbia University & HHMI: Organizational principles of *C. elegans* nervous system development

# 11:00-11:15 COFFEE

11:15 Opening remarks
Yoav Henis, Vice President for Research and Development
Karen Avraham, Vice Dean, Sackler Faculty of Medicine, Co-Director, SCA Core Lab

# 12:45 – 13:15 LUNCH

# 13:15 - 13:45

**Hila Sharim** (Ebenstein lab, TAU) - Single-molecule genome mapping of 5-hydroxymethylcytosine allows single cell like population clustering

# 13:45 - 14:15

**Franziska Paul** (Amit lab, WIS) - Single cell genomic analysis of hematopoietic progenitors: Towards a refined model of hematopoiesis

# 14:15 -14:45

epigenetic memory

**Amos Tanay** (WIS) - Single cell epigenomics and

11:30 – 12:15
Plenary speaker
Sarah Teichmann (European Bioinformatics Institute, EMBL-EBI) - Immunogenomics one cell at a time

12:15 – 12:45Oren Ram (HUJI) - From single cells to clones:Advances in droplet-based microfluidics

Organizers: Karen Avraham, Yuval Ebenstein, Irit Gat-Viks, Lior Mayo, Anat Moshayov, Naomi Caplan

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