



Icahn
School of
Medicine at
Mount
Sinai

Spring School for RNA-seq Sequence Analysis

May 15th – 19th, 2017

Meet with and learn
from researchers
in the field.

- Learn how to apply computational tools to genomic sequencing data.
- Gain practical knowledge of genome browser and data analysis tools for RNA sequencing using real-life data sets.

This program is intended for early career faculty and postdoctoral fellows in biology or computer science.

There is a competitive selection process for applicants, and for those selected there is no fee to attend the course.

In addition, housing will be provided at no cost for those requiring it, and travel stipends are available for attendees traveling from outside the NY Metro area.

Topics include:

- Introduction to Scientific Computing, including Unix, Scripting and Python
- Overview of the Human Genome and Genetic Variation
- Isoform-level Analysis of RNA-seq datasets
- Analysis of Rare Variant/Exome Datasets
- Intro to Galaxy ToolKit, UCSC Genome Browser, 1000 Genomes, GWAS and WES data

Course Sponsor:

NIH Big Data to Knowledge (BD2K) 1R25EB020393

Educational costs are covered by this grant. Students accepted outside of the NY Metro area will be provided with travel and housing stipends for the duration of the program.

Location:

Icahn School of Medicine at Mount Sinai
1468 Madison Avenue,
New York, NY 10029

Eligibility:

Applicants must be U.S. citizens or permanent residents.

For more information and to apply, please visit: mssm.edu/creeds

This program strives to enhance the diversity of the biomedical big data workforce through recruitment of individuals from diverse racial, ethnic, cultural, and socioeconomic backgrounds.