

# Single Cell Genomics Day 2026

**10:00 - 10:40AM**  
**(All times ET)**

**Rahul Satija**, *NYGC and NYU*

Single-cell genomics: Recent advances and future directions

**10:40 - 11:00AM**

**Dingchang Lin**, *Johns Hopkins University*

Genetically encoded assembly recorder temporally resolves cellular history

**11:00 - 11:40AM**

**Xin Jin**, *Scripps Research*

Towards a functional brain genome

**11:40 - 12:00PM**

**Coffee Break**

**12:00 - 12:20PM**

**Sydney Blattman**, *MSKCC*

Scalable genotyping in fixed transcriptomes resolves clonal heterogeneity

**12:20 - 12:40PM**

**Amanda Urke**, *Harvard University*

Protein-guided RNA barcoding links transcriptomes to synaptic architecture

**12:40 - 1:20PM**

**Anshul Kundaje**, *Stanford University*

Deep learning the regulatory code of fetal development and cellular reprogramming from single cell chromatin accessibility

**1:20 - 2:00PM**

**Lunch**

**2:00 - 2:45PM**

**Aviv Regev**, *Genentech*

From Cell Atlases to Medicines, with AI

**2:45 - 3:05PM**

**Alex Bradu**, *NYGC*

Genome-wide single-cell perturbation screens with VIPerturb-seq

**3:05 - 3:20PM**

**Coffee Break**

**3:20 - 3:40PM**

**Kevin Chao**, *Harvard University*

TimeVault: A genetically encoded device for transcriptome storage in mammalian cells

**3:40 - 4:20PM**

**Junyue Cao**, *Rockefeller University*

Scalable genomic approaches to investigate cellular dynamics in aging and disease

**4:20 - 4:40PM**

**Moritz Schaefer**, *Stanford University*

Biomedical Data Analysis with Language-Multimodal AI



Friday June 12

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