

Single Cell Core Grant Program

Informational Webinar - Coffee and snacks provided
Genomics Auditorium 1102A
Thursday, Sept 22 @ 10:00 AM

What You'll Learn by Attending

Single cell RNA-Seq has become a required tool in today's genomics research. Droplet-based approaches are effective, but struggle against flexibility, experimental design, and scale. Join us for an informational session in which we describe the Parse Biosciences split-pool combinatorial barcoding technology and a corresponding grant program we offer. By attending, you'll learn:

- Modern single cell approaches
- Details of the grant program
- How to apply for an award

What the Award Includes

- An Evercode Whole Transcriptome (WT) kit capable of up to 100,000 cells
- Two Cell or Nuclei fixation kits capable of up to 8 samples
- Parse Biosciences Data Analysis suite for data processing and visualization
- Parse Biosciences fixation training
- Parse Biosciences data analysis training and support
- Barcoding and library preparation services from your core
- NGS sequencing (one lane of an S4 flow cell)

Speakers



Wei Zhang
Genomics Coordinator
we Zhang@ucr.edu

Brandon Le
Bioinformatics Core
lbrandon@ucr.edu



Sam O'Neill
Technical Sales Manager
samantha@parsebiosciences.com



Bo Liu
Executive Sales Specialist
bliu@illumina.com

