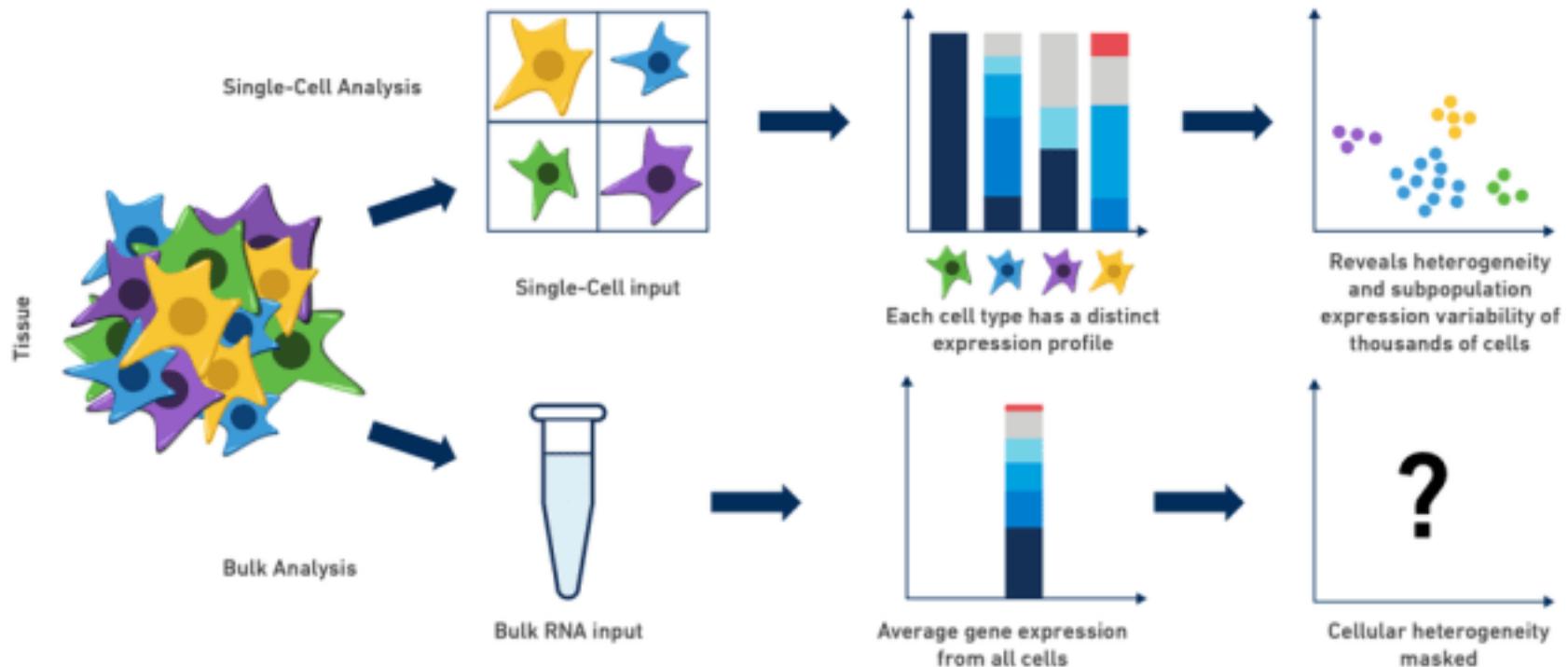


# Single Cell RNA Sequencing

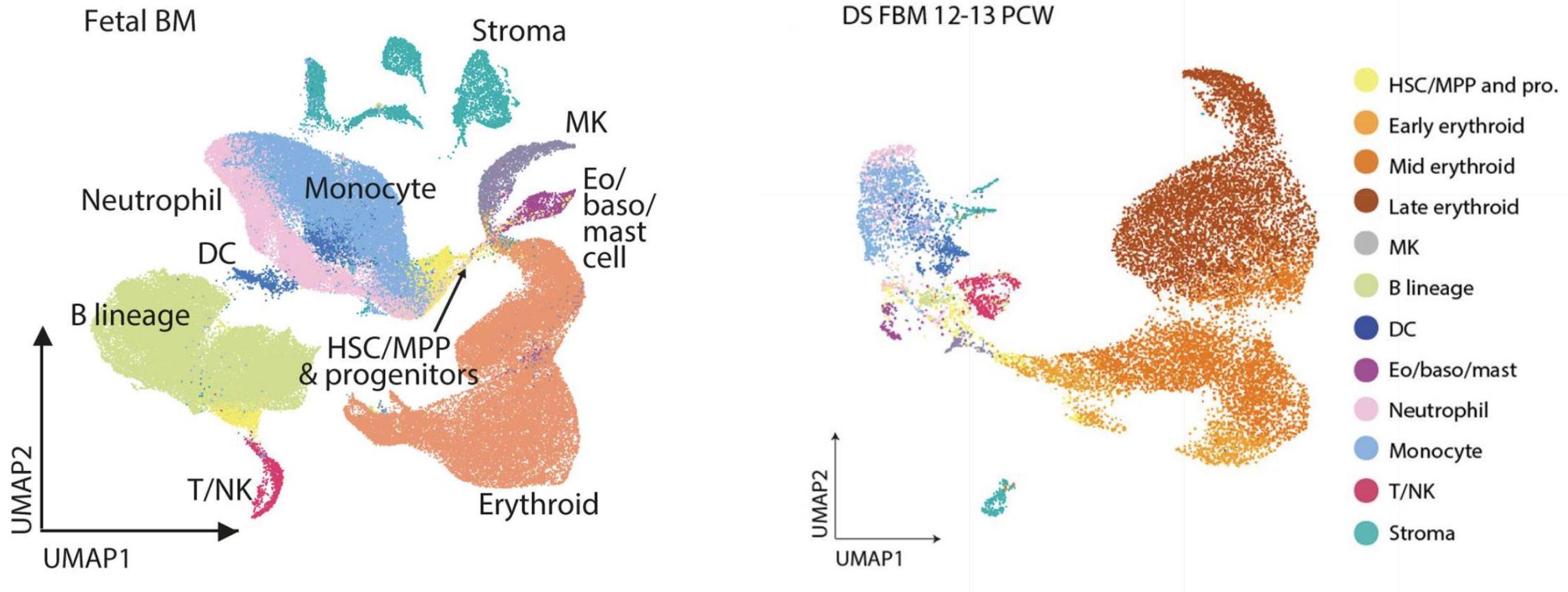
Chris Ozeroff

Short Read 2023

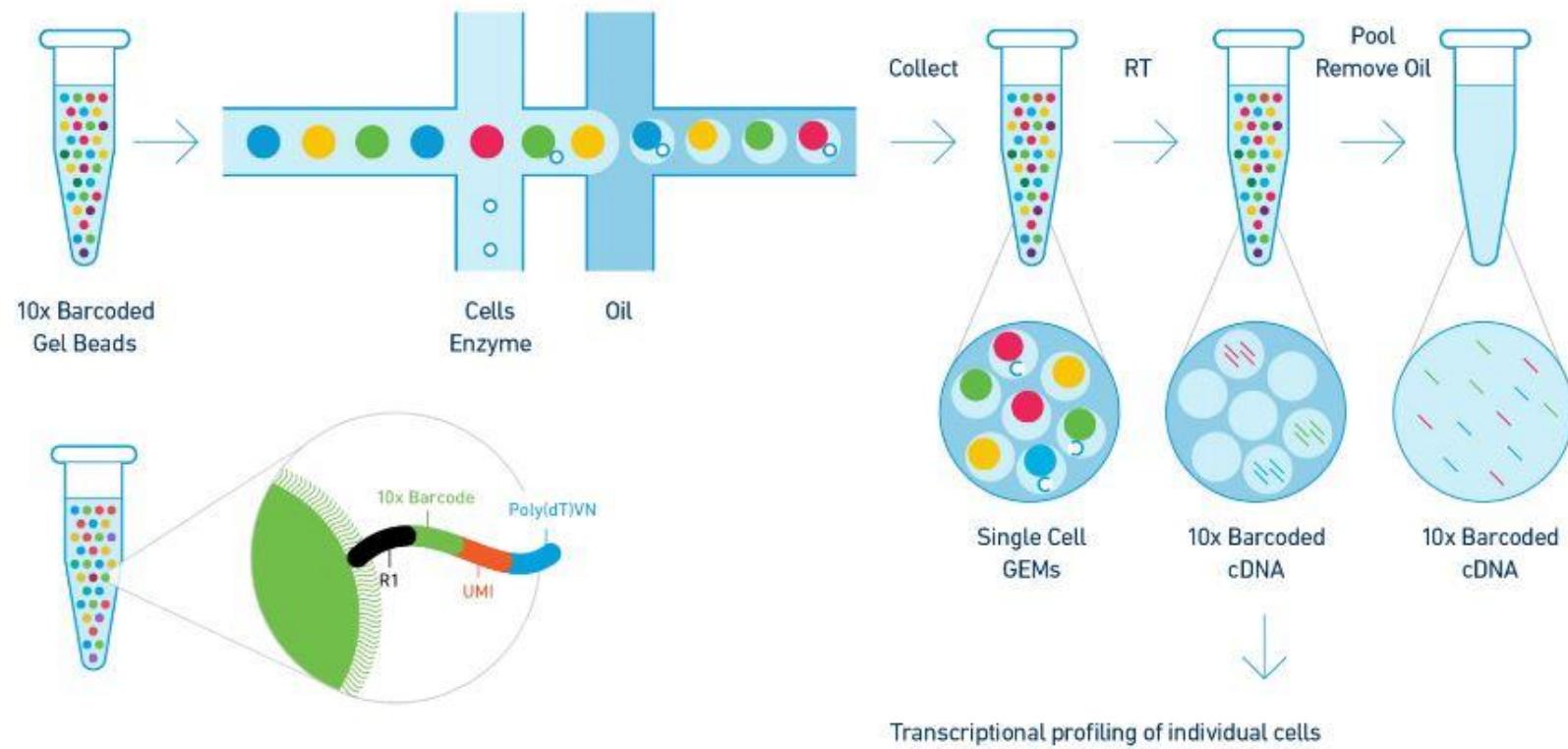
# Single Cell RNA Sequencing



# Down Syndrome vs. Typical Fetal Bone Marrow

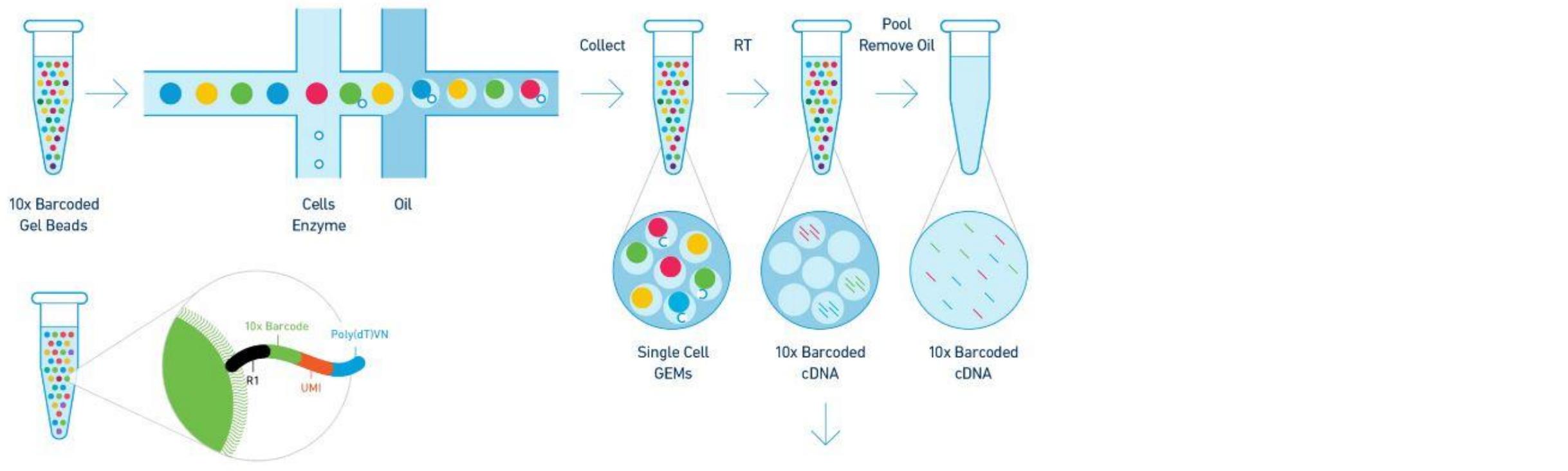


# 10X Genomics Droplet Method and Cell Ranger



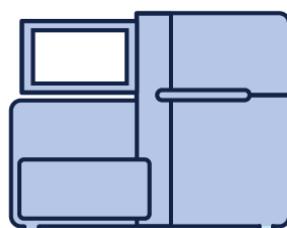
Genomics Technology Core,  
University of Missouri, 2021

# 10X Genomics Droplet Method and Cell Ranger



Genomics Technology Core,  
University of Missouri, 2021

## Sequencing



BCL



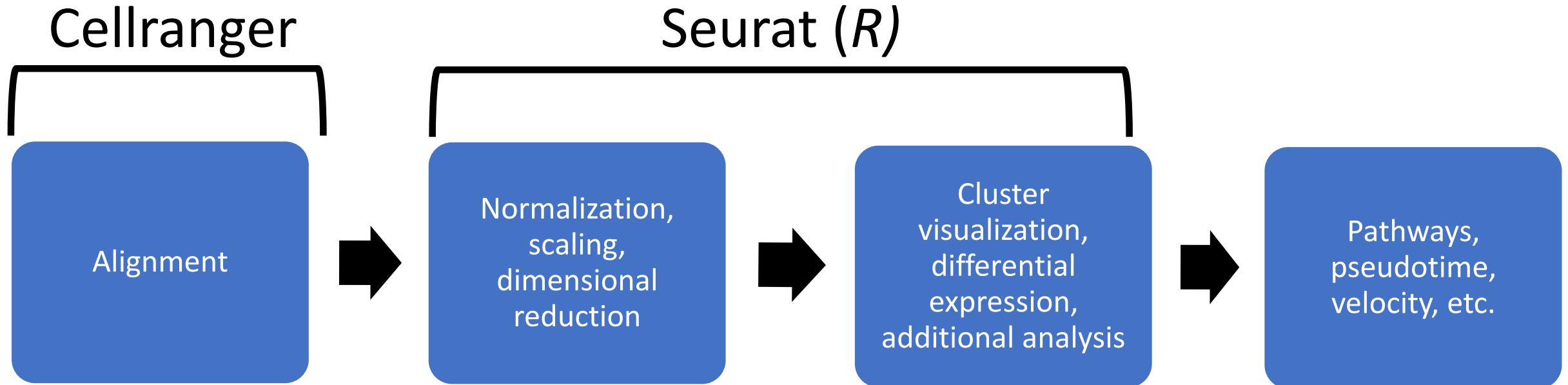
FASTQ



Output

## Cell Ranger Pipeline

# scRNA-seq workflow



**SEURAT**

R toolkit for single cell genomics



# Seurat

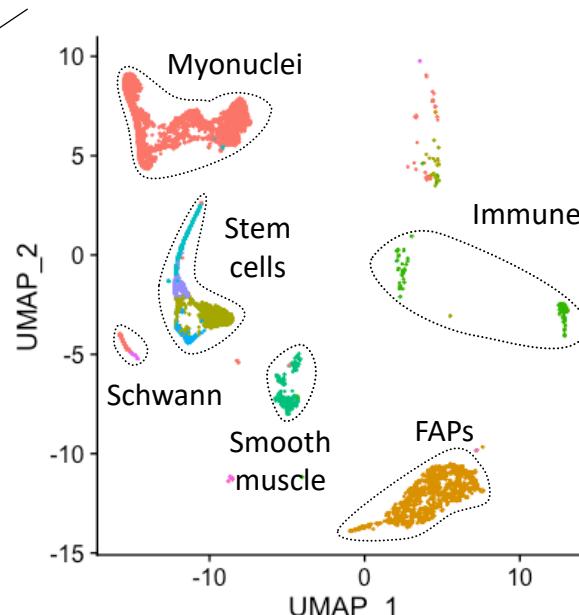
Remove debris/doublets

↓  
Normalization

↓  
Scaledata

↓  
PCA

↓  
UMAP/clustering



“...named for Georges Seurat to invoke the analogy between the intricate spatial patterning of single cells and a pointillist painting.”  
(doi: [10.1038/nbt.3192](https://doi.org/10.1038/nbt.3192))

# Seurat workflow

## Processing individual Seurat objects

- Add metadata
- Merge Seurat objects
- Normalize data
- Find genes with variable gene expression
- Scale data
- Run PCA reduction

## Integrating Seurat objects

Split Seurat objects and identify anchors (genes) to “integrate” objects

## Plotting and clustering

UMAP reduction and visualization