

# scAnalyzeR

Instructions

Upload Dataset

Pre-processing

Normalization

Dimensionality Reduction

Clustering

Differential Expression

Plots

Pathways Analysis

Trajectory Analysis

## Species

- ☒ Human  
☐ Mouse

## Source Dataset

- ☐ Use sample dataset(Mouse)  
☐ Upload dataset(.csv/.txt)  
☒ Upload files(cellranger output)

Choose 3 files (barcodes.tsv, genes.tsv, matrix.mtx)

Browse...

3 files

Upload complete

*Summary of The Uploaded Dataset:*  
*Total number of features(genes): 33694*  
*Total number of samples(cells): 8381*