

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