

# scStudio

A web portal for intuitive and flexible analysis of public scRNA-seq data

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User-friendly highly interactive scRNA-seq data analysis

Automatic retrieval of public datasets from the Gene Expression Omnibus

Modular structure to allow easy integration of new methods

## Methods



Gene Expression Omnibus

Platforms (1) [GPL19057](#) Illumina NextSeq 500 (Mus musculus)

Samples (10) [GSM5936967](#) Parabiosis

[More...](#) [GSM5936968](#) Hashed\_18h\_MQ

[GSM5936969](#) HASH-TIM43hr8hrMQ

This SubSeries is part of SuperSeries:

[GSE198057](#) Metabolic reprogramming of skeletal muscle by resident macrophages points to CSF1R inhibitors as muscular dystrophy therapeutics

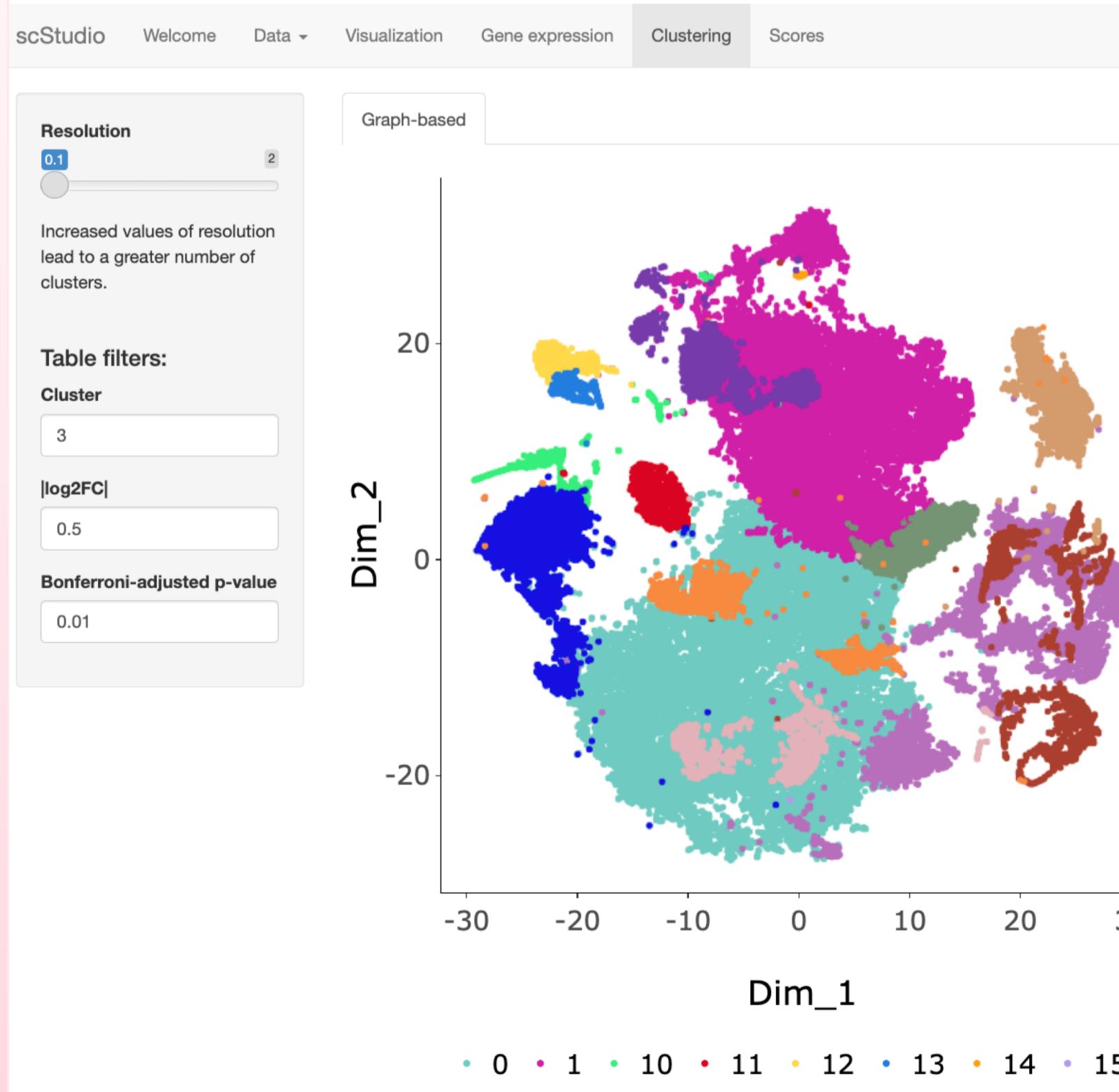
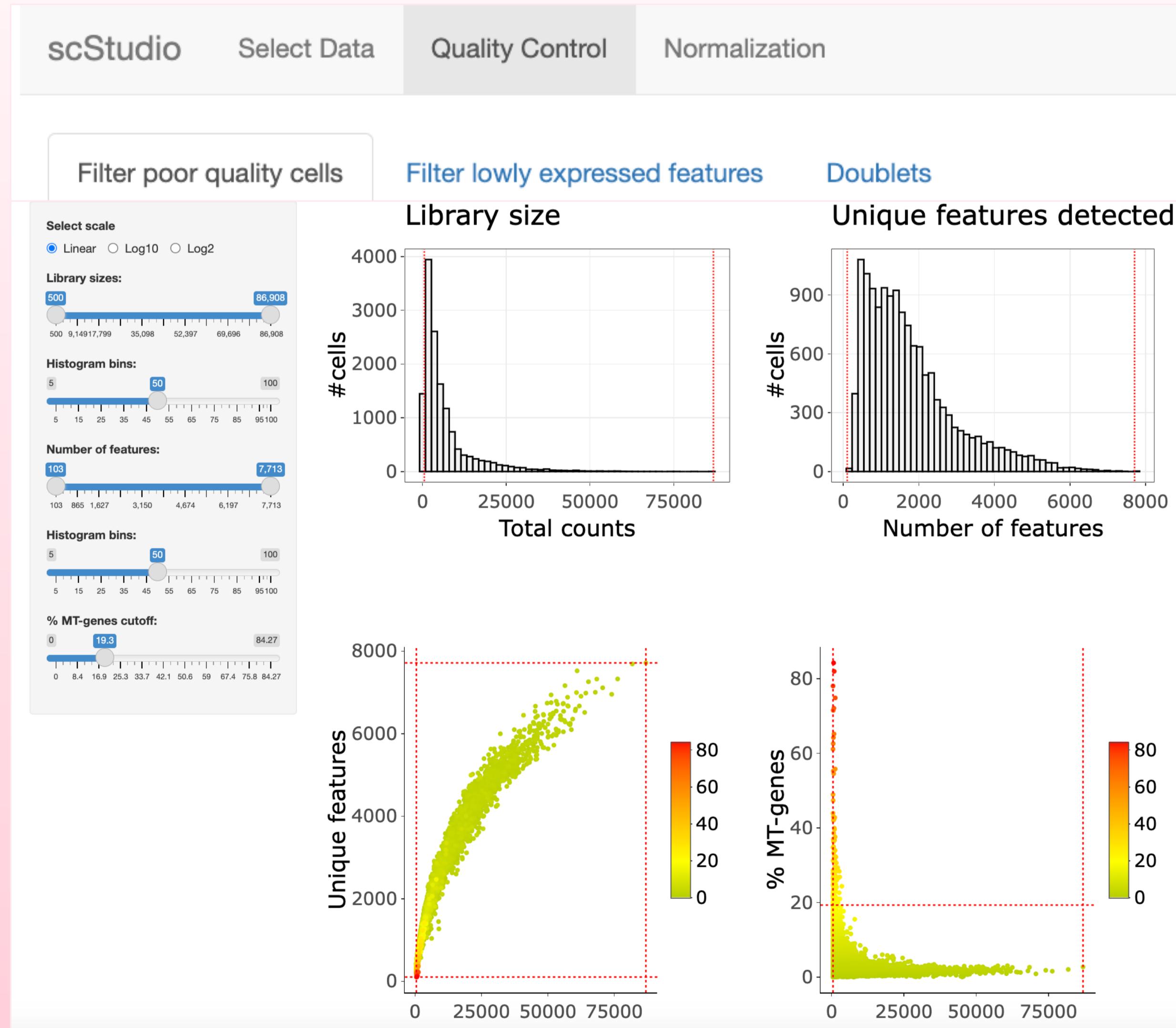
### Relations

BioProject [PRJNA813543](#)

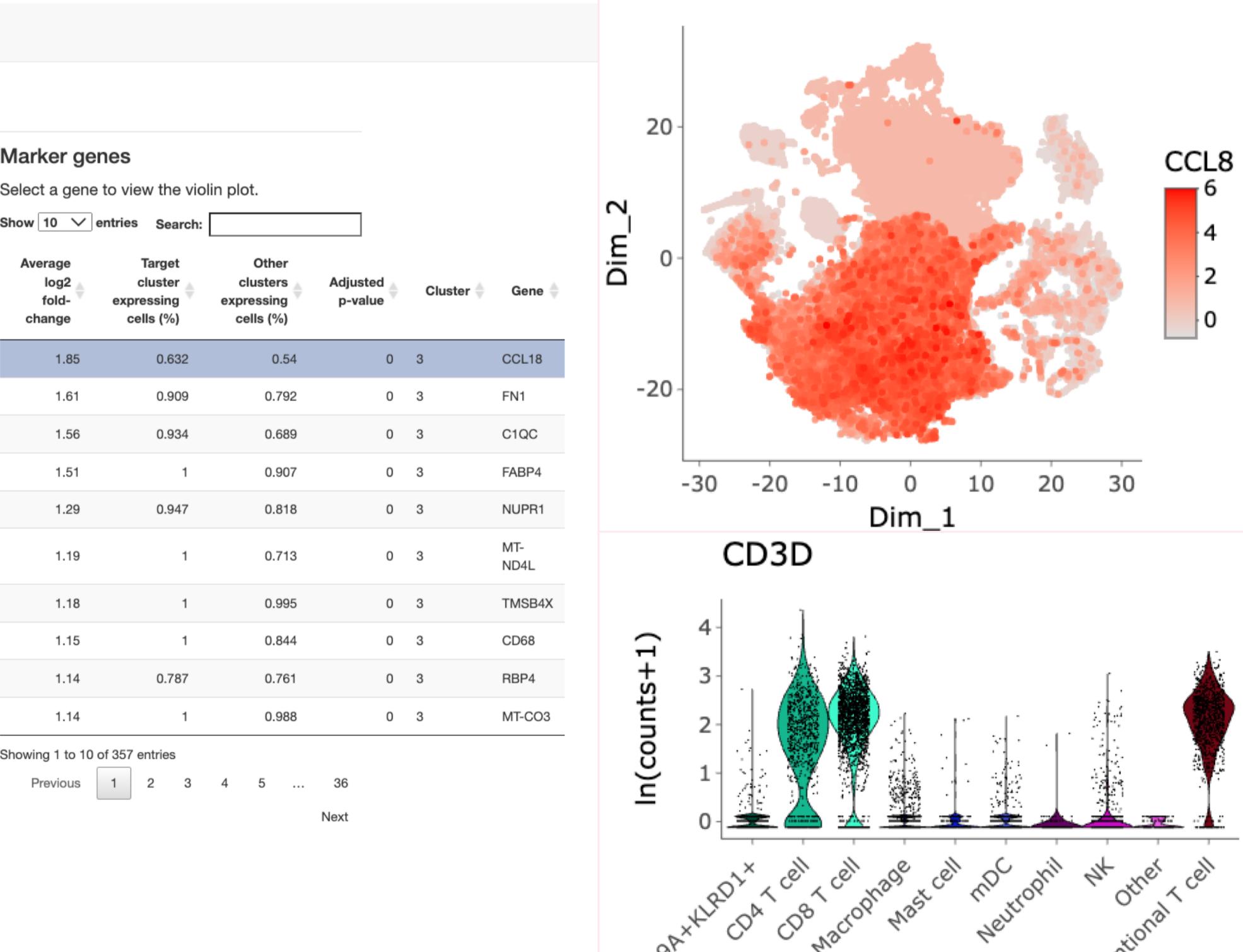
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SOFT formatted family file(s)	SOFT <a href="#">[?]</a>
MINIMI formatted family file(s)	MINIMI <a href="#">[?]</a>
Series Matrix File(s)	TXT <a href="#">[?]</a>

Supplementary file	Size	Download	File type/resource
GSE198055_RAW.tar	118.2 Mb	(http)(custom)	TAR (of H5)

## 2. Quality control and normalization



## 3. Downstream analysis



Try the demo version!

