

DECODING ULCERATIVE COLITIS AT THE CELLULAR LEVEL WITH OUR SINGLE-CELL RNA SEQUENCING DATA CURATION PORTAL

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INTRODUCTION

Our portal hosts meticulously curated Ulcerative Colitis (UC) datasets

KEY FEATURES

Request

data

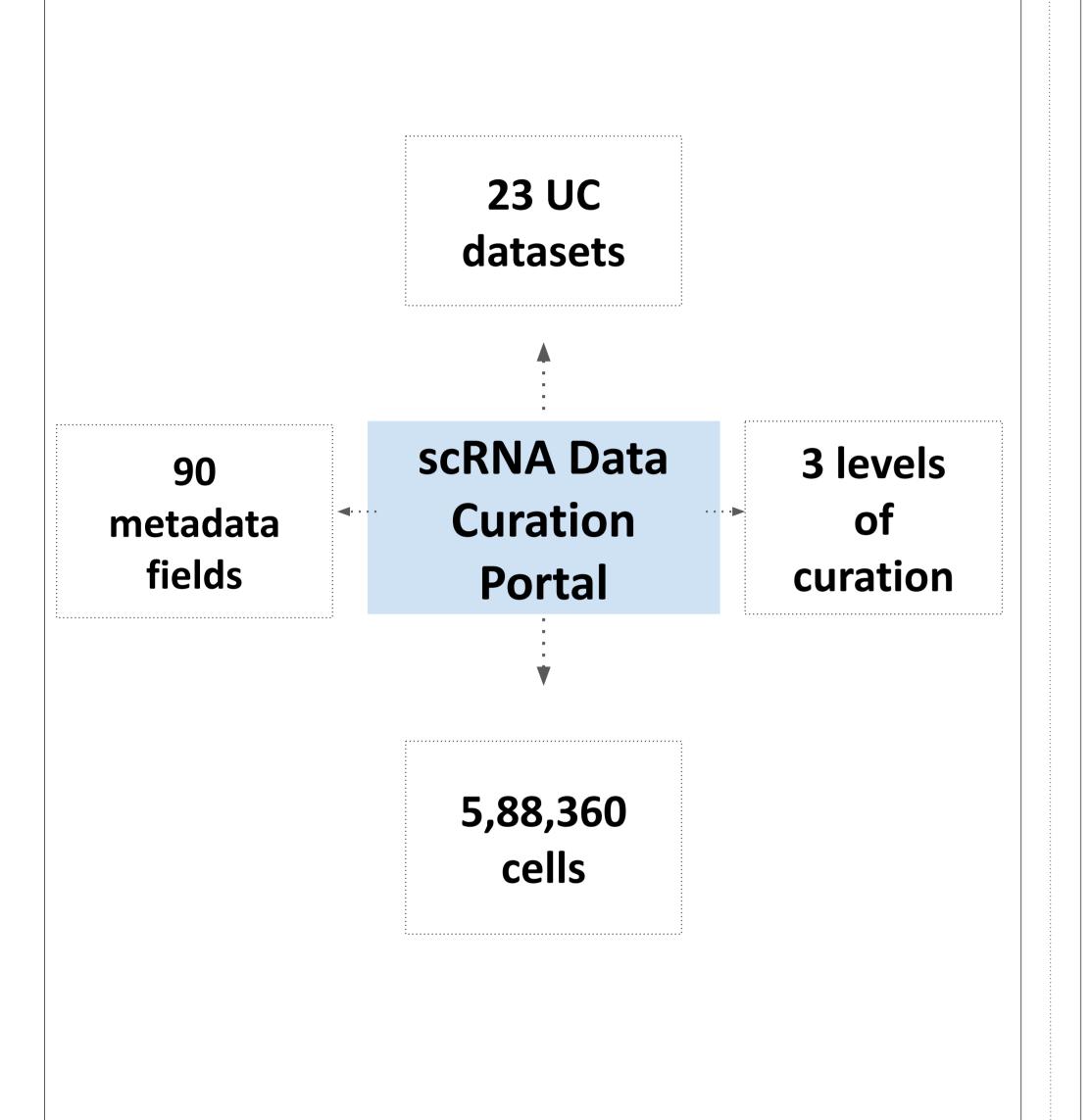
curation

Our portal has several avenues for use and is equipped with features for

FUTURE DIRECTIONS

As we continue to grow, we plan to include more disease categories and

sourced from NCBI's GEO platform.



effortless data navigation and filtering to help you identify the dataset of your choice in minimal time.

Datasets from publicly available resources	
Confidential/ in-house datasets	

Equipped with

a set of 27

filters from

diverse

metadata

fields

more resources for aggregating publicly available datasets.

> **Expand to disease categories** beyond UC, eg: **Neurodegenerative disorders**, metabolic disorders, etc

Expand to public repositories beyond GEO, eg:EMBL-EBI's ArrayExpress, The European **Nucleotide Archive**

Integrate curation with downstream analysis using our in-house analysis pipeline

CURATION & QC

Datasets are curated across three levels of curation namely study, experiment and sample levels with 10, 42 and 48 metadata fields respectively. All datasets are curated and validated independently through a 2-step process. The TAT per dataset is 24-48 hours.

Curation

Browse the data showcase

Allows simultaneous selection of multiple filters as well as multiple options within the same filter for easy navigation

CONCLUSION

Strand's scRNA data curation portal provides a platform for organizing and standardizing complex datasets, a better understanding of which, can aid in potential discoveries of biomarkers with prognostic, diagnostic or therapeutic purposes.

