

download_dataset
dataset: PapalexiSatija2021_eccite_RNA

download_dataset
dataset: NormanWeissman2019_filtered

download_dataset
dataset: ZhaoSims2021

run_your_method

run_your_method

run_your_method

all

```
graph TD; A1["download_dataset<br/>dataset: PapalexiSatija2021_eccite_RNA"] --> B1(run_your_method); A2["download_dataset<br/>dataset: NormanWeissman2019_filtered"] --> B2(run_your_method); A3["download_dataset<br/>dataset: ZhaoSims2021"] --> B3(run_your_method); B1 --> C(all); B2 --> C; B3 --> C;
```

The diagram is a flowchart with three parallel vertical paths. Each path starts with a red-outlined rounded rectangle containing the text 'download_dataset' followed by a specific dataset name. Arrows point from these rectangles to green-outlined rounded rectangles labeled 'run_your_method'. Finally, arrows from all three 'run_your_method' boxes point to a single light-green-outlined rounded rectangle at the bottom labeled 'all'.