

# Census object

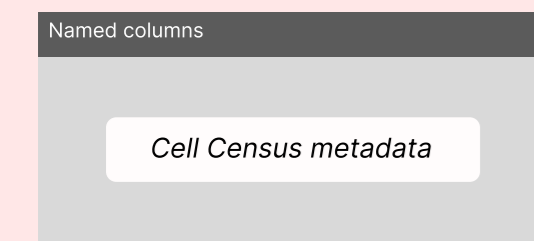
Collection

## "census\_info"

Collection

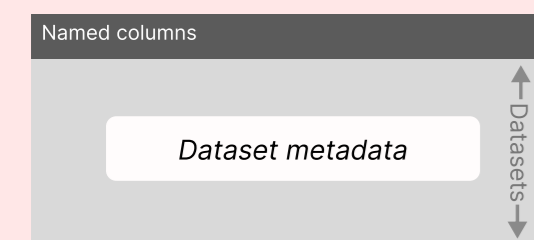
### "summary"

DataFrame



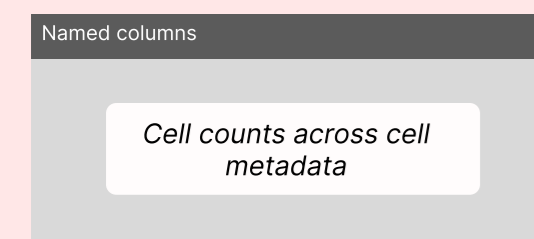
### "datasets"

DataFrame



### "summary\_cell\_counts"

DataFrame



## "census\_data"

Collection

### "homo\_sapiens" or "mus\_musculus"

Experiment (is\_a Collection)

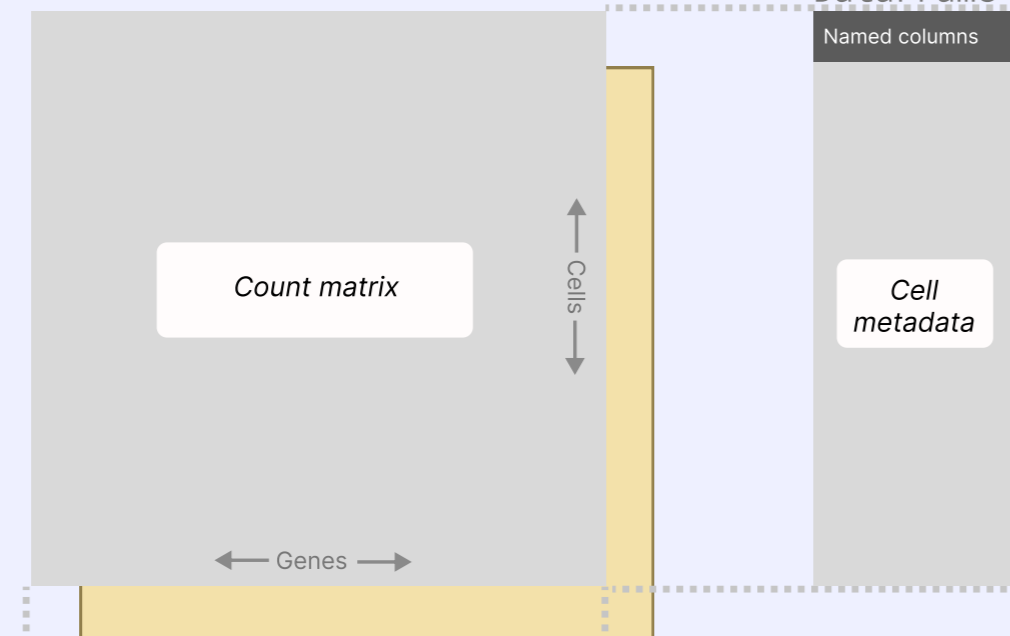
### ms["RNA"]

Measurement (is\_a Collection)

### X["raw"], X["normalized"]

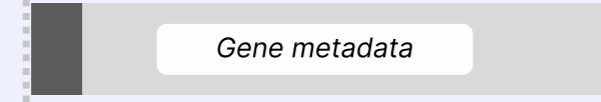
SparseNDarray

Only in >V1.1.0



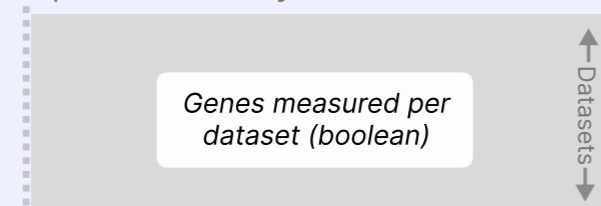
### var

DataFrame



### "feature\_dataset\_presence\_matrix"

SparseNDarray



## "census\_spatial\_sequencing" Contains Visium and Slide-seq data from CELLxGENE

Collection

### "homo\_sapiens" or "mus\_musculus"

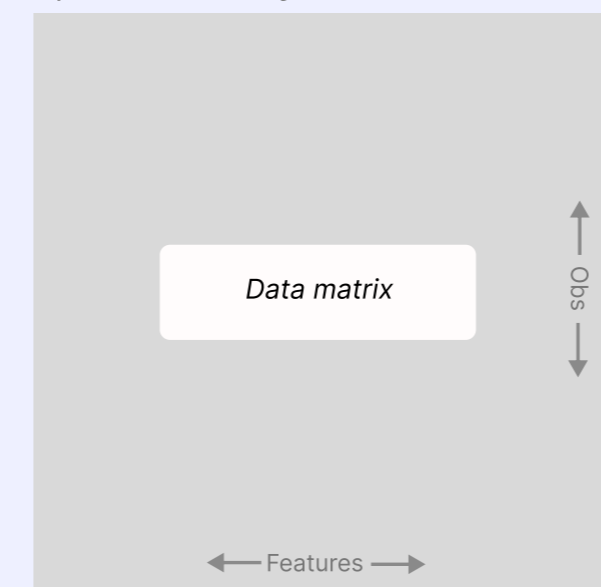
Experiment (is\_a Collection)

### ms["RNA"]

Measurement (is\_a Collection)

### X["raw"]

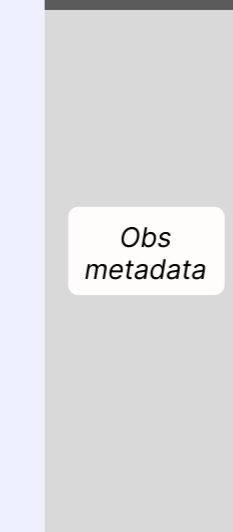
SparseNDarray



### obs

DataFrame

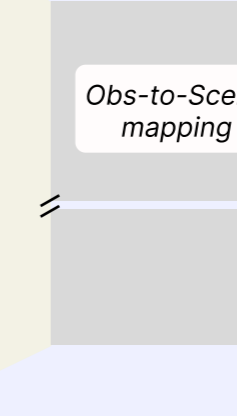
Named columns



### obs\_scene

DataFrame

Named columns

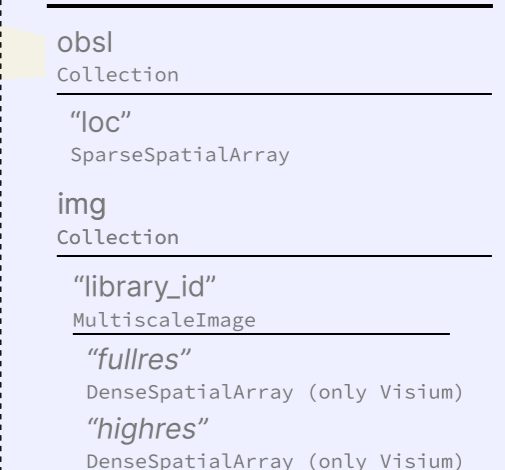


### spatial

Collection

### "library\_id"

Scene (is\_a Collection)



### "library\_id"

Scene (is\_a Collection)

