ScHARe Repository Introduction

November 20, 2024

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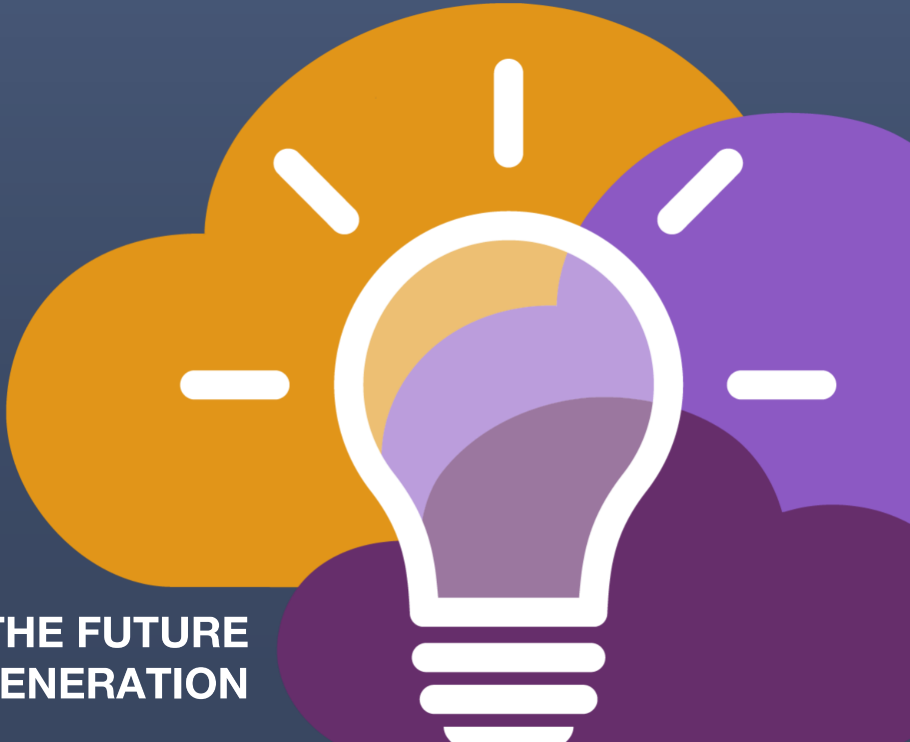
Elif Dede Yildirim, PhD · NIMHD
Mark Aronson, PhD · NIMHD

ScHARe

Data Aggregation



BE A PART OF THE FUTURE
OF KNOWLEDGE GENERATION

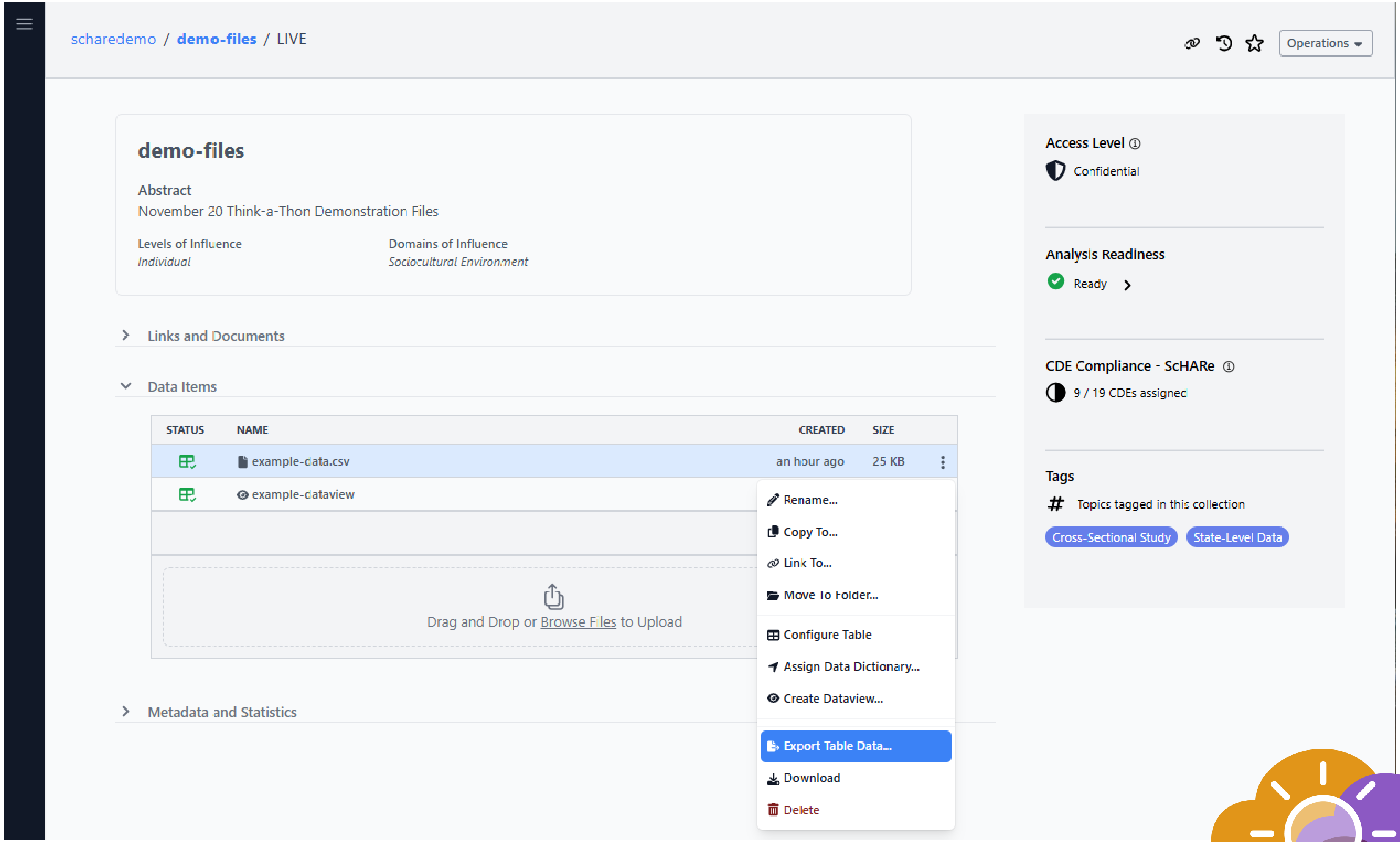
Downloading data to your own computer

Use the item's menu to download the original file

- Best for working directly with raw data

You can also export and download the tabular data in CSV, TSV, or Parquet format

- Best for working with mapped data, summary data, and other processed data



scharedemo / **demo-files** / LIVE

demo-files

Abstract
November 20 Think-a-Thon Demonstration Files

Levels of Influence: *Individual*

Domains of Influence: *Sociocultural Environment*

Access Level: Confidential

Analysis Readiness: Ready

CDE Compliance - ScHARe: 9 / 19 CDEs assigned

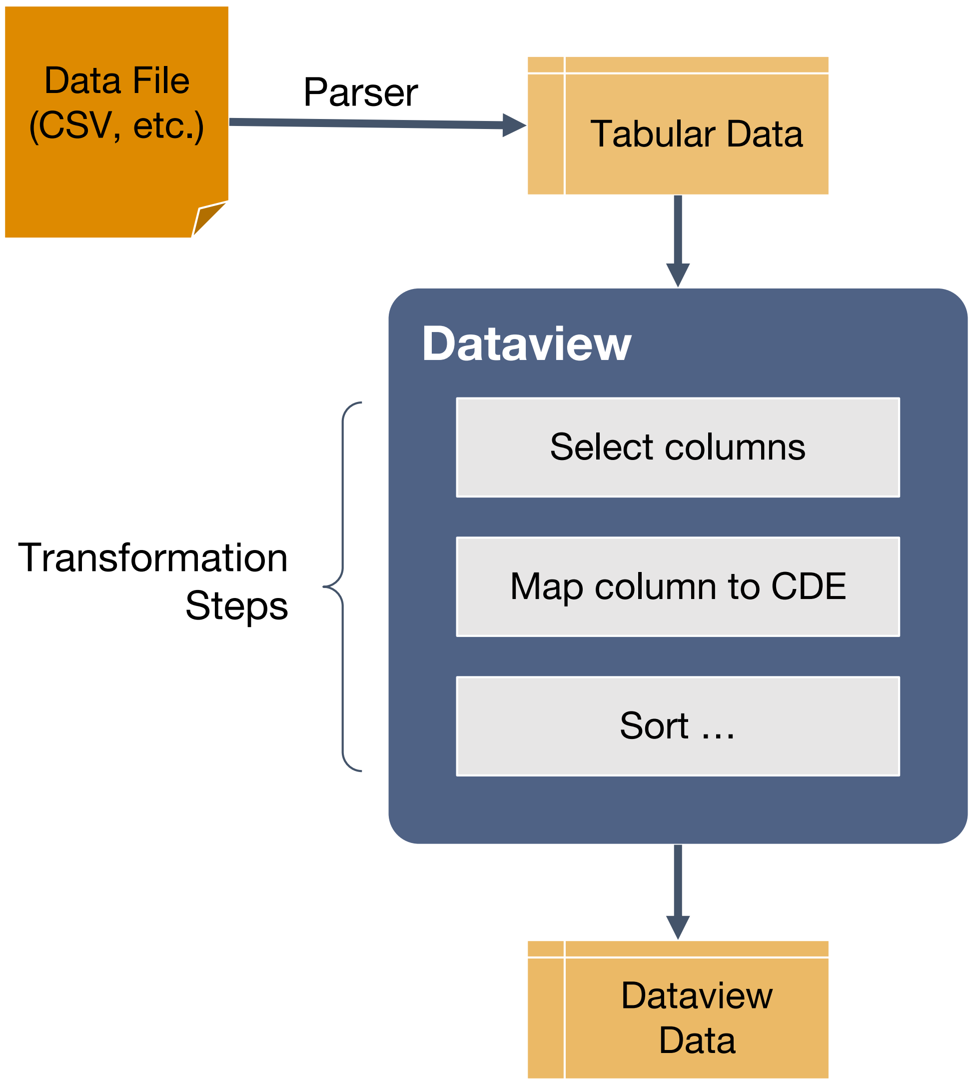
Tags: Topics tagged in this collection
Cross-Sectional Study, State-Level Data

Data Items

STATUS	NAME	CREATED	SIZE
	example-data.csv	an hour ago	25 KB
	example-dataview		

Drag and Drop or Browse Files to Upload

Using Dataviews to Aggregate Data

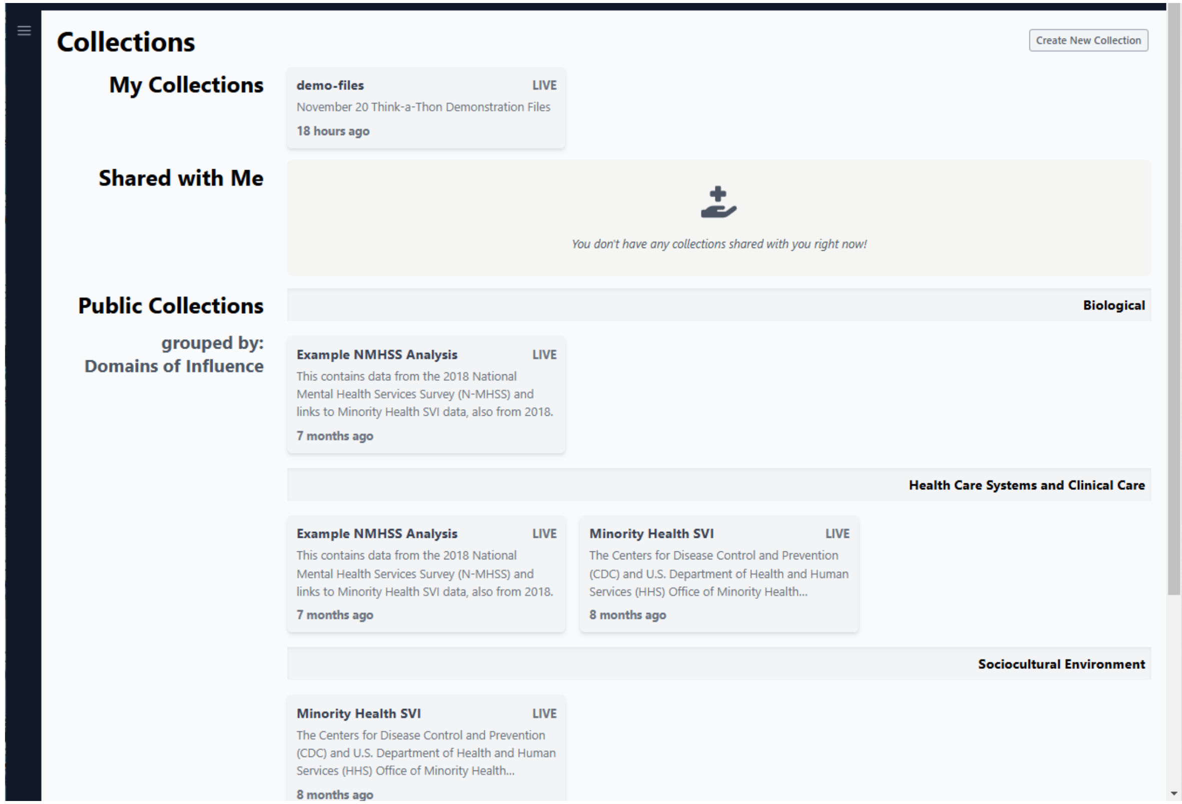


Dataviews take data from one or more sources,
apply a series of transformation steps to that data
(filtering, sorting, mapping, etc.)
resulting in a new table of data as output.

Uses of Dataviews:

- Creating subsets of data
- Hiding PHI/PII for publishing
- Summarizing individual-level data into subsets and estimates
- **Joining multiple datasets together**
- Mapping to CDEs
- … many others!

Browsing and Searching for Data



Collections

My Collections

demo-files LIVE
November 20 Think-a-Thon Demonstration Files
18 hours ago

Shared with Me

You don't have any collections shared with you right now!

Public Collections

grouped by: Domains of Influence

Biological

Example NMHSS Analysis LIVE
This contains data from the 2018 National Mental Health Services Survey (N-MHSS) and links to Minority Health SVI data, also from 2018.
7 months ago

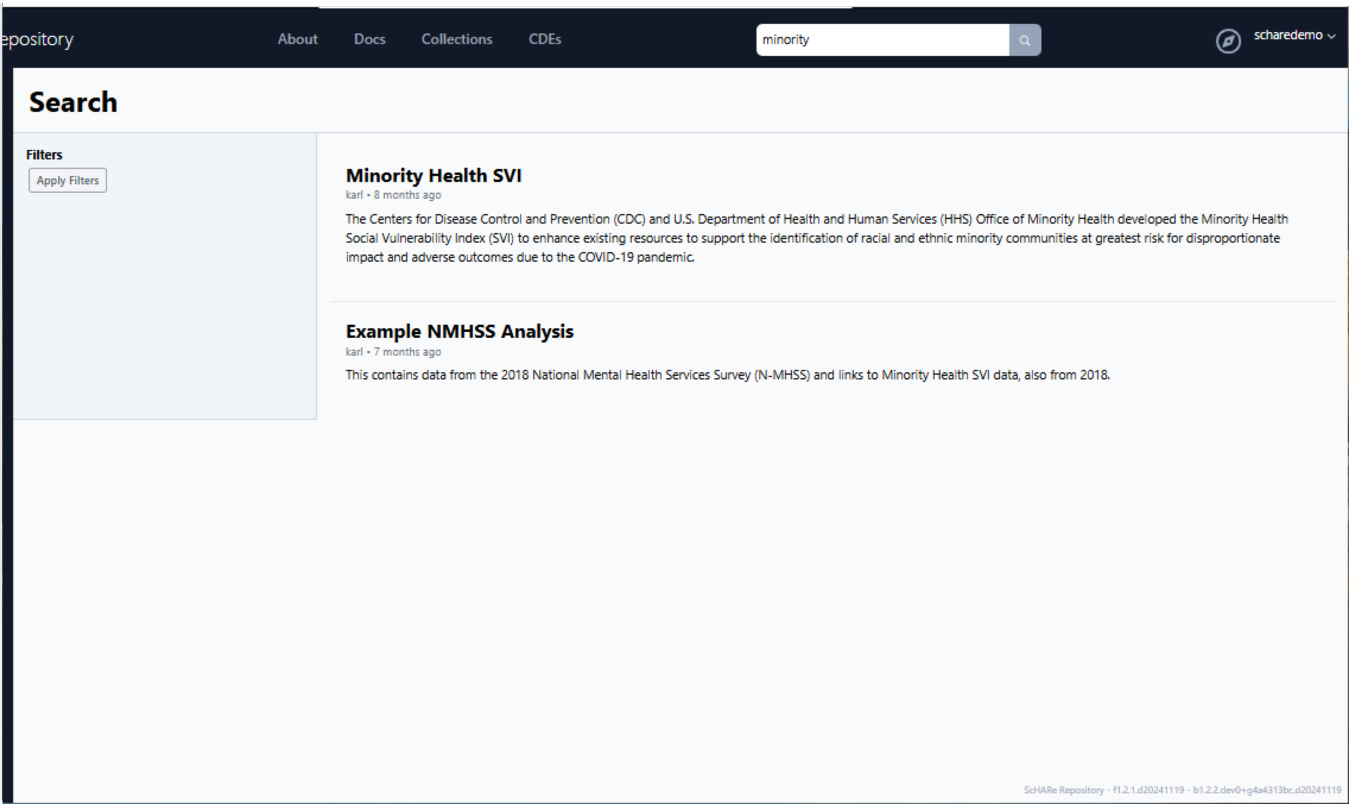
Health Care Systems and Clinical Care

Example NMHSS Analysis LIVE
This contains data from the 2018 National Mental Health Services Survey (N-MHSS) and links to Minority Health SVI data, also from 2018.
7 months ago

Minority Health SVI LIVE
The Centers for Disease Control and Prevention (CDC) and U.S. Department of Health and Human Services (HHS) Office of Minority Health...
8 months ago

Sociocultural Environment

Minority Health SVI LIVE
The Centers for Disease Control and Prevention (CDC) and U.S. Department of Health and Human Services (HHS) Office of Minority Health...
8 months ago



Search: minority

Search

Filters

Minority Health SVI
karl • 8 months ago
The Centers for Disease Control and Prevention (CDC) and U.S. Department of Health and Human Services (HHS) Office of Minority Health developed the Minority Health Social Vulnerability Index (SVI) to enhance existing resources to support the identification of racial and ethnic minority communities at greatest risk for disproportionate impact and adverse outcomes due to the COVID-19 pandemic.

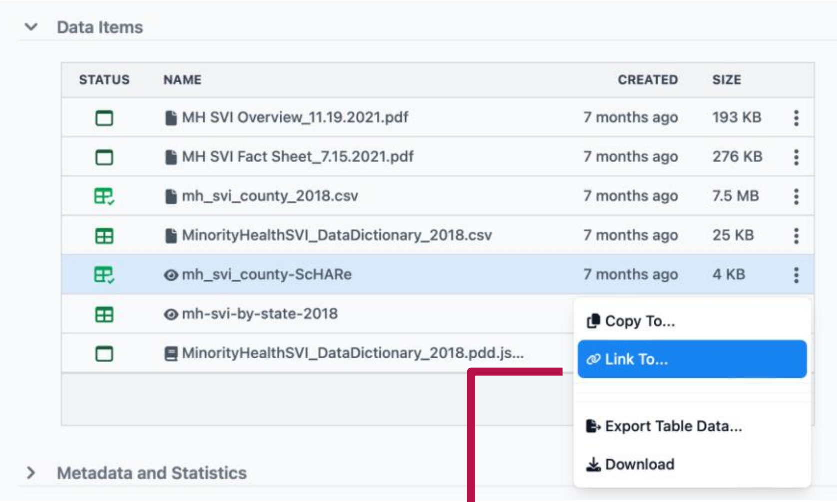
Example NMHSS Analysis
karl • 7 months ago
This contains data from the 2018 National Mental Health Services Survey (N-MHSS) and links to Minority Health SVI data, also from 2018.

ScHARe Repository - f1.2.1.d20241119 - b1.2.2.dev0+g4a4313bc.d20241119

Look for improvements to the data organization and search experience in 2025!

Linking and Aggregating Data

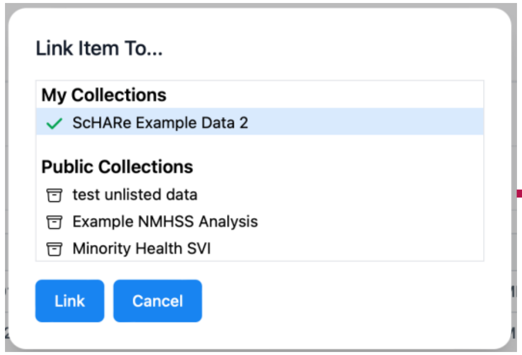
If you find public data that you would like to combine with your project data, you can link it into your own collection and use it just like a file that you had uploaded yourself.



Data Items

STATUS	NAME	CREATED	SIZE
☐	MH SVI Overview_11.19.2021.pdf	7 months ago	193 KB
☐	MH SVI Fact Sheet_7.15.2021.pdf	7 months ago	276 KB
▦	mh_svi_county_2018.csv	7 months ago	7.5 MB
▦	MinorityHealthSVI_DataDictionary_2018.csv	7 months ago	25 KB
▦	mh_svi_county-ScHARe	7 months ago	4 KB
▦	mh-svi-by-state-2018		
☐	MinorityHealthSVI_DataDictionary_2018.pdd.js...		

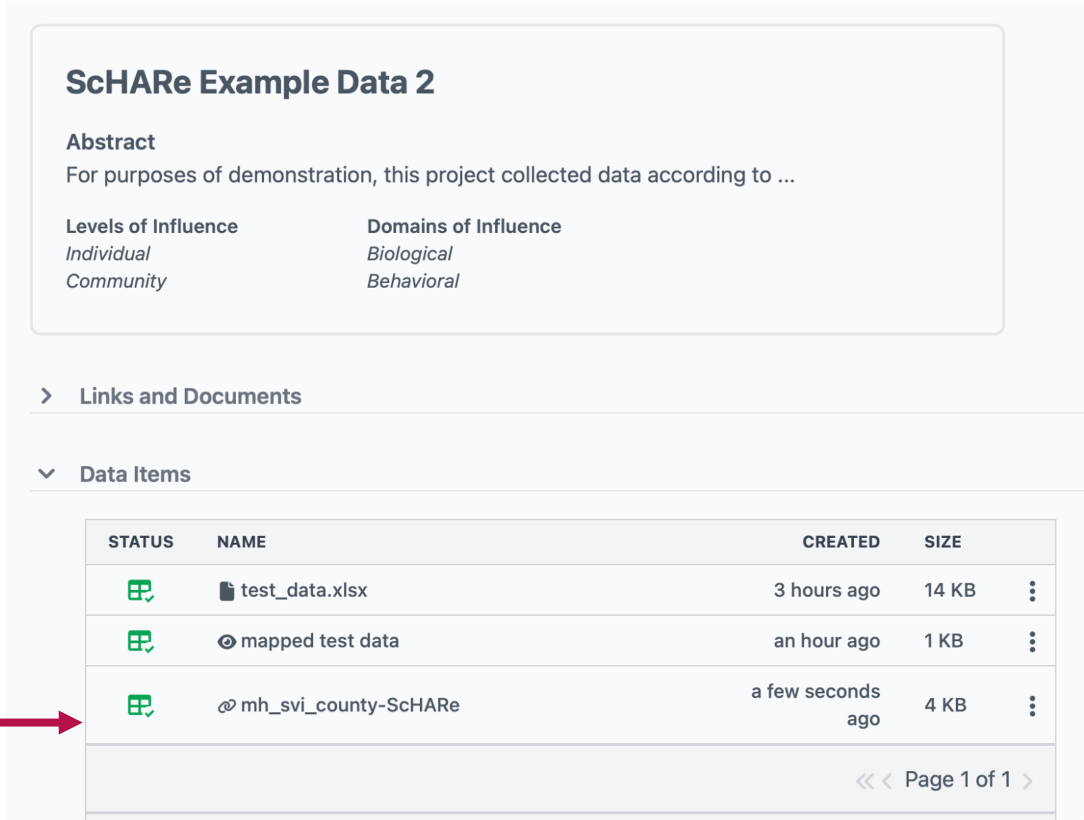
Metadata and Statistics



Link Item To...

My Collections
- ✓ ScHARe Example Data 2

Public Collections
- test unlisted data
- Example NMHSS Analysis
- Minority Health SVI



ScHARe Example Data 2

Abstract

For purposes of demonstration, this project collected data according to ...

Levels of Influence	Domains of Influence
Individual	*Biological*
Community	*Behavioral*

Links and Documents

Data Items

STATUS	NAME	CREATED	SIZE
▦	test_data.xlsx	3 hours ago	14 KB
▦	mapped test data	an hour ago	1 KB
▦	mh_svi_county-ScHARe	a few seconds ago	4 KB

Page 1 of 1

Linking and Aggregating Data

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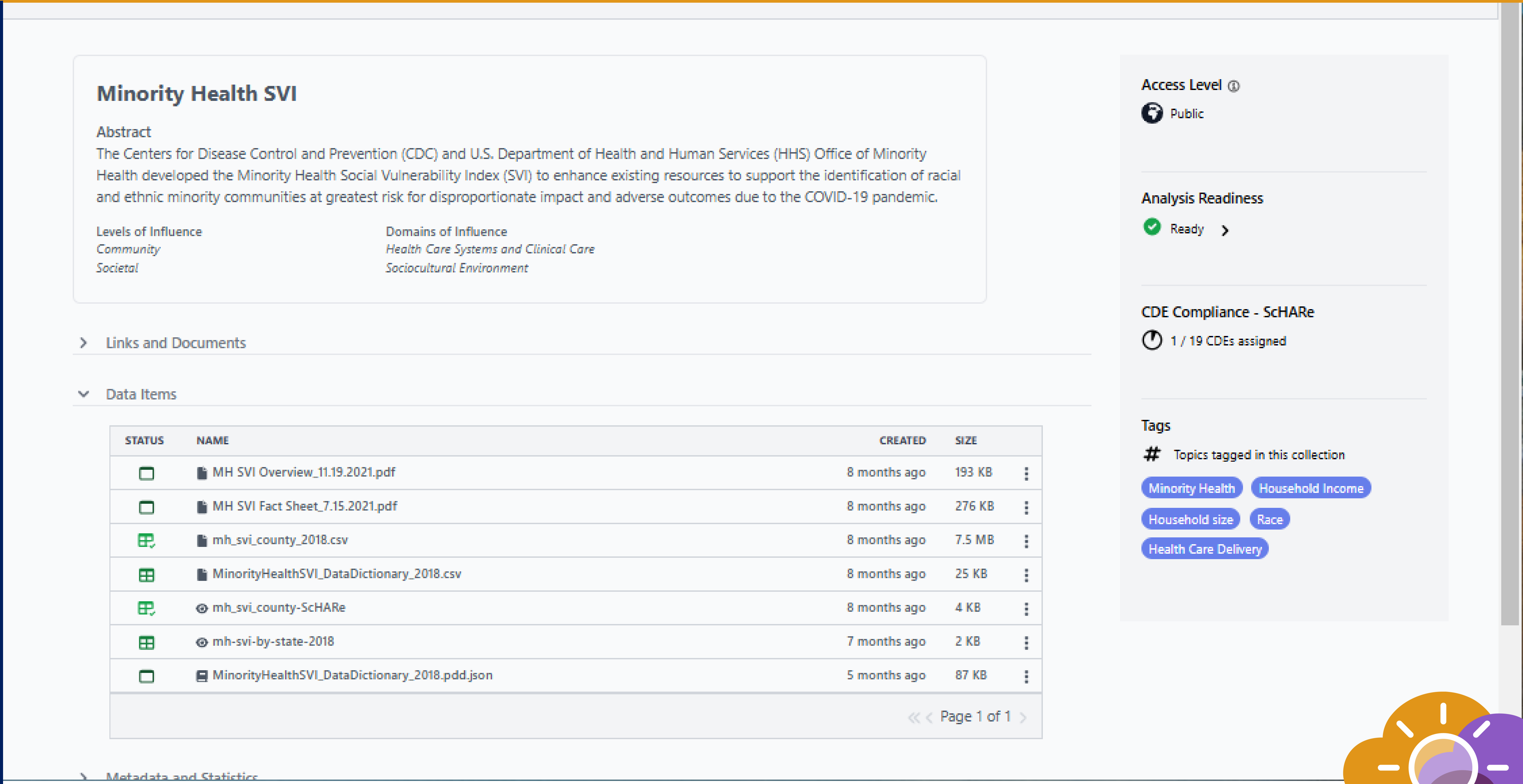


Minority Health SVI

Abstract
The Centers for Disease Control and Prevention (CDC) and U.S. Department of Health and Human Services (HHS) Office of Minority Health developed the Minority Health Social Vulnerability Index (SVI) to enhance existing resources to support the identification of racial and ethnic minority communities at greatest risk for disproportionate impact and adverse outcomes due to the COVID-19 pandemic.

Levels of Influence	Domains of Influence
Community	*Health Care Systems and Clinical Care*
Societal	*Sociocultural Environment*

› Links and Documents

⌄ Data Items

STATUS	NAME	CREATED	SIZE
☐	MH SVI Overview_11.19.2021.pdf	8 months ago	193 KB
☐	MH SVI Fact Sheet_7.15.2021.pdf	8 months ago	276 KB
▦✓	mh_svi_county_2018.csv	8 months ago	7.5 MB
▦	MinorityHealthSVI_DataDictionary_2018.csv	8 months ago	25 KB
▦✓	mh_svi_county-ScHARe	8 months ago	4 KB
▦	mh-svi-by-state-2018	7 months ago	2 KB
☐	MinorityHealthSVI_DataDictionary_2018.pdd.json	5 months ago	87 KB

« ‹ Page 1 of 1 ›

› Metadata and Statistics

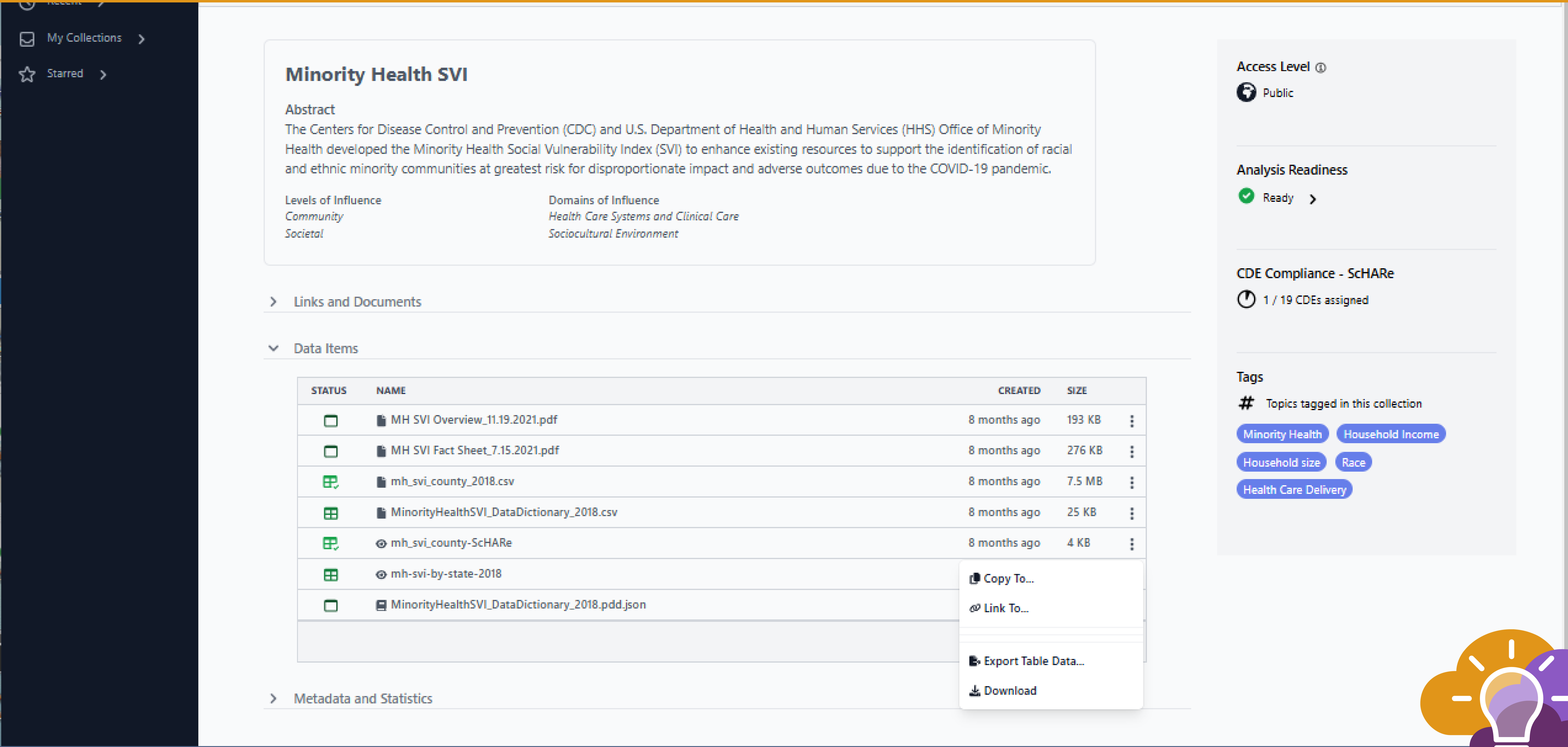
Access Level
Public

Analysis Readiness
✓ Ready ›

CDE Compliance - ScHARe
1 / 19 CDEs assigned

Tags
Topics tagged in this collection
Minority Health
Household Income
Household size
Race
Health Care Delivery

Linking Data



Minority Health SVI

Abstract
The Centers for Disease Control and Prevention (CDC) and U.S. Department of Health and Human Services (HHS) Office of Minority Health developed the Minority Health Social Vulnerability Index (SVI) to enhance existing resources to support the identification of racial and ethnic minority communities at greatest risk for disproportionate impact and adverse outcomes due to the COVID-19 pandemic.

Levels of Influence	Domains of Influence
Community	*Health Care Systems and Clinical Care*
Societal	*Sociocultural Environment*

STATUS	NAME	CREATED	SIZE
▭	MH SVI Overview_11.19.2021.pdf	8 months ago	193 KB
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▦	mh_svi_county-ScHARe	8 months ago	4 KB
▦	mh-svi-by-state-2018		
▭	MinorityHealthSVI_DataDictionary_2018.pdd.json		

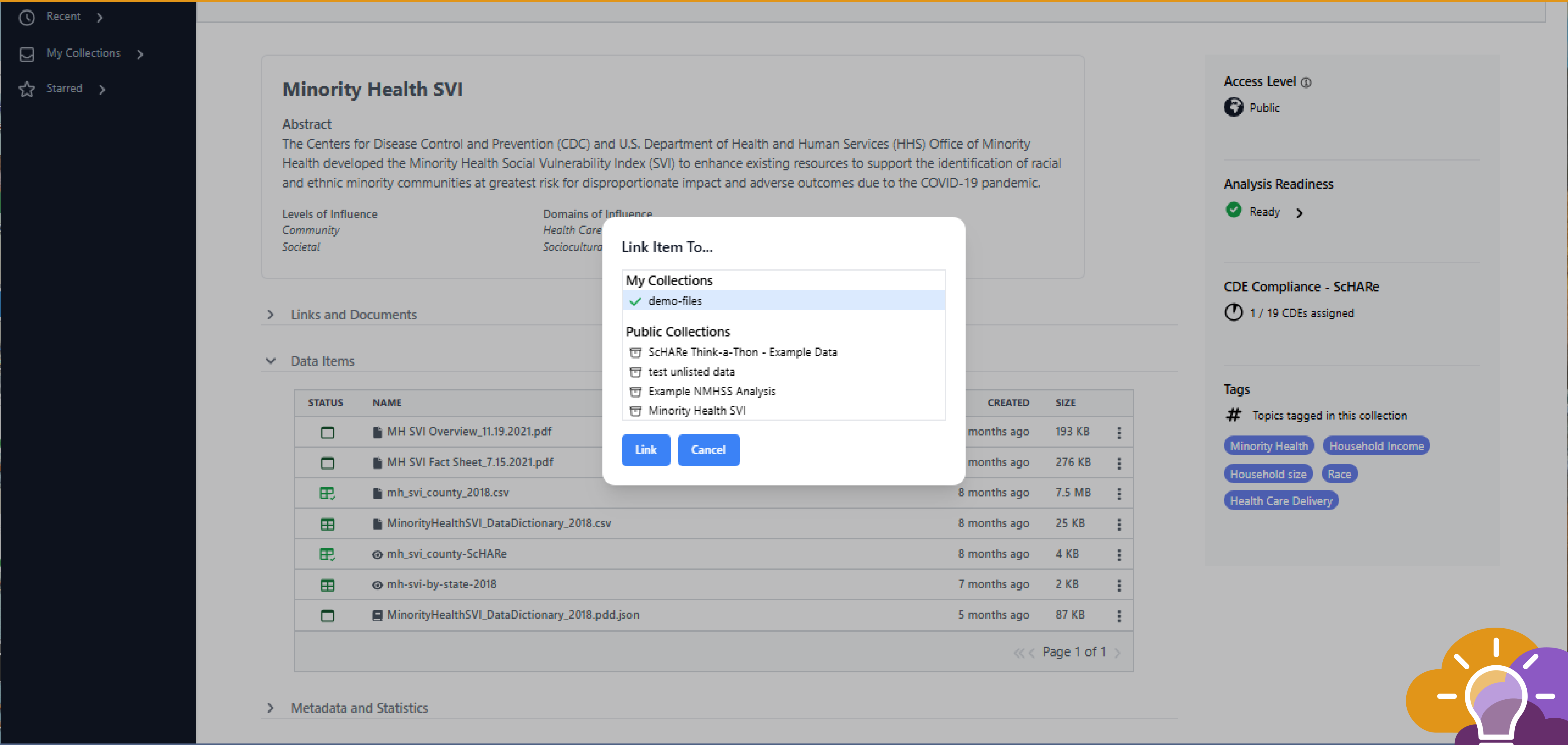
Access Level: Public

Analysis Readiness: Ready

CDE Compliance - ScHARe: 1 / 19 CDEs assigned

Tags
Topics tagged in this collection
Minority Health, Household Income, Household size, Race, Health Care Delivery

Linking Data



Minority Health SVI

Abstract

The Centers for Disease Control and Prevention (CDC) and U.S. Department of Health and Human Services (HHS) Office of Minority Health developed the Minority Health Social Vulnerability Index (SVI) to enhance existing resources to support the identification of racial and ethnic minority communities at greatest risk for disproportionate impact and adverse outcomes due to the COVID-19 pandemic.

Levels of Influence
Community
Societal

Domains of Influence
Health Care
Sociocultura

Link Item To...

My Collections
✓ demo-files

Public Collections
ScHARe Think-a-Thon - Example Data
test unlisted data
Example NMHSS Analysis
Minority Health SVI

Links and Documents

Data Items

STATUS	NAME	CREATED	SIZE
	MH SVI Overview_11.19.2021.pdf	months ago	193 KB
	MH SVI Fact Sheet_7.15.2021.pdf	months ago	276 KB
	mh_svi_county_2018.csv	8 months ago	7.5 MB
	MinorityHealthSVI_DataDictionary_2018.csv	8 months ago	25 KB
	mh_svi_county-ScHARe	8 months ago	4 KB
	mh-svi-by-state-2018	7 months ago	2 KB
	MinorityHealthSVI_DataDictionary_2018.pdd.json	5 months ago	87 KB

Page 1 of 1

Metadata and Statistics

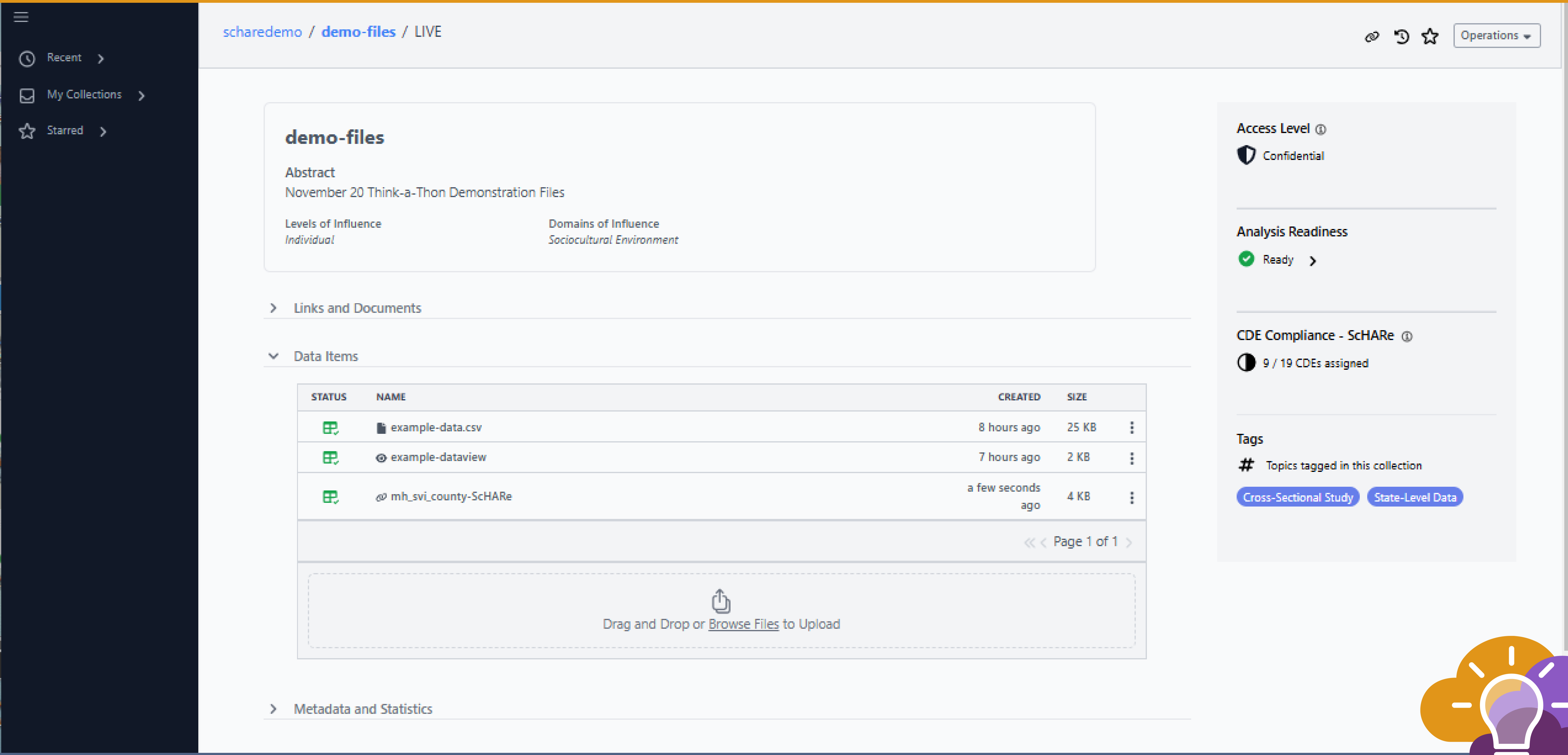
Access Level
Public

Analysis Readiness
Ready

CDE Compliance - ScHARe
1 / 19 CDEs assigned

Tags
Topics tagged in this collection

Linking Data



scharedemo / demo-files / LIVE

demo-files

Abstract
November 20 Think-a-Thon Demonstration Files

Levels of Influence
Individual

Domains of Influence
Sociocultural Environment

Data Items

STATUS	NAME	CREATED	SIZE
✓	example-data.csv	8 hours ago	25 KB
✓	example-dataview	7 hours ago	2 KB
✓	mh_svi_county-ScHARe	a few seconds ago	4 KB

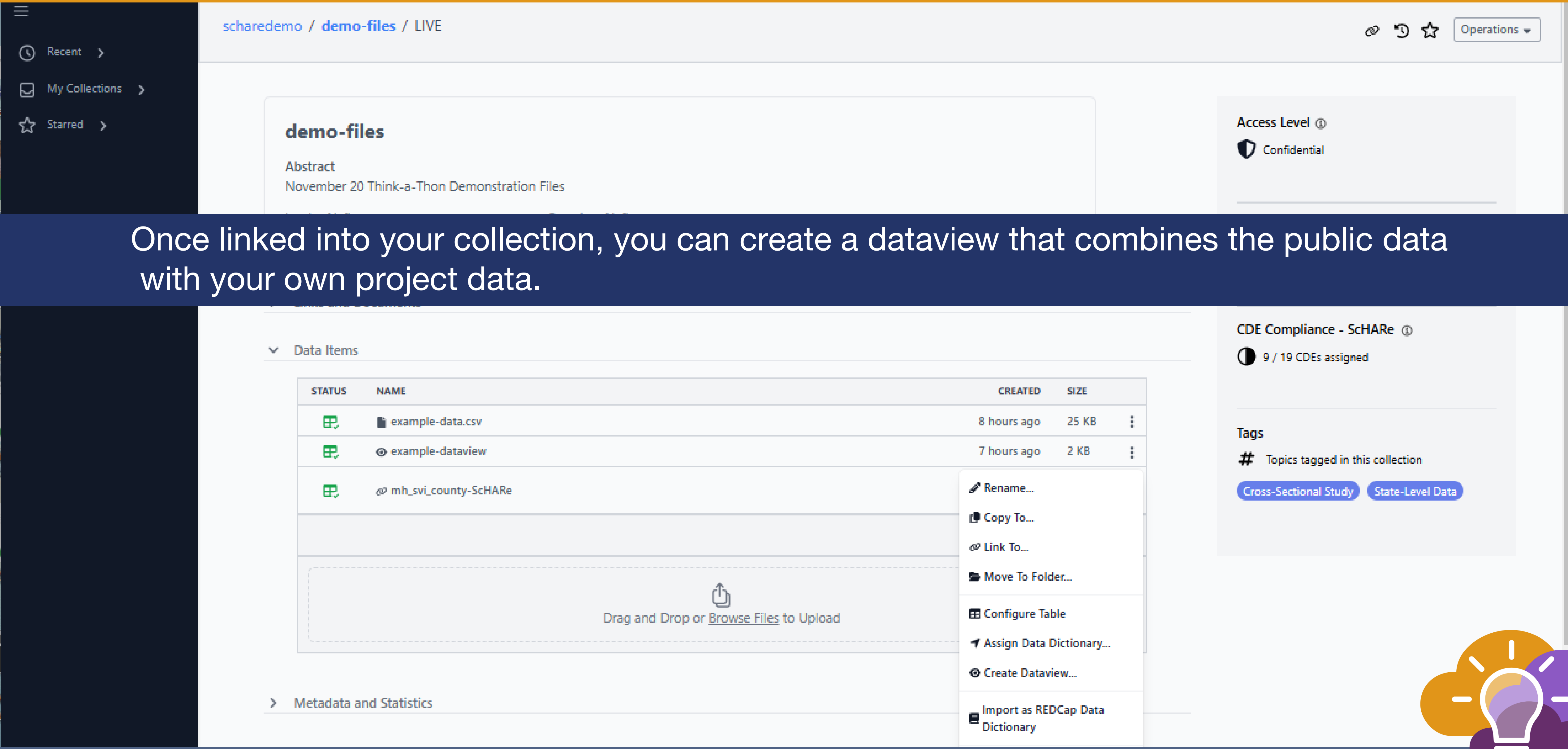
Access Level
Confidential

Analysis Readiness
Ready

CDE Compliance - ScHARe
9 / 19 CDEs assigned

Tags
Topics tagged in this collection
Cross-Sectional Study
State-Level Data

Aggregating Data



demo-files

Abstract

November 20 Think-a-Thon Demonstration Files

Access Level: Confidential

Data Items

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✓	example-data.csv	8 hours ago	25 KB
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✓	mh_svi_county-ScHARe		

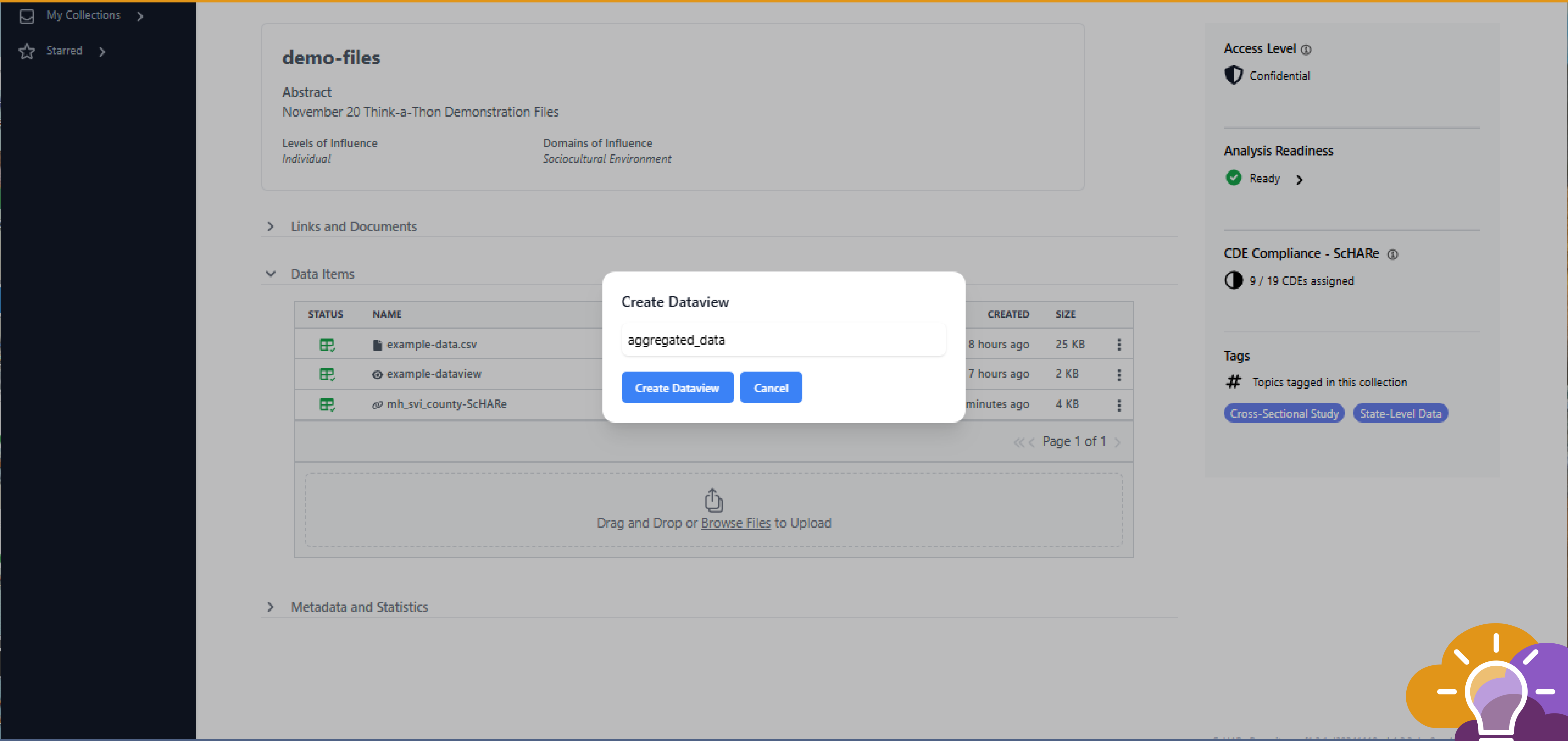
CDE Compliance - ScHARe: 9 / 19 CDEs assigned

Tags: Topics tagged in this collection

Cross-Sectional Study, State-Level Data

Once linked into your collection, you can create a dataview that combines the public data with your own project data.

Aggregating Data



demo-files

Abstract
November 20 Think-a-Thon Demonstration Files

Levels of Influence: *Individual*

Domains of Influence: *Sociocultural Environment*

STATUS	NAME	CREATED	SIZE
	example-data.csv	8 hours ago	25 KB
	example-dataview	7 hours ago	2 KB
	mh_svi_county-ScHARe	minutes ago	4 KB

Page 1 of 1

Drag and Drop or Browse Files to Upload

Create Dataview

aggregated_data

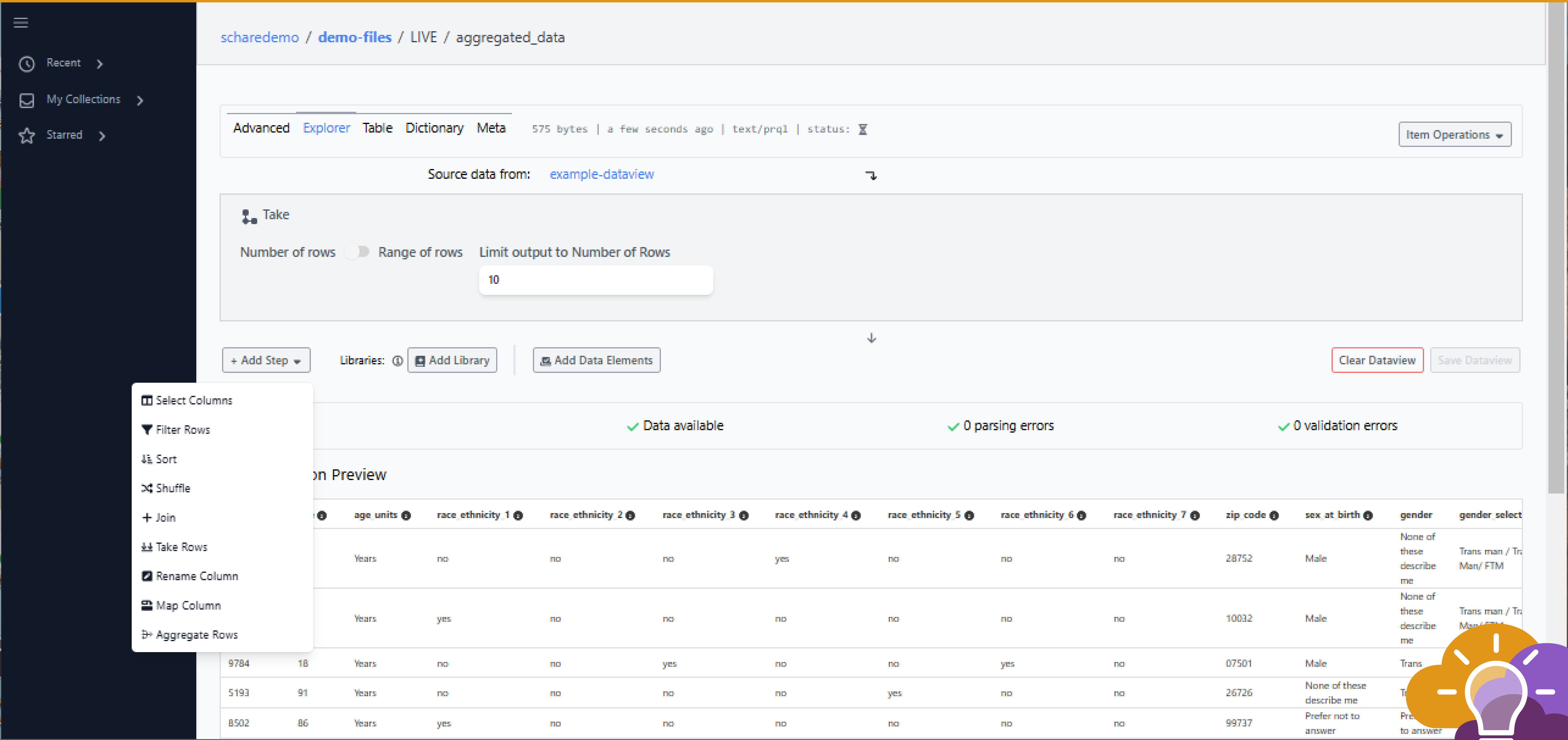
Access Level: Confidential

Analysis Readiness: Ready

CDE Compliance - ScHARe: 9 / 19 CDEs assigned

Tags
Topics tagged in this collection
Cross-Sectional Study
State-Level Data

Aggregating Data



scharedemo / demo-files / LIVE / aggregated_data

575 bytes | a few seconds ago | text/prql | status: ⌛

Source data from: example-dataview

Number of rows ○ Range of rows

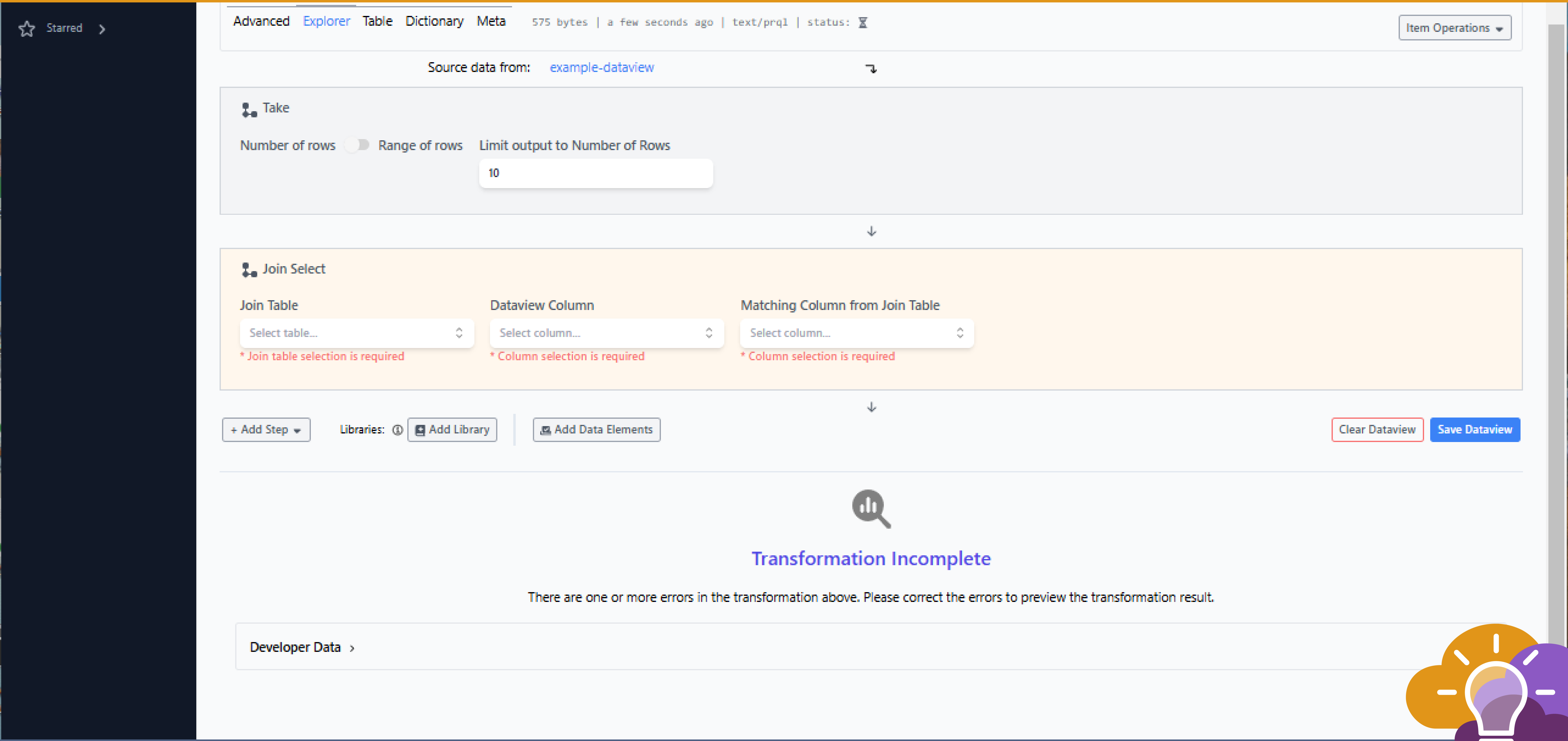
Limit output to Number of Rows: 10

✓ Data available ✓ 0 parsing errors ✓ 0 validation errors

on Preview

		age_units	race_ethnicity_1	race_ethnicity_2	race_ethnicity_3	race_ethnicity_4	race_ethnicity_5	race_ethnicity_6	race_ethnicity_7	zip_code	sex_at_birth	gender	gender_select
		Years	no	no	no	yes	no	no	no	28752	Male	None of these describe me	Trans man / Tr Man/ FTM
		Years	yes	no	no	no	no	no	no	10032	Male	None of these describe me	Trans man / Tr
9784	18	Years	no	no	yes	no	no	yes	no	07501	Male	Trans	
5193	91	Years	no	no	no	no	yes	no	no	26726	None of these describe me	T	
8502	86	Years	yes	no	no	no	no	no	no	99737	Prefer not to answer	Pre to answer	

Aggregating Data



575 bytes | a few seconds ago | text/prql | status:

Source data from: example-dataview

Take

Number of rows ○ Range of rows

Limit output to Number of Rows: 10

Join Select

Join Table: Select table...
* Join table selection is required

Dataview Column: Select column...
* Column selection is required

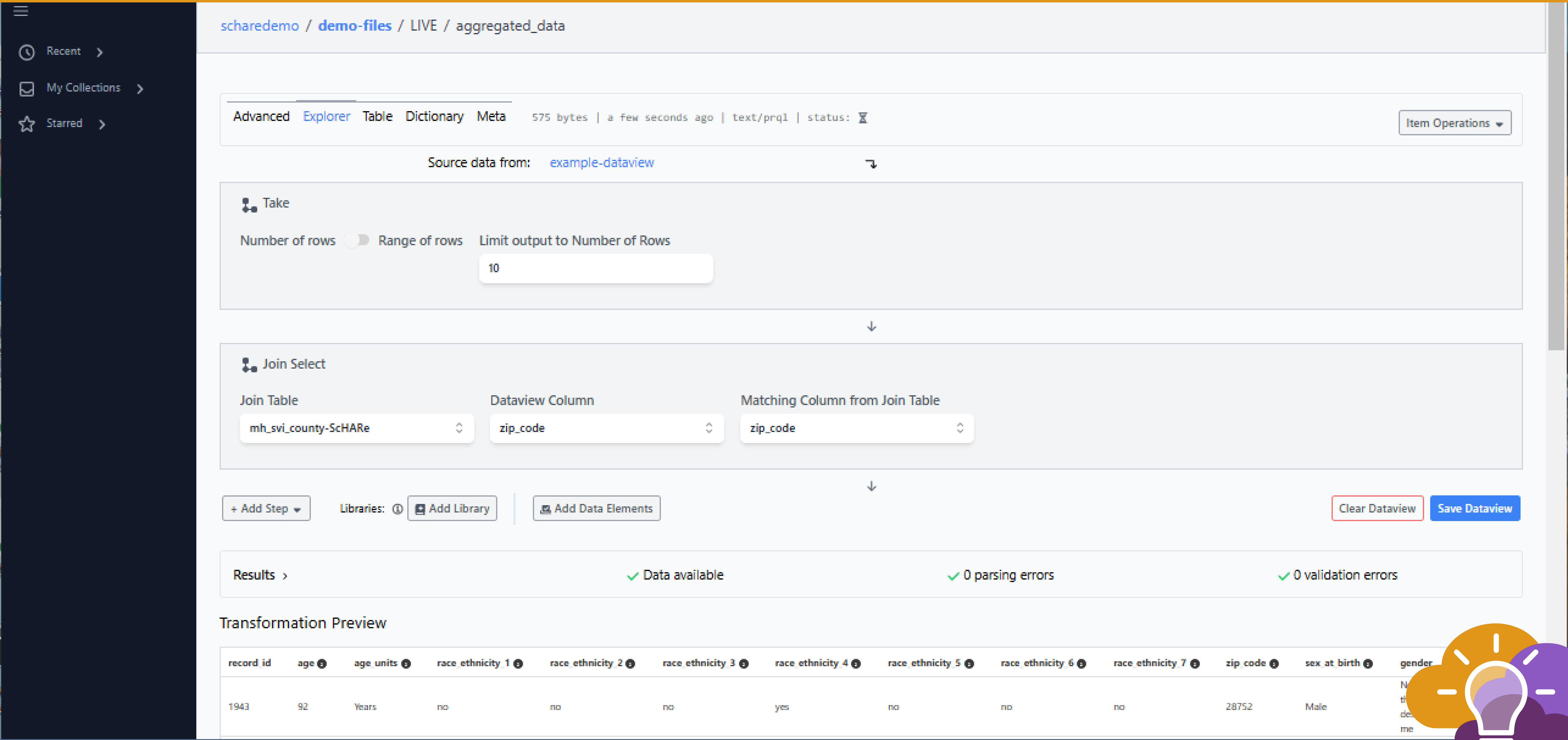
Matching Column from Join Table: Select column...
* Column selection is required

Libraries:

Transformation Incomplete

There are one or more errors in the transformation above. Please correct the errors to preview the transformation result.

Aggregating Data



scharedemo / demo-files / LIVE / aggregated_data

575 bytes | a few seconds ago | text/prql | status: ⏳

Source data from: example-dataview

Take

Number of rows ○ Range of rows

Limit output to Number of Rows: 10

Join Select

Join Table: mh_svi_county-ScHARe

Dataview Column: zip_code

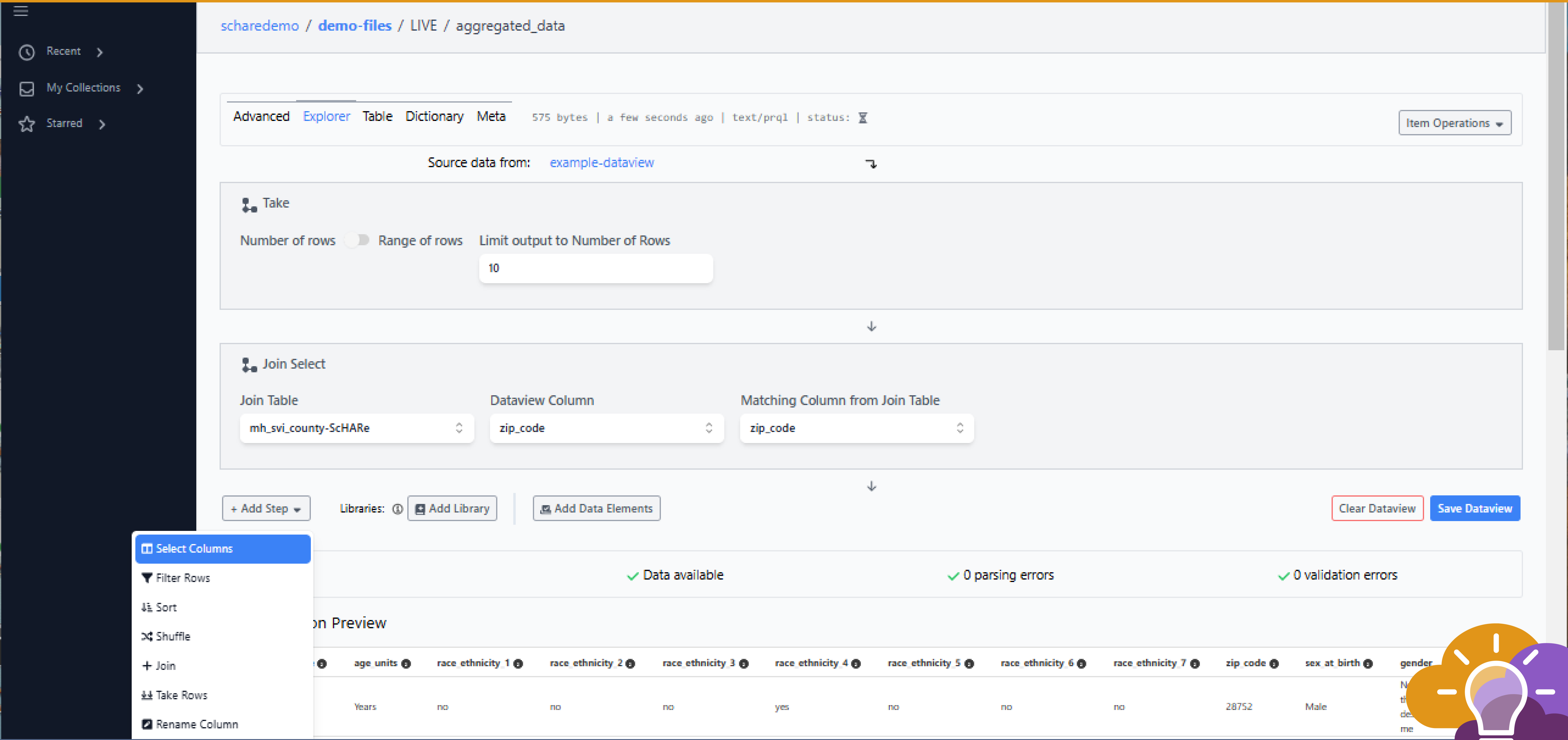
Matching Column from Join Table: zip_code

Results › ✓ Data available ✓ 0 parsing errors ✓ 0 validation errors

Transformation Preview

record_id	age	age_units	race_ethnicity_1	race_ethnicity_2	race_ethnicity_3	race_ethnicity_4	race_ethnicity_5	race_ethnicity_6	race_ethnicity_7	zip_code	sex_at_birth	gender
1943	92	Years	no	no	no	yes	no	no	no	28752	Male	N th de me

Aggregating Data



scharedemo / demo-files / LIVE / aggregated_data

575 bytes | a few seconds ago | text/prql | status: ⧗

Source data from: example-dataview

Take

Number of rows ○ Range of rows

Limit output to Number of Rows: 10

Join Select

Join Table: mh_svi_county-ScHARe

Dataview Column: zip_code

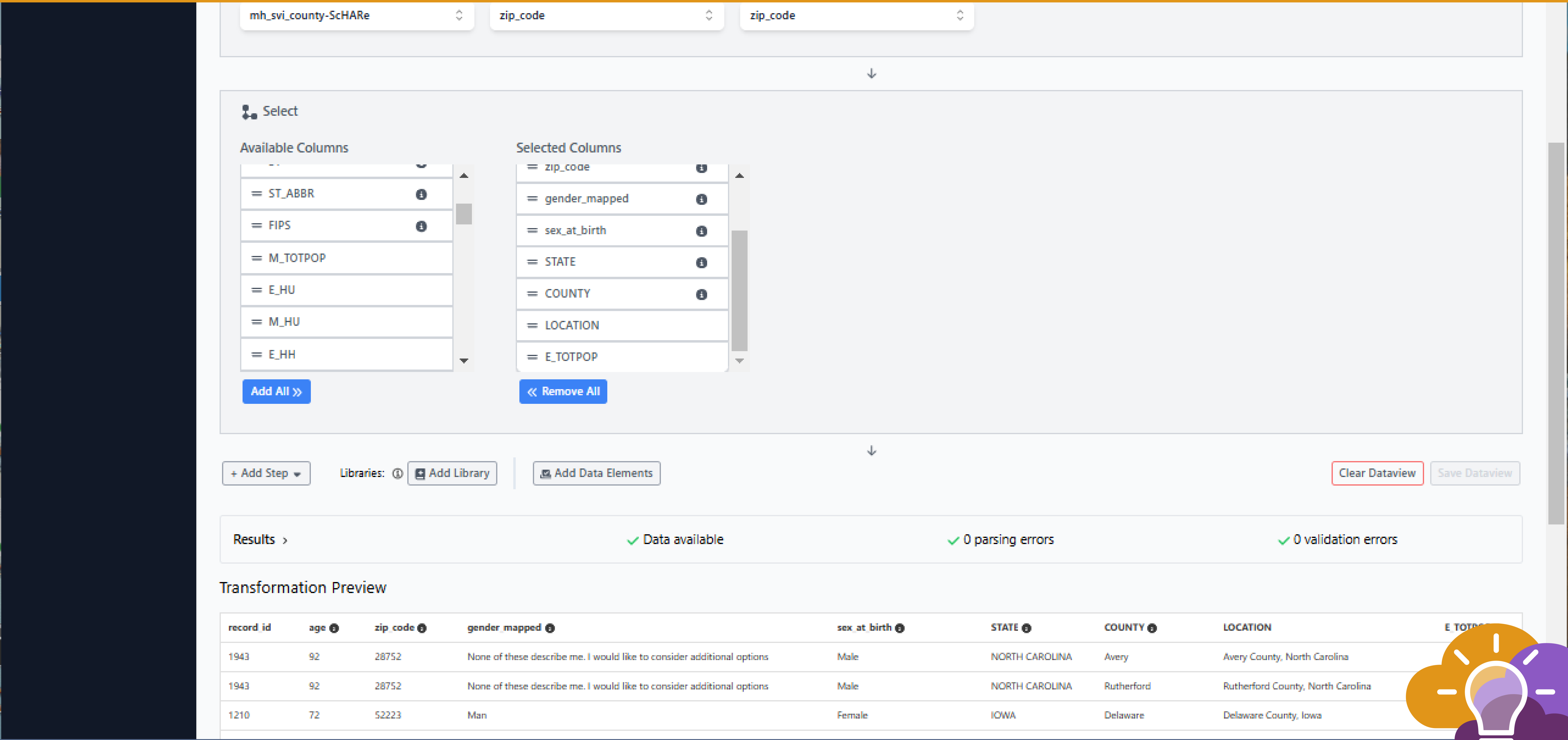
Matching Column from Join Table: zip_code

✓ Data available ✓ 0 parsing errors ✓ 0 validation errors

on Preview

age_units	race_ethnicity_1	race_ethnicity_2	race_ethnicity_3	race_ethnicity_4	race_ethnicity_5	race_ethnicity_6	race_ethnicity_7	zip_code	sex_at_birth	gender
Years	no	no	no	yes	no	no	no	28752	Male	

Selecting Variables



✓ Data available

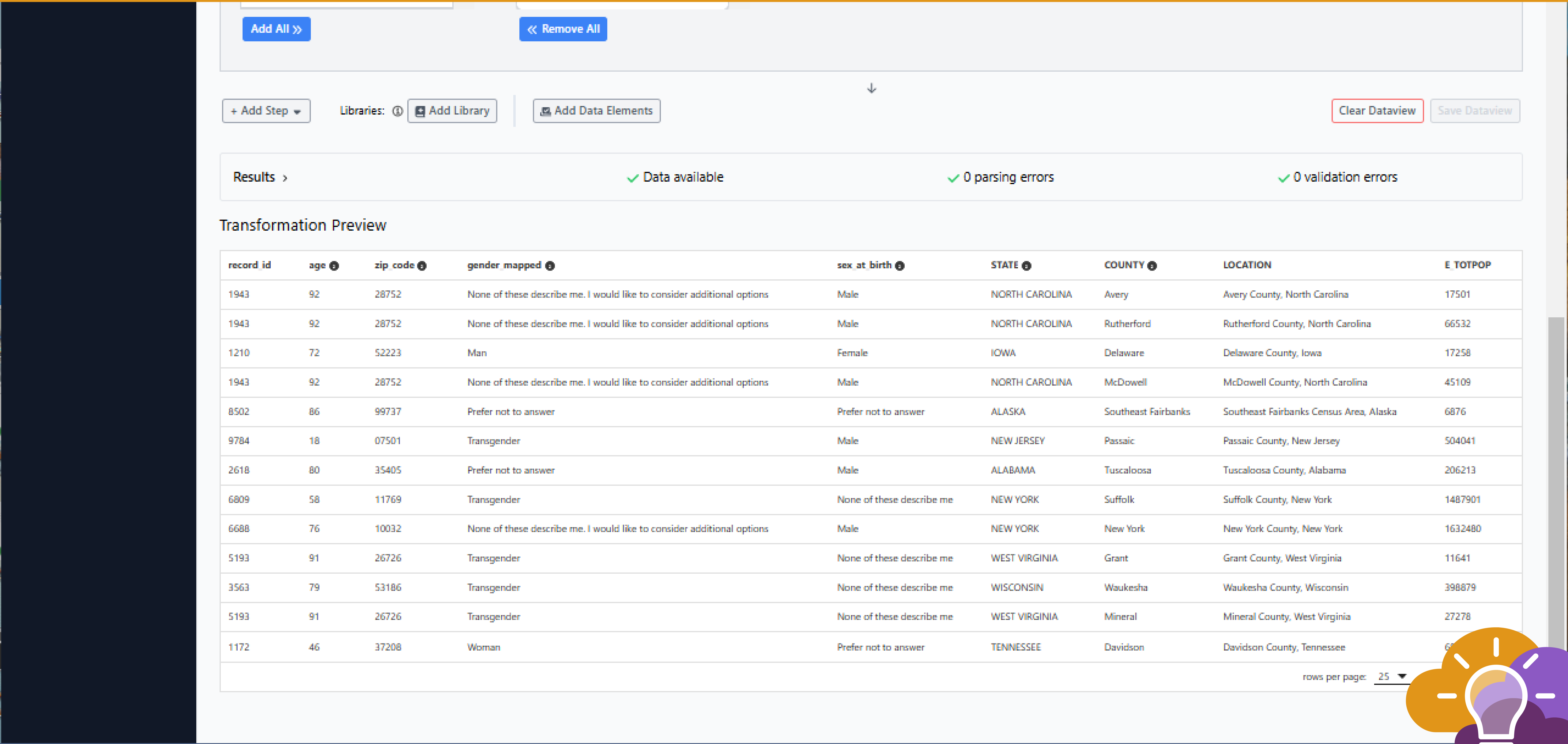
✓ 0 parsing errors

✓ 0 validation errors

Transformation Preview

record_id	age	zip_code	gender_mapped	sex_at_birth	STATE	COUNTY	LOCATION	E_TOTP
1943	92	28752	None of these describe me. I would like to consider additional options	Male	NORTH CAROLINA	Avery	Avery County, North Carolina	
1943	92	28752	None of these describe me. I would like to consider additional options	Male	NORTH CAROLINA	Rutherford	Rutherford County, North Carolina	
1210	72	52223	Man	Female	IOWA	Delaware	Delaware County, Iowa	

Selecting Variables



Results › ✓ Data available ✓ 0 parsing errors ✓ 0 validation errors

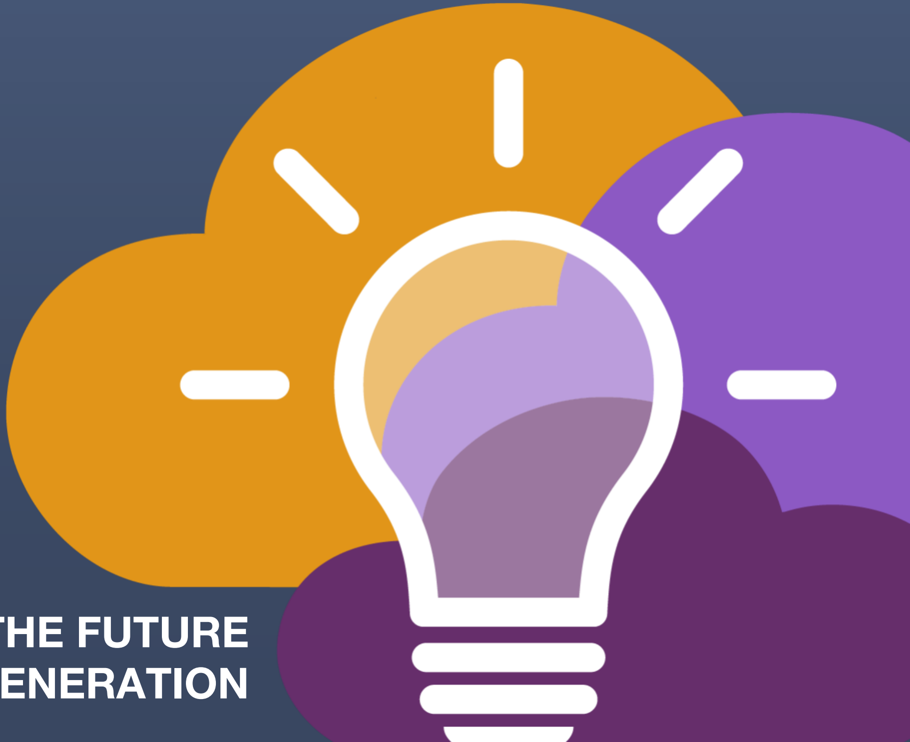
Transformation Preview

record_id	age	zip_code	gender_mapped	sex_at_birth	STATE	COUNTY	LOCATION	E_TOTPOP
1943	92	28752	None of these describe me. I would like to consider additional options	Male	NORTH CAROLINA	Avery	Avery County, North Carolina	17501
1943	92	28752	None of these describe me. I would like to consider additional options	Male	NORTH CAROLINA	Rutherford	Rutherford County, North Carolina	66532
1210	72	52223	Man	Female	IOWA	Delaware	Delaware County, Iowa	17258
1943	92	28752	None of these describe me. I would like to consider additional options	Male	NORTH CAROLINA	McDowell	McDowell County, North Carolina	45109
8502	86	99737	Prefer not to answer	Prefer not to answer	ALASKA	Southeast Fairbanks	Southeast Fairbanks Census Area, Alaska	6876
9784	18	07501	Transgender	Male	NEW JERSEY	Passaic	Passaic County, New Jersey	504041
2618	80	35405	Prefer not to answer	Male	ALABAMA	Tuscaloosa	Tuscaloosa County, Alabama	206213
6809	58	11769	Transgender	None of these describe me	NEW YORK	Suffolk	Suffolk County, New York	1487901
6688	76	10032	None of these describe me. I would like to consider additional options	Male	NEW YORK	New York	New York County, New York	1632480
5193	91	26726	Transgender	None of these describe me	WEST VIRGINIA	Grant	Grant County, West Virginia	11641
3563	79	53186	Transgender	None of these describe me	WISCONSIN	Waukesha	Waukesha County, Wisconsin	398879
5193	91	26726	Transgender	None of these describe me	WEST VIRGINIA	Mineral	Mineral County, West Virginia	27278
1172	46	37208	Woman	Prefer not to answer	TENNESSEE	Davidson	Davidson County, Tennessee	6

rows per page: 25

ScHARe

Data Analysis



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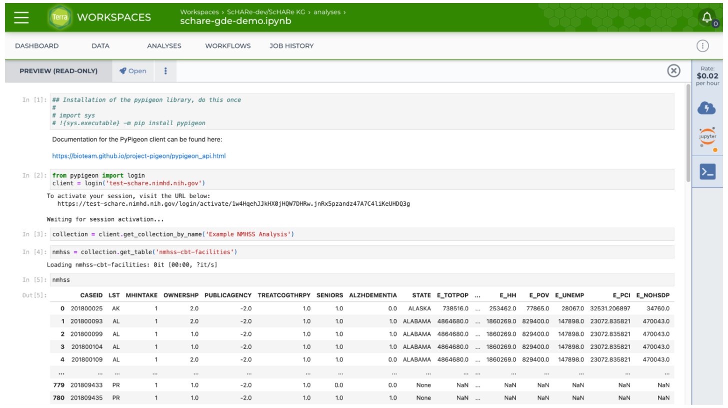
OF KNOWLEDGE GENERATION

How do I analyze or aggregate data from the ScHARe Repository?

Three quick options:

Connect to a Terra Jupyter notebook



```
In [1]: ## Installation of the pypigeon library, do this once
        #
        # import sys
        # !{sys.executable} -m pip install pypigeon
```

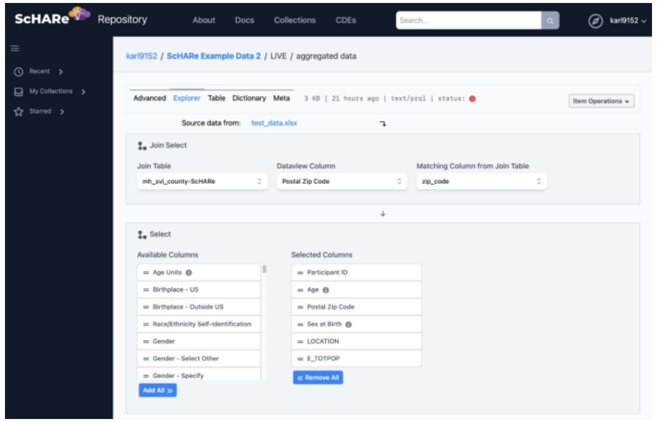
Documentation for the PyPigeon client can be found here:

https://bioteam.github.io/project-pigeon/pypigeon_api.html

```
In [2]: from pypigeon import login
        client = login('test-schare.nimhd.nih.gov')

To activate your session, visit the URL below:
  https://test-schare.nimhd.nih.gov/login/activate/1w4HqehJJkHX0jHQW7DHRw.jnRx5pzandz47A7C4liKeUHDQ3g

Waiting for session activation...

In [3]: collection = client.get_collection_by_name('Example NMHSS Analysis')

In [4]: nmhss = collection.get_table('nmhss-cbt-facilities')
Loading nmhss-cbt-facilities: 0it [00:00, ?it/s]

In [5]: nmhss

Out[5]:
        CASEID LST MHINTAKE OWNERSHP PUBLICAGENCY TREATCOGTHRPY SENIORS ALZHDEMENTIA   STATE  E_TOTPOP ...      E_HH    E_POV  E_UNEMP        E_PCI E_NOHSDP
  0  201800025  AK        1      2.0         -2.0           1.0     1.0          0.0  ALASKA  738516.0 ...  253462.0  77865.0  28067.0 32531.206897  34760.0
  1  201800093  AL        1      2.0         -2.0           1.0     1.0          1.0 ALABAMA 4864680.0 ... 1860269.0 829400.0 147898.0 23072.835821 470043.0
  2  201800099  AL        1      1.0         -2.0           1.0     1.0          1.0 ALABAMA 4864680.0 ... 1860269.0 829400.0 147898.0 23072.835821 470043.0
  3  201800104  AL        1      1.0         -2.0           1.0     1.0          1.0 ALABAMA 4864680.0 ... 1860269.0 829400.0 147898.0 23072.835821 470043.0
  4  201800109  AL        1      2.0         -2.0           1.0     1.0          0.0 ALABAMA 4864680.0 ... 1860269.0 829400.0 147898.0 23072.835821 470043.0
...        ...  ...     ...      ...          ...           ...     ...          ...     ...       ... ...       ...      ...      ...          ...      ...
779  201809433  PR        1      1.0         -2.0           1.0     0.0          0.0    None       NaN ...       NaN      NaN      NaN          NaN      NaN
780  201809435  PR        1      1.0         -2.0           1.0     1.0          1.0    None       NaN ...       NaN      NaN      NaN          NaN      NaN
```

Download to your own computer

Use Dataviews on the Repository



karl9152 / ScHARe Example Data 2 / LIVE / aggregated data

3 KB | 21 hours ago | text/prql | status:

Source data from: test_data.xlsx

Join Table: mh_svi_county-ScHARe

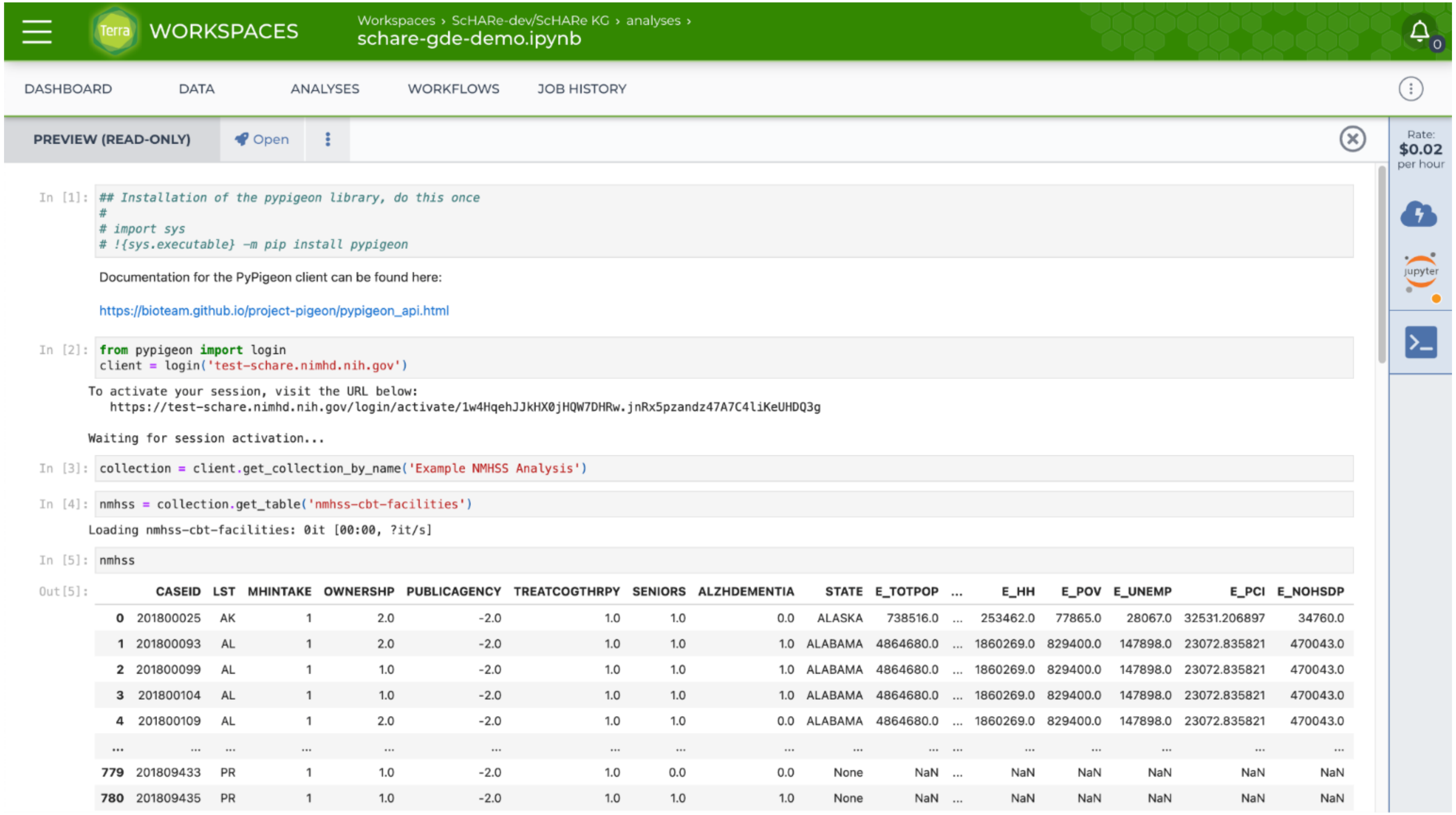
Dataview Column: Postal Zip Code

Matching Column from Join Table: zip_code

Available Columns	Selected Columns
Age Units	Participant ID
Birthplace - US	Age
Birthplace - Outside US	Postal Zip Code
Race/Ethnicity Self-Identification	Sex at Birth
Gender	LOCATION
Gender - Select Other	E_TOTPOP
Gender - Specify	

Look for a future Think-a-Thon where we will share more about how to link your data to Terra and how to use the ScHARe Repository to create aggregated data sets!

Analyzing Data on Terra (Jupyter Notebooks)



In [1]:

```
## Installation of the pypigeon library, do this once
#
# import sys
# !{sys.executable} -m pip install pypigeon
```

Documentation for the PyPigeon client can be found here:

https://bioteam.github.io/project-pigeon/pypigeon_api.html

In [2]:

```
from pypigeon import login
client = login('test-schare.nimhd.nih.gov')
```

```
To activate your session, visit the URL below:
  https://test-schare.nimhd.nih.gov/login/activate/1w4HqehJJkHX0jHQW7DHRw.jnRx5pzandz47A7C4liKeUHDQ3g

Waiting for session activation...
```

In [3]:

```
collection = client.get_collection_by_name('Example NMHSS Analysis')
```

In [4]:

```
nmhss = collection.get_table('nmhss-cbt-facilities')
```

```
Loading nmhss-cbt-facilities: 0it [00:00, ?it/s]
```

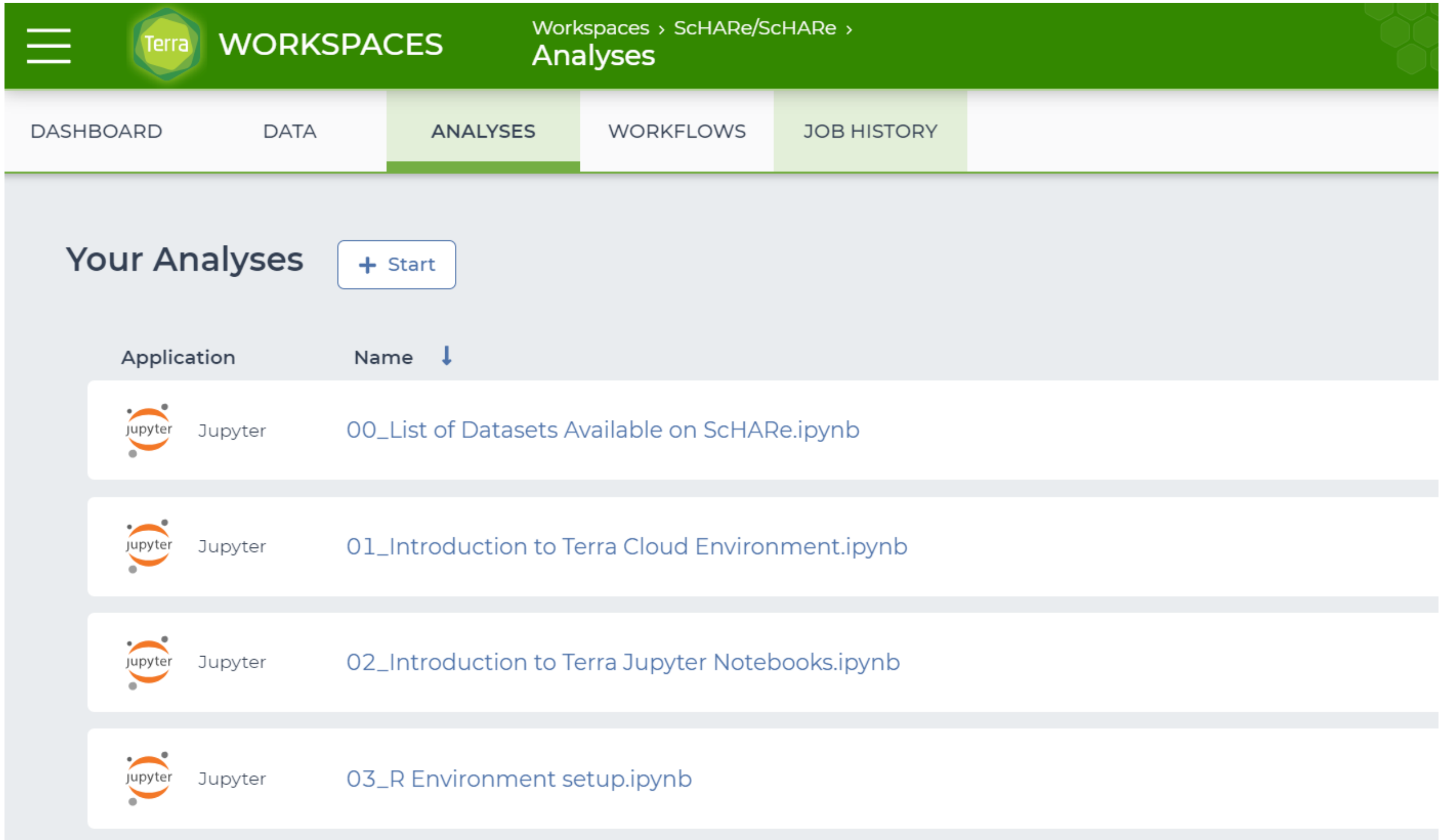
In [5]:

```
nmhss
```

Out[5]:

	CASEID	LST	MHINTAKE	OWNERSHP	PUBLICAGENCY	TREATCOGTHRPY	SENIORS	ALZHDEMENTIA	STATE	E_TOTPOP	...	E_HH	E_POV	E_UNEMP	E_PCI	E_NOHSDP
0	201800025	AK	1	2.0	-2.0	1.0	1.0	0.0	ALASKA	738516.0	...	253462.0	77865.0	28067.0	32531.206897	34760.0
1	201800093	AL	1	2.0	-2.0	1.0	1.0	1.0	ALABAMA	4864680.0	...	1860269.0	829400.0	147898.0	23072.835821	470043.0
2	201800099	AL	1	1.0	-2.0	1.0	1.0	1.0	ALABAMA	4864680.0	...	1860269.0	829400.0	147898.0	23072.835821	470043.0
3	201800104	AL	1	1.0	-2.0	1.0	1.0	1.0	ALABAMA	4864680.0	...	1860269.0	829400.0	147898.0	23072.835821	470043.0
4	201800109	AL	1	2.0	-2.0	1.0	1.0	0.0	ALABAMA	4864680.0	...	1860269.0	829400.0	147898.0	23072.835821	470043.0
...	...	...	...	...	...	...	...	...	...	...	...	...	...	...	...	...
779	201809433	PR	1	1.0	-2.0	1.0	0.0	0.0	None	NaN	...	NaN	NaN	NaN	NaN	NaN
780	201809435	PR	1	1.0	-2.0	1.0	1.0	1.0	None	NaN	...	NaN	NaN	NaN	NaN	NaN

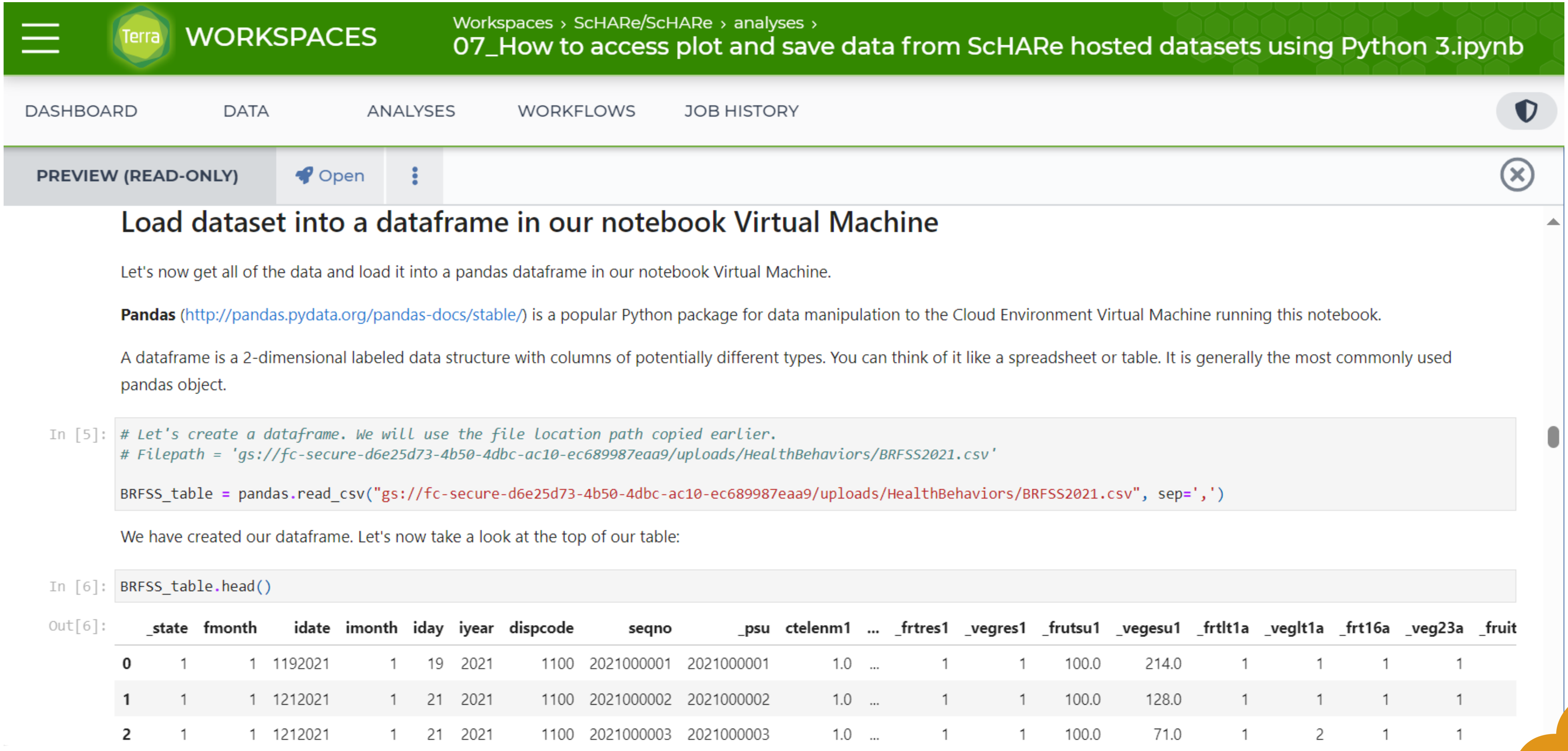
Analyzing Data on Terra (Jupyter Notebooks)



Your Analyses

Application	Name ↓
Jupyter	00_List of Datasets Available on ScHARe.ipynb
Jupyter	01_Introduction to Terra Cloud Environment.ipynb
Jupyter	02_Introduction to Terra Jupyter Notebooks.ipynb
Jupyter	03_R Environment setup.ipynb

Analyzing Data on Terra (Jupyter Notebooks)



Load dataset into a dataframe in our notebook Virtual Machine

Let's now get all of the data and load it into a pandas dataframe in our notebook Virtual Machine.

Pandas (http://pandas.pydata.org/pandas-docs/stable/) is a popular Python package for data manipulation to the Cloud Environment Virtual Machine running this notebook.

A dataframe is a 2-dimensional labeled data structure with columns of potentially different types. You can think of it like a spreadsheet or table. It is generally the most commonly used pandas object.

In [5]:

```
# Let's create a dataframe. We will use the file location path copied earlier.
# Filepath = 'gs://fc-secure-d6e25d73-4b50-4dbc-ac10-ec689987eaa9/uploads/HealthBehaviors/BRFSS2021.csv'

BRFSS_table = pandas.read_csv("gs://fc-secure-d6e25d73-4b50-4dbc-ac10-ec689987eaa9/uploads/HealthBehaviors/BRFSS2021.csv", sep=',')
```

We have created our dataframe. Let's now take a look at the top of our table:

In [6]:

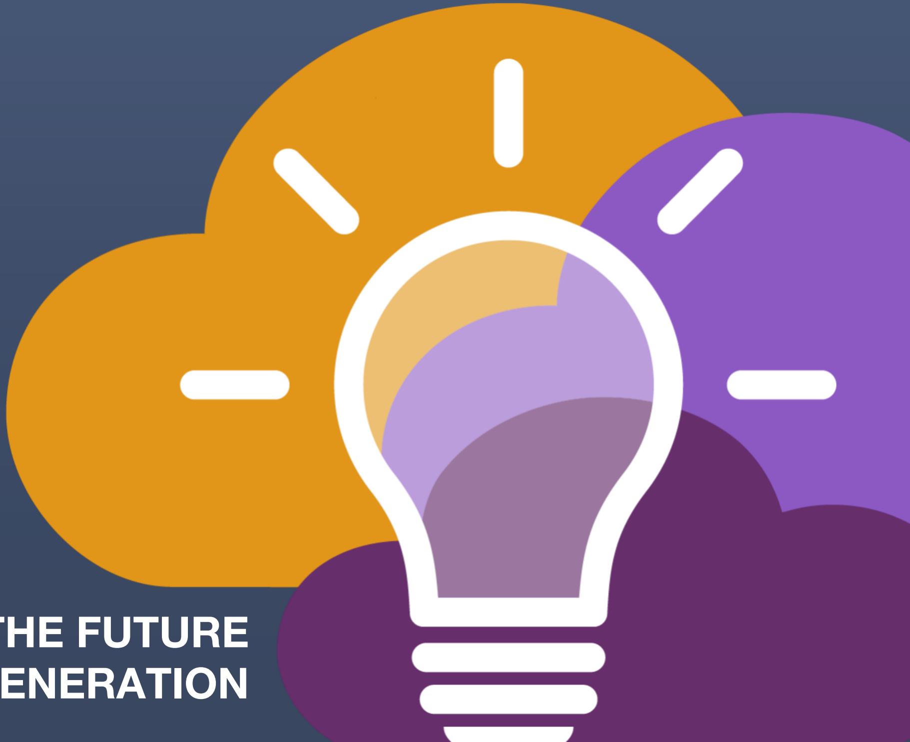
```
BRFSS_table.head()
```

Out[6]:

	_state	fmonth	idate	imonth	iday	iyear	dispcode	seqno	_psu	ctelenm1	...	_frtres1	_vegres1	_frutsu1	_vegesu1	_frtlt1a	_veglt1a	_frt16a	_veg23a	_fruit
0	1	1	1192021	1	19	2021	1100	2021000001	2021000001	1.0	...	1	1	100.0	214.0	1	1	1	1	
1	1	1	1212021	1	21	2021	1100	2021000002	2021000002	1.0	...	1	1	100.0	128.0	1	1	1	1	
2	1	1	1212021	1	21	2021	1100	2021000003	2021000003	1.0	...	1	1	100.0	71.0	1	2	1	1	

ScHARe

Conclusion



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OF KNOWLEDGE GENERATION

ScHARe Terra

In Summary:

- Register for the ScHARe/Terra Platform
- Access federated data – especially SDoH and population science
- Create your own workspace – can share with others
- Assess computational tools, such as python and R coding

ScHARe Repository

In Summary: (Comply with the NIH Data Management and Sharing Policy Requirement)

- When getting started, first create a collection, provide metadata, and upload documents.
- Upload your data and use the system to help you map to the ScHARe CDEs.
- View your data, see CDE compliance and analysis readiness.
- Data can be shared with your colleagues, and can be made publicly available after review.
- You can use dataviews to create subsets of your data, join datasets together and more.
- Data can also be analyzed on the ScHARe Terra workspace.

We are here to support you!

We want to hear your questions, issues and comments about the ScHARe Repository!

For any questions regarding how to use the Repository, please reach out to:
✉ schare@mail.nih.gov
Response time: within 24 hours

For Technical Support inquiries, contact:
✉ schare-repository-support@bioteam.net
🕓 Office Hours: Wednesdays, 4-5 PM US Eastern

Questions?

Slido Poll

What concerns or barriers do you have regarding sharing your data publicly?

Think-a-Thon poll

1. Rate how useful this session was:

☐ Very useful
☐ Useful
☐ Somewhat useful
☐ Not at all useful

Think-a-Thon poll

2. Rate the pace of the instruction for yourself:

☐ Too fast

☐ Adequate for me

☐ Too slow

Think-a-Thon poll

3. How likely will you participate in the next Think-a-Thon?

☐ Very interested, will definitely attend

☐ Interested, likely will attend

☐ Interested, but not available

☐ Not interested in attending any others

ScHARe

Thank you

ScHARe

Next Think-a-Thons:

bit.ly/think-a-thons

Register for ScHARe:

bit.ly/join-schare

✉ schare@mail.nih.gov