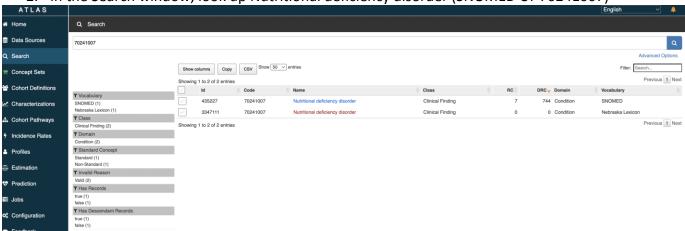
### Session 3: In-class exercises and homework

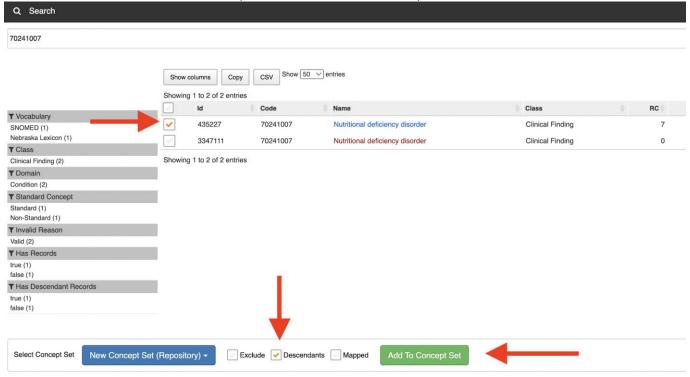
### Exercise 1:

Using Atlas, build a starting concept set for malnutrition.

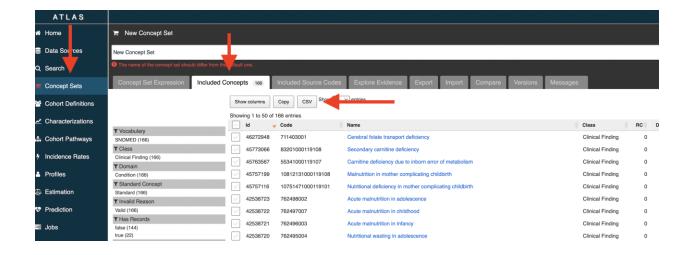
- 1. Go to Atlas: <a href="https://atlas-demo.ohdsi.org/">https://atlas-demo.ohdsi.org/</a>
- 2. In the Search window, look up Nutritional deficiency disorder (SNOMED CT 70241007)



3. Include all descendant concepts and click "Add to Concept Set"



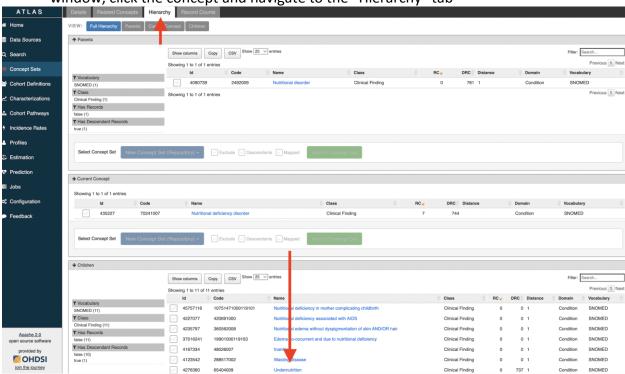
4. View concept set and extract it as a csv using the "Included Concepts" tab



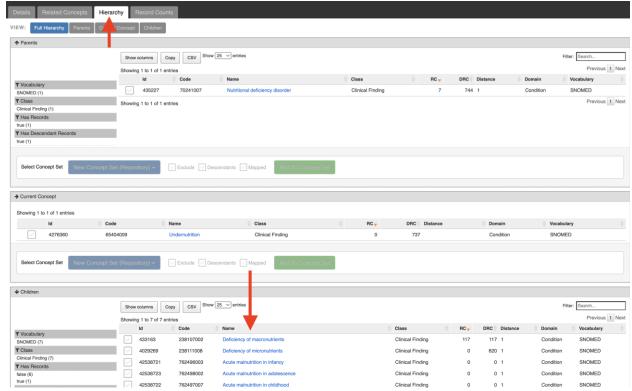
#### Exercise 2:

Using Atlas, refine previous concept set to include only Deficiency of Macronutrients.

1. Starting with Nutritional deficiency disorder (SNOMED CT 70241007) from the search window, click the concept and navigate to the "Hierarchy" tab



- 2. Go to the descendant concept "Undernutrition" and navigate to the "Hierarchy" tab
- 3. Create a new concept set using Deficiency of macronutrients (SNOMED CT 238107002) and add all descendant concepts



4. Extract concept set as a csv

### Exercise 3

# Single-Table Query

- 1. Create a new folder in your workspace and call it Session 3. Within this, create a new code workbook called "Malnutrition Conditions."
- 2. Import the concept table from the data catalog:
  - 1. Click "Import dataset"
  - 2. Navigate to the data catalog and find the folder "OMOP Concepts"
  - 3. Select concept and import into your code workbook
- 3. Select the concept table and select "new transform" then select "SQL code." Name the transform exercise 3
- 4. Write a query to return the concept\_id for Nutritional deficiency using the vocabulary\_id "SNOMED." Save the results to a dataset by toggling "Save as dataset" and run the transform.

SQL code below (in white font, so highlight it):

#### Exercise 4:

## Single-Table Query

- 1. Import the concept\_ancestor table from the data catalog:
  - 1. Click "Import dataset"
  - 2. Navigate to the data catalog and find the folder "OMOP Concepts"
  - 3. Select concept ancestor and import into your code workbook
- Select the concept\_ancestor table and select "new transform" then select "SQL code." Name the transform exercise 4
- 3. Write a query to return every descendant\_concept\_id for Nutritional deficiency using concept\_id returned in exercise 3. Save the results to a dataset by toggling "Save as dataset" and run the transform.

SQL code below (in white font, so highlight it):

### Exercise 5:

## Multiple-Table Query

- 1. Select the resulting transformation from exercise 4 (should be called exercise\_4) and select "new transform" then select "SQL code." Name the transform exercise 5.
- 2. On the new exercise\_5 transformation, add an additional input for the dataset from for the concept table used in exercise 3.
- 3. Write a query to return the resulting set from exercise 4 using a JOIN with the concept table and only include those concepts that are standard concepts using the standard\_concept column (standard\_concept = 'S'). Save the results to a dataset by toggling "Save as dataset" and run the transform. (Note: don't be surprised when this is the same result as exercise 4 since we are sticking within one vocabulary)

#### Exercise 6:

## Multiple-Table Query

- 1. Import the condition\_occurrence table from the data catalog:
  - 1. Click "Import dataset"
  - 2. Navigate to the data catalog and find the folder "De-Identified Data"
  - 3. Select condition occurrence and import into your code workbook
- Select the condition\_occurrence table and select "new transform" then select "SQL code." Name the transform exercise 6.

- 3. Add a second input on the new transformation using the resulting transform from exercise 5 (should be called exercise\_5). This can be done by clicking the "+" icon in the SQL transform or going to the "Inputs" tab at the bottom of the transform.
- 4. Select a data partner to be discussed in class and rite a query to return every row in the condition\_occurrence table from that data partner with a matching condition\_occurrence\_concept\_id from the resulting set of exercise 4. You will need to filter on data\_partner\_id (either in the WHERE clause or in the JOIN). Only include malnutrition records prior to the start of the COVID-19 pandemic in the US (hint: condition\_occurrence\_start\_date prior to '2020-01-01'). Save the results to a dataset by Go through the entire process (exercise 1 through 6) with the parent code for rheumatoid arthritis (69896004): <a href="https://snomedbrowser.com/Codes/Details/69896004">https://snomedbrowser.com/Codes/Details/69896004</a>. Save the results in a new
  - workbook and name it "Rheumatoid Arthritis."
- 5. toggling "Save as dataset" and run the transform.

SQL code below (in white font, so highlight it):

### Homework!

- 1. Follow the steps above, but this time for the micronutrient deficiency SNOMED CT code you identified in exercise 2. SNOMED CT: 70241007. Save the results in a new workbook and name it "Micronutrient Deficiency."
- Go through the entire process (exercise 1 through 6) with the parent code for rheumatoid arthritis (69896004): <a href="https://snomedbrowser.com/Codes/Details/69896004">https://snomedbrowser.com/Codes/Details/69896004</a>. Save the results in a new workbook and name it "Rheumatoid Arthritis."
- 3. Deploy the Logic Liaison All Patients template
  - a. Create a new code workbook and name it Logic Liaison All Patients
  - b. Skip the step to import a dataset
  - c. Import the template: [LOGIC LIAISON TEMPLATE] L2 and L3 Fact Tables: All Patients
  - d. Keep the template as is, but toggle all datasets to save
  - e. Run all transforms