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Sept 30th, 2025



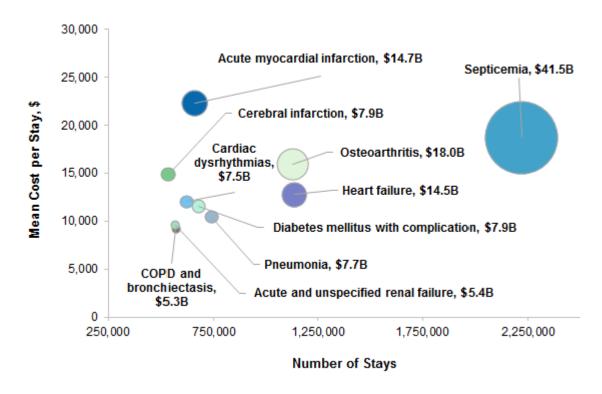


Hasso Plattner Institute for Digital Health at Mount Sinai

Diagnosis Information



- During a visit (encounter) to the hospital, a patient may be diagnosed with a condition, such as diabetes, pneumonia, etc.
- Typically, diagnoses are stored using standard coding systems such as SNOMED and ICD-10
- These coding systems have been built over years by international consortia (for ICD-10) and are constantly being revised
- When you look for diagnoses and pull up lists of conditions, you can consult with clinicians (physicians, specialists, etc) who will have a good idea of how frequent conditions are, how they rank, etc. – this is the process of clinically validating your query results

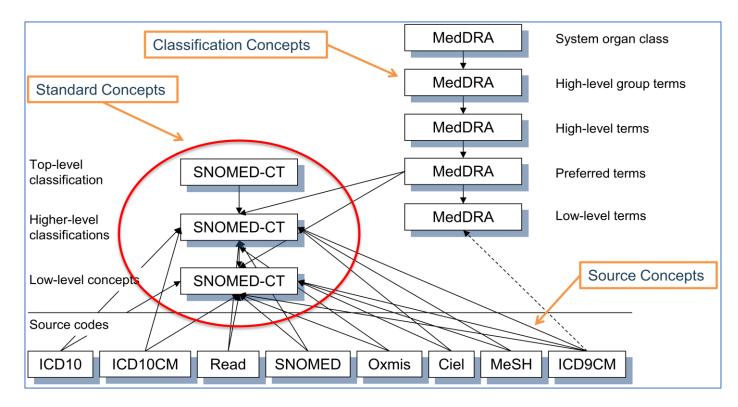


https://hcup-us.ahrq.gov/reports/statbriefs/sb277-Top-Reasons-Hospital-Stays-2018.jsp

ICD-10 and **SNOMED**



- ICD-10: International Classification of Diseases, version 10 (developed internationally by the WHO)
- SNOMED: Systematized Nomenclature of Medicine (developed in the USA)
- There are 357,000 unique concepts in SNOMED, ICD-10-CM has about 70,000
- ICD codes are mapped to the low level and higher-level SNOMED-CT codes
- AIRMS has a custom mapping produced by a 3rd party vendor between Epic IDs and SNOMED / ICD codes



https://ohdsi.github.io/TheBookOfOhdsi/StandardizedVocabularies.html

Examples – ICD-10



E11: Diabetes mellitus without complications

E11.0: Diabetes mellitus with hyperosmolarity

E11.1: Diabetes mellitus with ketoacidosis...

W61.62: Struck by duck

W61.62XA: Struck by duck, initial

encounter

W61.62XD: Struck by duck, subsequent encounter...

E00-E89 Endocrine, nutritional and metabolic diseases

E08-E13 Diabetes mellitus

E11 Type 2 diabetes mellitus

E11.1 Type 2 diabetes mellitus with ketoacidosis

E11.11 with coma



```
# Approach B
# ICD10CM I48* AF codes -> SNOMED standard concepts
code = 'I48%'
vocabulary = 'ICD10CM'
domain = 'Condition'
relationship = 'Maps to'
sql maps to = f"""
SELECT
                       AS source code.
    c1.concept code
    c1.concept name
                       AS source name,
    c1.vocabulary id
                       AS source vocab,
                       AS standard concept id,
    c2.concept id
                       AS standard name,
    c2.concept name
    c2.vocabulary id
                       AS standard vocab
FROM CDMDEID.CONCEPT c1
JOIN CDMDEID.CONCEPT RELATIONSHIP cr
 ON cr.concept id 1 = c1.concept id
JOIN CDMDEID.CONCEPT c2
  ON c2.concept id = cr.concept id 2
WHERE c1.vocabulary id = '{vocabulary}'
  AND c1.concept code LIKE '{code}'
 AND cr.relationship id = '{relationship}'
 AND c2.standard concept = 'S'
 AND c2.domain id = '{domain}'
mapped af = airms.conn.sql(sql maps to).collect()
mapped af
```

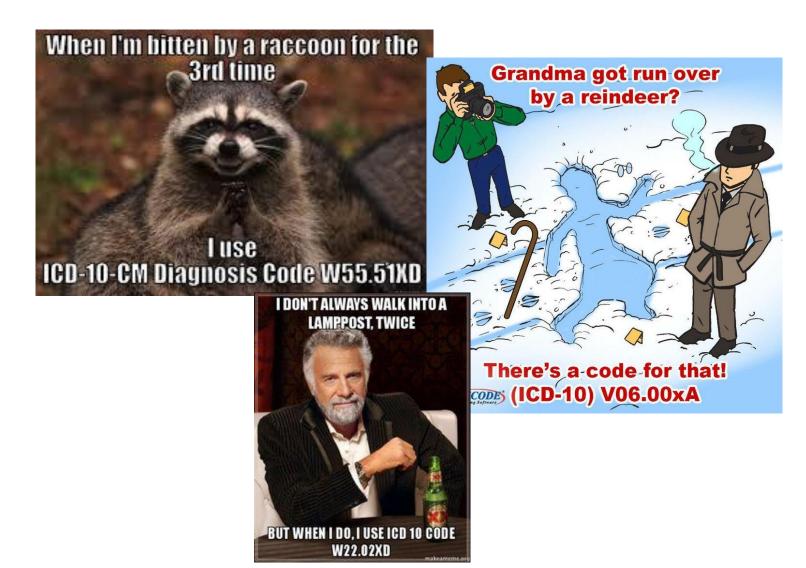
```
# Approach C
# Use mapped SNOMED parents -> expand to descendants
sql af conceptset = f"""
WITH mapped snomed AS (
  {sql_maps_to}
SELECT DISTINCT
    ca.descendant concept id
                                          AS concept id,
    c.concept name
                                          AS concept name,
    c.vocabulary id
                                          AS vocabulary id,
    c.domain id
                                          AS domain id,
    c.standard concept
                                          AS standard flag
FROM CDMDEID.CONCEPT ANCESTOR ca
JOIN CDMDEID.CONCEPT c
  ON c.concept id = ca.descendant concept id
WHERE ca.ancestor_concept_id IN (
  SELECT standard concept id FROM mapped snomed
ORDER BY c.vocabulary id, c.concept name
af concepts = airms.conn.sql(sql af conceptset).collect()
af concepts
```



```
# OUALITY CONTROL STEP
# Identify which source concepts (ICD10CM etc.) actually contributed
# to the atrial fibrillation (AF) cohort in CONDITION_OCCURRENCE.
# This list should be reviewed by a clinical collaborator to confirm
# that all included codes are relevant, and that nothing important
# was missed.
sql sources for review = f"""
WITH af cs AS (
  {sql af conceptset}
af hits AS (
  SELECT
      co.condition_source_concept_id
  FROM CDMDEID.CONDITION OCCURRENCE co
  WHERE co.condition concept id IN (SELECT concept id FROM af cs)
    AND co.condition source concept id IS NOT NULL
    AND co.condition source concept id LIKE REGEXPR '^[0-9]+$'
SELECT
    c.concept id
                                AS source concept id,
    c.concept_code
                                AS source code,
    c.concept name
                                 AS source name,
    c.vocabulary id
                                 AS source vocab,
    COUNT(*)
                                 AS n occurrences
FROM af hits h
JOIN CDMDEID.CONCEPT c
  ON c.concept id = TO INTEGER(h.condition source concept id)
GROUP BY c.concept id, c.concept code, c.concept name, c.vocabulary id
ORDER BY n occurrences DESC, source vocab, source code
af_source_review = airms.conn.sql(sql_sources_for_review).collect()
af source review
```

Other ICD-10 Examples

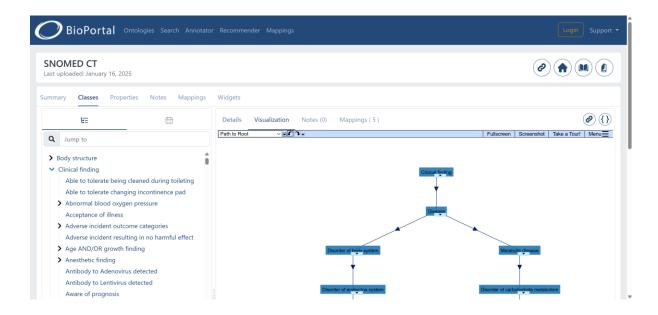




SNOMED CT



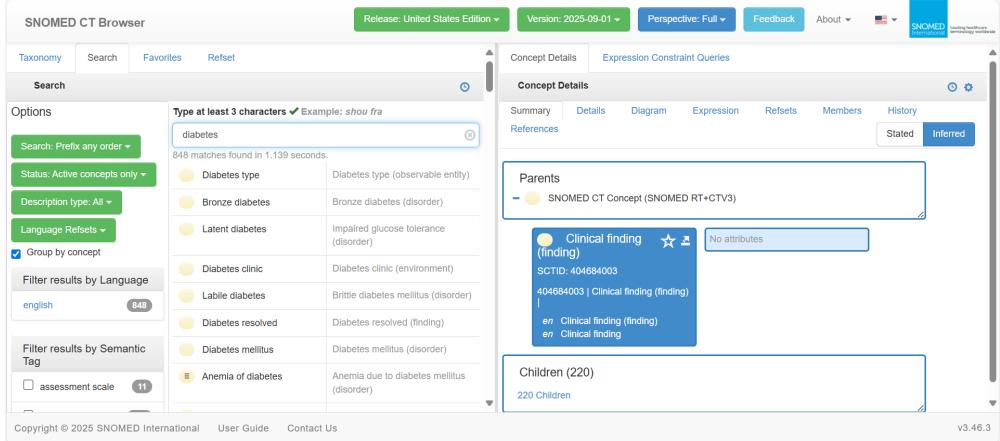
- SNOMED CT stands for Systematized Nomenclature of Medicine - Clinical Terms
- This is a comprehensive terminology for standardizing medical terms, including diagnoses, procedures, and other information
- SNOMED CT is organized into a hierarchical structure
- The main top-level categories include "Clinical Findings", "Procedures", "Organisms", etc.
- Relationships are also defined in SNOMED CT, the most common relationship being "is_a"
- Use Bioportal to browse these relationships: http://purl.bioontology.org/ontology/SNOMEDCT/ 313436004



https://bioportal.bioontology.org/

Examples – SNOMED





https://browser.ihtsdotools.org/

What about Epic IDs?



- Epic has its own internal identification system (Epic ID's) which map to diseases, procedures, medications, etc. These are not standard codes – for research, we need standardized codes
- It also has mapping between Epic IDs and SNOMED CT, ICD-10, etc.
- However, these mapping are commercial, and cannot be used for research purposes, so external
 consultants and companies have to produce them.
- This is why you typically won't find a mapping on the Internet of Epic ID to SNOMED, ICD-10, etc.
- AIRMS OMOP contains its own mapping between Epic IDs and SNOMED/ICD, which was produced with the help of a consulting company

Diagnoses

visit_occurrence				condition_occi	observation			
Patient A	Outpatient Visit 1					Patient A	Past Medical History F	- 6
							Past Medical History H	K11
Patient B	Outpatient Visit 2	Р	Patient B	Outpatient Visit 2	Encounter Diagnosis D4	 Patient B	Past Medical History [04
					Encounter Diagnosis H8		Past Medical History l	.12
Patient C	Inpt Hospitalization 3	Р	Patient C	Inpt Hospitalization 3	Encounter Diagnosis A1			
					Encounter Diagnosis B2			
					Hospital Problem C3			
					Hospital Problem D4			
					Problem List E5			
Patient D	Inpt Hospitalization 4	Р	Patient D	Inpt Hospitalization 4	Encounter Diagnosis F6			
					Billing Diagnosis G7			
		Р	Patient E		Problem List I9			
					Problem List J10			$\wedge \wedge$

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Diagnosis Record Types

xtn_condition_type_ source_concept_id	xtn_condition_type_source_concept_name	row_count
200000108	Billing Diagnosis	73,040,692
200000129	Encounter Diagnosis	120,323,291
200000122	Hospital Problem	3,837,039
200000120	Problem List	13,032,738

Record counts as of April 21, 2025

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```
# How many patients have AF?
print("AF cohort size:", len(af index))
# Make a year column for visualization
af index["year"] = pd.to datetime(af index["AF INDEX DATE"]).dt.year
import matplotlib.pyplot as plt
plt.figure(figsize=(10, 5))
ax = af index["year"].value counts().sort index().plot(
   kind="bar",
   color="#1f77b4",
   edgecolor="black"
plt.title("Number of patients with first AF diagnosis per year", fontsize=14, pad=15)
plt.xlabel("Year of AF index", fontsize=12)
plt.ylabel("Patient count", fontsize=12)
plt.xticks(rotation=45, ha="right")
plt.grid(axis="y", linestyle="--", alpha=0.7)
plt.tight layout()
plt.show()
```

How Do You Pick the Right Codes?



- It's tempting to do a keyword search for a condition, but this is not correct
- For example, if you query for "stroke" you'll miss "cerebral ischemia" (2025 ICD-10-CM Diagnosis Code I67.82)
- As a result: look to existing published studies, or consult a clinician when building a list of conditions to look for
- Clinicians can also help you validate your query results: if you query for a few diseases among a specific demographic (ex. >65 year old males), certain diseases should come up as frequent

Epic to OMOP Procedure Mapping





Procedure Data: CPT



- CPT stands for Current Procedural Terminology
- A commercial code set of codes maintained by the American Medical Association (AMA), requires a license to use
- These codes have played an important role in medical billing, documentation, and reporting
- Currently there are > 11,000 CPT codes
- Commercial will need subscription to tools like Codify
- Mount Sinai has a license, which the AIRMS data uses – limitation on distributing mapping

Code Range	Category		
00100-01999	Anesthesia		
95700-95811	Sleep Medicine Testing Procedures		
10004-69990	Surgery		
70010-79999	Radiology Procedures		
80047-89398	Pathology and Laboratory Procedures		
90281-99607	Medicine Services and Procedures		
98000-99499	Evaluation and Management		
0001F-9007F	Category II Codes		
0002M-0020M	Multianalyte Assay		
0042T-0987T	Category III Codes		

Category II Codes – supplemental performance tracking codes (4 digits with "F" suffix)

Category III Codes – temporary codes for emerging technologies, etc. (4 digits with "T" suffix)

Procedures

Procedure	Surgical Procedure	pro	cedure_occurrence	Flowsheet			
Procedure 1		Patient A	Procedure 1				
Procedure 2		Patient B	Procedure 2				
Procedure 3			Procedure 3				
	Surgical Procedure 4	Patient C	Surgical Procedure 4				
	Surgical Procedure 5	Patient D	Surgical Procedure 5				
		Patient E	Inferred Procedure 6	Patient E	Flowsheet Metric 6-1		
					Flowsheet Metric 6-2		
					Flowsheet Metric 6-3		

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Procedure Record Types

xtn_procedure_type_ source_concept_id	xtn_procedure_type_source_concept_ name	row_count
200000111	General Procedure	326,372,912
2002067235	Procedure Inferred from Flowsheet	281,426
200000097	Surgical Procedure	962,017

Record counts as of April 21, 2025

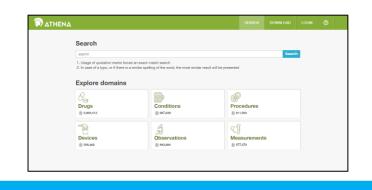
```
# Let's say we are satisfied with all codes identified above, Let's see if we have additional sub-codes to extract
sql pm conceptset = f"""
WITH mapped_std AS (
 {sql_pm_recon}
SELECT DISTINCT
    ca.descendant_concept_id AS concept_id,
                        AS concept_name,
   c.concept name
   c.vocabulary_id
                          AS vocabulary id,
   c.domain_id
                             AS domain_id,
   c.standard_concept
                             AS standard_flag
FROM CDMDEID.CONCEPT ANCESTOR ca
JOIN CDMDEID.CONCEPT c
 ON c.concept_id = ca.descendant_concept_id
WHERE ca.ancestor concept id IN (SELECT concept id FROM mapped std)
ORDER BY c.vocabulary_id, c.concept_name
pm_concepts = airms.conn.sql(sql_pm_conceptset).collect()
pm_concepts
```

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Epic to OMOP Medication Mapping







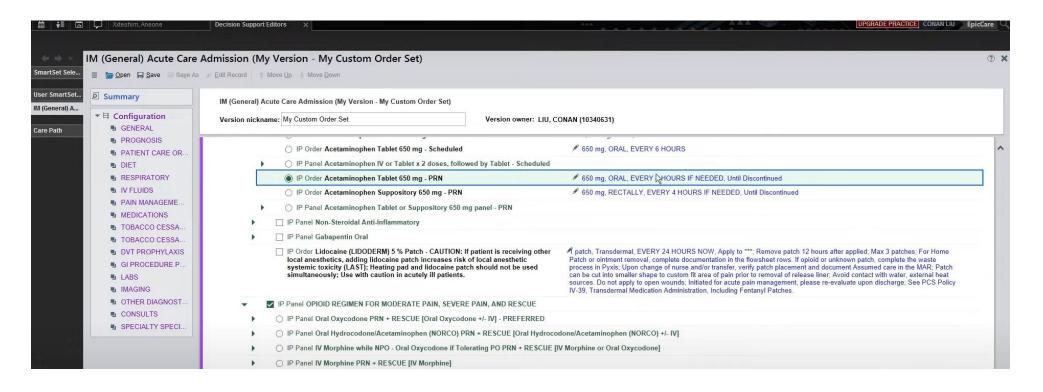
https://athena.ohdsi.org/





Medication Data





- Medications are stored in AIRMS, and contain a wealth of data
- This information is mapped from ATC codes (in Epic) to RxNorm (in OMOP)

Medication Data: RxNorm



- RxNorm this coding standard is used in the OMOP CDM
 - It is developed by the National Library of Medicine (NLM)
 - A comprehensive system that presents a standardized way of representing medications
 - They use unique identifiers, namely a Concept Identifier (RXCUI), that is linked to each medication concept.
 - Identifiers are consistent across different sources and versions

```
Ingredient (IN)

    Precise Ingredient (PIN)

       Clinical Drug Form (CDF)
        Clinical Drug (SCD)

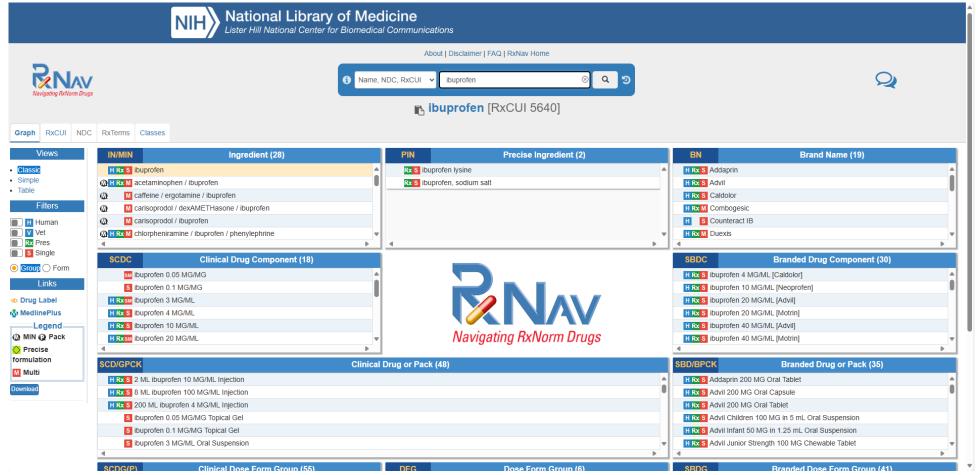
    Branded Drug Component (SBDC)

          ── Branded Drug (SBD)
                Branded Pack (BPCK)
Multiple Ingredients (MIN)
  — Clinical Drug (SCD)
     Branded Drug (SBD)
```

Pack (BPCK)

Medication Data: RxNorm





https://mor.nlm.nih.gov/RxNav/search

RxNorm Examples



1. Ingredient (IN)

The basic chemical component of a drug.

Example: Ibuprofen

2. Precise Ingredient (PIN)

A more specific form of an ingredient, like a salt or ester.

Example: Ibuprofen sodium

3. Multiple Ingredients (MIN)

Represents a combination of ingredients.

Example: Ibuprofen / Hydrocodone

4. Clinical Drug Form (CDF) [Optional level in hierarchy]

Combines an ingredient with a dosage form (without strength).

Example: Ibuprofen Oral Tablet

5. Clinical Drug (SCD = Semantic Clinical Drug)

Includes the ingredient, strength, and dose form, but no brand.

Example: Ibuprofen 200 MG Oral Tablet

6. Branded Drug Component (SBDC = Semantic Branded Drug Component)

A branded version of the SCD with one ingredient.

Example: Advil 200 MG Oral Tablet [Ibuprofen]

7. Branded Drug (SBD = Semantic Branded Drug)

A branded medication with all details (strength, dose form, etc.).

Example: Advil 200 MG Oral Tablet

8. Branded Pack (BPCK) / Clinical Pack (GPCK)

Represents packaging of one or more SBDs or SCDs.

Example: Advil 200 MG Oral Tablet Pack

Medication Data: ATC Codes

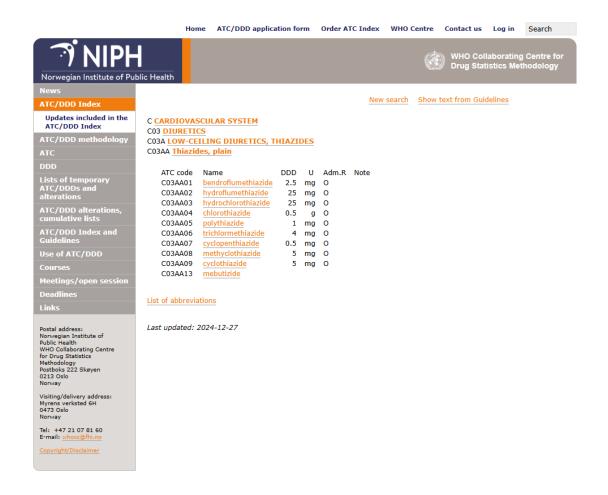


- Anatomical Therapeutic Chemical (ATC)
 Classification System classifies drugs based on the organ system they act on, and their properties (chemical, pharmacological, or therapeutic)
- Produced by the World Health Organization Collaborating Center for Drug Statistics Methodology (WHOCC)
- First published in 1976
- The lowest level of the ATC classification contains 5067 codes
- Simple drug browser available from https://atcddd.fhi.no (Norwegian Institute of Public Health)

Α	Alimentary tract and metabolism (1st level, anatomical main group)
A10	Drugs used in diabetes (2nd level, therapeutic subgroup)
A10B	Blood glucose lowering drugs, excl. insulins (3rd level, pharmacological subgroup)
A10BA	Biguanides (4th level, chemical subgroup)
A10BA02	metformin (5th level, chemical substance)

Medication Data: ATC Codes





https://atcddd.fhi.no/atc ddd index/?code=B02BC&showdescription=no

Medications

Medication Order		Medication Dispense		Medication Administration	Immunization Event			drug_exposure			
						Patier	it A	Vaccine Z	Patient A	Immuniz Z1	Vaccine Z
Patient B	Inpt Med Order					Inpt M Order		Vaccine Y	Patient B	Inpt Med Order 2	Vaccine Y
Patient C	Inpt Med Order 3				Inpt Med Order 3 Med Admin 3-1				Patient C	Med Admin 3-1	Drug X
					Med Admin 3-2					Med Admin 3-2	Drug X
Patient D	Inpt Med Order 4	Inpt Me	d Order Dispense 4-1		Inpt Med Order 4 Med Admin 4-1				Patient D	Med Admin 4-1	Drug W
			Dispense 4-2		Med Admin 4-2					Med Admin 4-2	Drug W
Patient E	Inpt Med Order 5	Inpt Me 5	d Order Dispense 5-1						Patient E	Dispense 5-1	Drug V
Patient F	Outpt Med Order 6	Outpt M Order 6	Dispense 6-1						Patient F	Dispense 6-1	Drug U
Patient G	Outpt Med Order 7	Outpt M Order 7	DISPENSE /-T						Patient G	Dispense 7-1	Drug T
			Dispense 7-2							Dispense 7-1	Drug T
Patient H	Outpt Med Order 8			-					Patient H	Outpt Med Order 8	Drug S

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Medication Record Types

xtn_drug_type_source _concept_id	xtn_drug_type_source_concept_name	row_count
2002056202	Immunization Administration	7,835,276
2002056203	Immunization from Medication Order	289,032
2002056204	Immunization from Procedure Order	3,488,313
200000121	Medication Order	13,670,853
200000110	Medication Order with Administration	138,120,557
200000098	Medication Order with Dispense	16,351,836
200000096	Outpatient Medication Order	48,588,821
200000109	Outpatient Medication Order with Dispense	151,086

Record counts as of April 21, 2025

```
A Confirm the ancestor concept (ATC class)
antiplatelet_concept = 35807468
sql ap atc info = f""
SELECT concept id, concept name, vocabulary id, concept class id, domain id
FROM COMDETO, CONCEPT
WHERE concept_id = {antiplatelet_concept}
airms.conn.sql(sql_ap_atc_info).collect()
# Pull RxNorm Ingredients that descend from the ATC Antiplatelet class (concept_id = 35101523)
vocabulary = 'RxNorm'
concept_class = 'Ingredient'
sql_ap_ingredients_from_atc = f"""
   ca.max_levels_of_separation,
   c.concept_id
                         AS ingredient id.
                         AS ingredient name,
   c.concept_name
   c.vocabulary_id,
   c.concept_class_id,
   c.domain_id,
   c.standard_concept
FROM COMDEID.CONCEPT_ANCESTOR ca-
DOIN COMDEID.CONCEPT C
  ON c.concept_id + ca.descendant_concept_id
WHERE ca.ancestor_concept_id = (antiplatelet_concept)
  AND c.vocabulary_id = '{vocabulary}'
  AND c.concept_class_id = '(concept_class)'
  AND c.invalid_reason IS NULL
ORDER BY ca.max_levels_of_separation, c.concept_name
ap_ingredients = airms.conn.sql(sql_ap_ingredients_from_atc).collect()
ap_ingredients
```

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Lab Results & Vital Signs (measurements)

Lab Order		Lab Con		mponent Result		measurement			Flowsheet		
Patient A	Lab Order 1	L	Lab Order 1	WBC Count		Patient A	WBC Count				
				RBC Count			RBC Count				
				Hct Result			Hct Result				
				Hgb Result			Hgb Result				
Patient B	Lab Order 2	L	Lab Order 2	Glucose Result		Patient B	Glucose Result				
				Calcium Result			Calcium Result				
				Sodium Result			Sodium Result				
				BUN Result			BUN Result				
						Patient C	Height		Patient C	Height	
							Weight			Weight	
							Temp			Temp	
						Patient D	Metric 3		Patient D	Metric 3	
						Patient E	Metric 4		Patient E	Metric 4	

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Measurement Record Types

xtn_measurement_type _source_concept_id	xtn_measurement_type_source _concept_name	row_count
2002067233	Flowsheet Measurement	254,696,901
200000100	Lab Component Result	1,107,913,246
200000123	Vital Signs	692,312,136

Record counts as of April 21, 2025

```
## BMI Measurement near AF Index

In OMOP, anthropometric values like BMI live in the **MEASUREMENT** table.

- Each row represents a quantitative measurement (`value_as_number`) recorded at a date (`measurement_date`).

- Concepts come from vocabularies like **LOINC** or **SNOMED**.

For BMI, we use the standard concept **"Body mass index"** and its descendants.

**Approach:**

1. Identify BMI concepts (LOINC + SNOMED descendants).

2. Pull all BMI rows for AF patients.

3. For each patient, find the **BMI value closest in time to AF index**.
```

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Find BMI concepts in OMOP (LOINC + SNOMED)

SELECT concept id, concept name, vocabulary id, domain id, concept class id

concept name = 'body mass index'

domain = 'Measurement'

FROM CDMDEID.CONCEPT

sql bmi conceptset = f"""

WHERE domain_id = '{domain}'
AND standard concept = 'S'

concept class = 'Clinical Observation'

Content Standardization via

Concept Mapping

OMOP Data Example (synthetic)



condition_occurrence	Value	Definition
condition_type_concept_id	32827	OMOP's standard record type "EHR encounter record"
xtn_condition_type_source_concept_id	200000129	MSDW's source record type "Encounter Diagnosis"
visit_occurrence_id	999888777	Unique identifier for the encounter
person_id	987654321	Unique identifier for the patient
provider_id	123456789	Unique identifier for the provider recording the diagnosis
condition_concept_id	4193704	OMOP's identifier for SNOMED code 313436004
condition_source_concept_id	2000602205	MSDW's identifier for Epic diagnosis code 521601
condition_start_date	1/1/2022	The date of condition onset or documentation per provider
condition_end_date	NULL	The date on which the condition resolved (if any)

concept_relationship	Value	Definition
concept_1	2000602205	MSDW's identifier for Epic diagnosis code 521601
relationship_id	Maps to non-standard	Text string denoting the type of mapping relationship between concept_id_1 & concept_id_2
concept_2	35206882	OMOP's identifier for ICD-10-CM code E11.9

OMOP Data Example (synthetic)



concept

condition_occurrence	Value	concept_id	vocabulary_id	concept_code	concept_name
condition_type_concept_id	32827	32827	Type Concept	OMOP4976900	EHR encounter record
xtn_condition_type_source_concept_id	200000129	200000129	MSDW Src Rec Type	Encounter Diagnosis	Encounter Diagnosis
visit_occurrence_id	999888777				
person_id	987654321				
provider_id	123456789				
condition_concept_id	4193704	4193704	SNOMED	313436004	Type 2 diabetes mellitus without complication
condition_source_concept_id	2000602205	2000602205	EPIC EDG .1	521601	Type 2 diabetes mellitus without complications
condition_start_date	1/1/2022				
condition_end_date	NULL				
concept_relationship	Value				
concept_1	2000602205	2000602205	EPIC EDG .1	521601	Type 2 diabetes mellitus without complications
relationship_id	Maps to non- standard				
concept_2	35206882	35206882	ICD10CM	E11.9	Type 2 diabetes mellitus without complications

Visualizing and Analyzing Data in Jupyter



Age and Sex from PERSON

We now extend the cohort with **demographics**:

- **Sex** is stored in PERSON.gender_concept_id. We join to the CONCEPT table for a readable label.
- Age at AF index is calculated from PERSON.birth_date relative to AF_INDEX_DATE.

This information is critical for describing the cohort and adjusting analyses.

```
# Querying age and sex
sql_pm_af_ap_age_sex = f"""
WITH cohort base AS (
 {sql_pm_af_ap_bmi}
SELECT
   cb.*,
   p.birth_datetime,
   g.concept_name AS sex,
   FLOOR(MONTHS_BETWEEN(p.birth_datetime, cb.af_index_date) / 12) AS age_at_af
FROM cohort base cb
JOIN CDMDEID.PERSON p
 ON p.person id = cb.person id
JOIN CDMDEID.CONCEPT g
 ON g.concept id = p.gender concept id
ORDER BY cb.af index date
0.00
pm af ap age sex = airms.conn.sql(sql pm af ap age sex).collect()
pm af ap age sex.head()
```



Exploratory Data Analyses

We summarize cohort size, demographics, therapies near diagnosis, device procedures, and BMI capture. Figures are de-identified and derived from the **de-identified OMOP** dataset.

Cohort anchor: first AF diagnosis (AF_INDEX_DATE).

Therapy window: antiplatelet exposure overlapping [AF_INDEX_DATE, +6 months].

Pacemaker outcome: first pacemaker procedure on/after AF_INDEX_DATE.

BMI: nearest measurement within ±180 days of AF_INDEX_DATE.

Preparing the data ¶



```
# Find BMI concepts in OMOP (LOINC + SNOMED)
concept name = 'body mass index'
concept class = 'Clinical Observation'
domain = 'Measurement'
sql bmi conceptset = f"""
SELECT concept id, concept name, vocabulary id, domain id, concept class id
FROM CDMDEID.CONCEPT
WHERE domain id = '{domain}'
 AND standard concept = 'S'
  AND concept_class_id = '{concept_class}'
  AND LOWER(concept name) LIKE '%{concept name}%'
  AND invalid reason IS NULL
bmi_concepts = airms.conn.sql(sql_bmi_conceptset).collect()
bmi concepts
# We want the actual ration that was observed (i.e. 3038553)
sql bmi conceptset = 'SELECT * FROM CDMDEID.CONCEPT WHERE CONCEPT ID=3038553'
# Build on your existing cohort (sql pm after af) and append BMI nearest to AF index
# Assumes sql pm after af returns: person id, af index date, first pm date, days to pacemaker,
                                 on antiplatelet within 6mo, first ap start in window, last ap end in window
sql pm af ap bmi = f"""
WITH cohort_base AS (
  {sql_pm_af_ap}
bmi cs AS (
  {sql bmi conceptset}
```

Observations

Patient	Demographics	Surgio	cal History	Soc	ial History	Famil	y History	Patient Allergy	ok	oservation
Patient A	Race								Patient A	Race
	Ethnicity									Ethnicity
	Language									Language
	Preference Sexual									Preference Sexual
	Orientation									Orientation
Patient B	Race								Patient B	Race
	Marital Status									Marital Status
	Gender Identity									Gender Identity
Patient C	Ethnicity								Patient C	Ethnicity
	Religious									Religious
	Affiliation									Affiliation
		Patient A	Procedure Z						Patient A	Procedure Z
			Procedure Y							Procedure Y
		Patient B	Procedure X						Patient B	Procedure X
			Procedure W							Procedure W
				Patient C	Soc Hx Item 1-1				Patient C	Soc Hx Item 1-1
					Soc Hx Item 1-2					Soc Hx Item 1-2
					Soc Hx Item 1-3					Soc Hx Item 1-3
						Patient D	Fam Hx 2-1		Patient D	Fam Hx 2-1
							Fam Hx 2-2			Fam Hx 2-2
								Patient E Allergy 3-1	Patient E	Allergy 3-1

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Observation Record Types

xtn_observation_type_ source_concept_id	xtn_observation_type_source_concept _name	row_count
2002056205	Allergy	4,277,961
200000128	Family History	65,724,728
2002067234	Flowsheet Observation	114,620,841
200000124	Past Medical History	8,610,801
200000116	Patient Demographics	64,233,860
200000118	Social History	145,273,185
200000126	Surgical History Procedure	3,681,599

Record counts as of April 21, 2025

```
t.xtn_observation_type_source_concept_id

, t.xtn_observation_type_source_concept_name

, FORMAT(COUNT_BIG(*), 'NO') AS row_count

FROM omop.cdm_phi.observation t

WHERE t.observation_id <> 0

GROUP BY

t.xtn_observation_type_source_concept_id

, t.xtn_observation_type_source_concept_name

ORDER BY

t.xtn observation type source concept name
```

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Putting it Together: Queries & Phenotypes

Querying the Data & Phenotypes

- When you're researching a disease (ex. Lyme disease) just looking for a particular International Classification of Diseases (ICD) code or disease keyword ("lyme") is not enough
 - You need to consult clinical practice guidelines
 - Sometimes there's an initial diagnosis that's wrong you may need to look for lab results (ex. 2 separate diagnoses of diabetes within 6 months of each other, and an HbA1c level)
 - So, you might write a query that looks for patients with three Type 2 diabetes codes within a 6-month period, and two HbA1c levels above a certain threshold, rather than simply looking for patients with one Type 2 diabetes code
 - When possible, consult physicians and clinicians when devising how to identify patients with a particular disease or phenotype, peer-reviewed publications are also a good source of information
 - For example NIH N3C specific codes for COVID-19: https://tinyurl.com/4dwruasx

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Checklist

ltem	Comments
Check if Patient IDs are deleted, merged, orphan, or substituted, where possible check against MPI	MPI = master patient index
When selecting codes, consult a clinician or use peer- reviewed literature	
Do not use keyword searches on vocabularies	Ex. "stroke" vs. "cerebral ischemia"
Use mapping/ vocabulary browser tools when necessary: Athena, Bioportal, RxNav, etc.	Remember: Epic IDs are licensed, so you won't find public mappings for Epic IDs vs. other coding systems
When constructing your digital phenotype, use clinical practice guidelines to inform the development of your phenotype	Get a clinician to help you understand a clinical practice guideline

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Thank You