

— — OHDSI

Observational Health Data Sciences and Informatics (OHDSI)

2025-02-24



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OHDSI OHDSI OHDSI

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- I. OHDSI□□□□□□
  - II. □□□□□□□□□□
  - III. □□□□□
  - IV. □□□□□□□
  - V. OHDSI□□

Hamed Abedtash	Mustafa Ascha	Mark Beno
Clair Blacketer	David Blatt	Brian Christian
Gino Cloft	Frank DeFalco	Sara Dempster
Jon Duke	Sergio Eslava	Clark Evans
Thomas Falconer	George Hripcsak	Vojtech Huser
Mark Khayter	Greg Klebanov	Kristin Kostka
Bob Lanese	Wanda Lattimore	Chun Li
David Madigan	Sindhoosha Malay	Harry Menegay
Akihiko Nishimura	Ellen Palmer	Nirav Patil
Jose Posada	Nicole Pratt	Dani Prieto-Alhambra
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Erica Voss	Kristin Waite	Mike Warfe
Jamie Weaver	James Wiggins	Andrew Williams
Seng Chan You		

OHDSI

- ACHILLES: 1.6.6
- ATLAS: 2.7.3
- EUNOMIA: 1.0.0
- OMOP: 1.0.0

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RMarkdown bookdown  
<https://github.com/OHDSI/TheBookOfOhdsiInJapanese/>  
 “travis”

Table 1: Methods Library

CaseControl	1.6.0
CaseCrossover	1.1.0
CohortMethod	3.1.0
Cyclops	2.0.2
DatabaseConnector	2.4.1
EmpiricalCalibration	2.0.0
EvidenceSynthesis	0.0.4
FeatureExtraction	2.2.4
MethodEvaluation	1.1.0
ParallelLogger	1.1.0
PatientLevelPrediction	3.0.6
SelfControlledCaseSeries	1.4.0
SelfControlledCohort	1.5.0
SqlRender	1.6.2



**Part I**

**OHDSI**



# Chapter 1

— — OHDSI

□□□□□□□□□□□□□□□□ & □□□□□□□□□□

1.1

(Olsen et al., 2007) of Medicine Roundtable on Evidence-Based Medicine 2020 90 (Olsen et al., 2007)

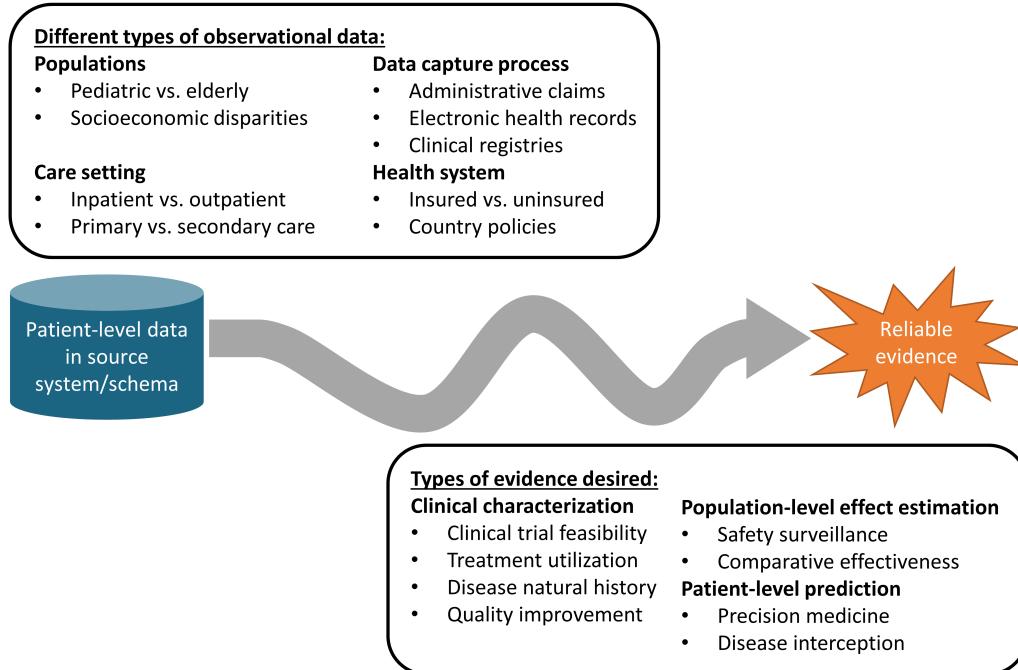
1.2

OMOP OMOP  
Institutes of Health (Stang et al., 2010) OMOP

(Overhage et al., 2012) OMOP ## HDSI

Observational Health Data Sciences and Informatics (OHDSI) □□□□□□□□□□

et al., 2015) OHDSI



**Figure 1.1:** □□□□□□□□□□□□

### 1.2.1

1.2.2

A horizontal row of 20 empty white rectangular boxes, likely used for input fields or placeholder text in a form.

1,2,3

- **OMOP:** [http://www.commondatamodel.org](#)
  - **OMOP:** [http://www.commondatamodel.org](#)
  - **OMOP-CDM:** [http://www.commondatamodel.org](#) **OHDSI** [http://www.ohdsi.org](#)
  - **OMOP:** [http://www.commondatamodel.org](#)
  - **OMOP:** [http://www.commondatamodel.org](#)
  - **OMOP:** [http://www.commondatamodel.org](#)

### 1.3 OHDSI

OHDSI<sup>1</sup> 2014  
OHDSI<sup>2</sup> 1.2



Figure 1.2: 2019<sup>3</sup> 8<sup>4</sup> OHDSI<sup>5</sup>

2019<sup>3</sup> 8<sup>4</sup> OHDSI<sup>5</sup> 20<sup>6</sup> 100<sup>7</sup>  
CDM<sup>8</sup> 10<sup>9</sup>

OHDSI<sup>10</sup> OMOP CDM<sup>11</sup>  
<sup>12</sup>)  
OHDSI<sup>13</sup> CDM<sup>14</sup> OHDSI<sup>15</sup>

OHDSI<sup>16</sup>  
Academy of Science<sup>17</sup> 2500<sup>18</sup> 11<sup>19</sup>  
OHDSI<sup>20</sup> (Tian et al., 2018)<sup>21</sup>  
et al., 2017)<sup>22</sup> (Vashisht et al., 2018)<sup>23</sup>  
et al., 2018b)<sup>24</sup> OHDSI<sup>25</sup>  
et al., 2018)<sup>26</sup> (Johnston et al., 2019; Cepeda et al.,  
2018; Reps et al., 2019)

<sup>1</sup><https://www.ohdsi.org/who-we-are/collaborators/>

<sup>2</sup><https://github.com/OHDSI>

## 1.4 OHDSI

OHDSI OHDSI  
2 OHDSI

## 1.5



- OHDSI
- OHDSI
- OHDSI

# Chapter 2

— —

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□ □ □ □ □ □ □ □ □ □ - □ □

OHDSI 



# Chapter 3

2.1

OHDSI 

### 3.0.1 2.1.1 OHDSI

OHDSI  <sup>1</sup> OHDSI 



to OHDSI! - Please introduce yourself OHDSI - 2)



### 3.0.2 2.1.2 OHDSI

OHDSI 

OHDSI 

<sup>1</sup><https://forums.ohdsi.org>

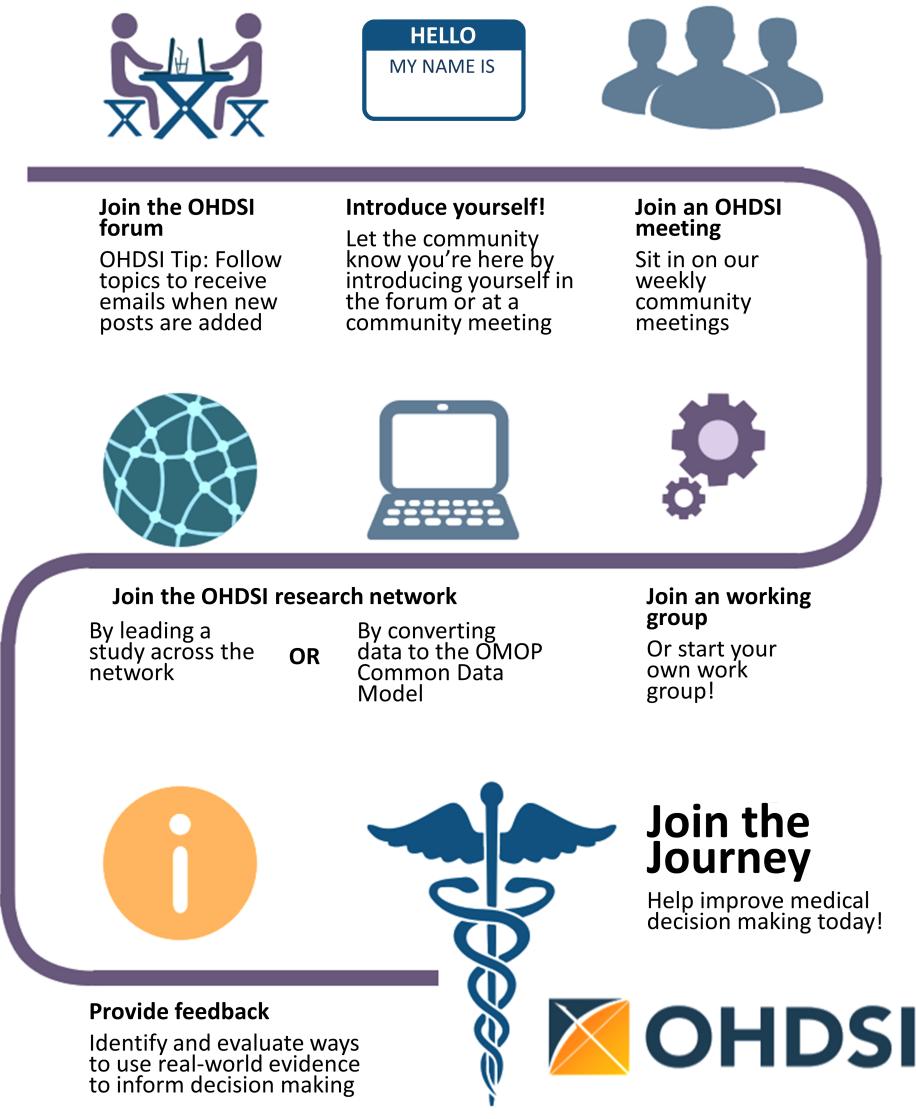


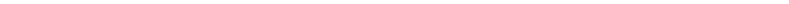
Figure 3.1: □□□□□□□ - OHDSI□□□□□□□□□□□□□

# OHDSI

### 3.0.3 2.1.3 OHDSI

OHDSI □□□□□□□□□□□□□□□ OHDSI □□□□□□□□□□□□□□□

### 3.0.4 2.1.4 OHDSI

OHDSI  OHD

Wiki OHDSIWiki](<https://www.ohdsi.org/web/wiki/doku.php?id=projects:overview>) □□□□□□□□□□

□: □□□□□ OHDSI □□□□□□

Atlas & WebAPI	Atlas	WebAPI	OMOP	Atlas/WebAPI & JavaScript
CDM & Vocabulary	OMOP	OMOP	OMOP	OMOP
Genomics	OMOP CD-	OMOP	OMOP	OMOP
	M	OMOP	OMOP	OMOP
	CDM	OMOP	OMOP	OMOP
Population-Level Estimation	OMOP	OMOP	OMOP	OMOP
Natural Language Processing	OHDSI	OHDSI (EHR)	OHDSI	OHDSI
Gold Standard Phenotype Library	OHDSI	OHDSI	OHDSI	OHDSI
FHIR Workgroup	OHDSI	FHIR	OHDSI	EHR
GIS	OMOP CD-	OMOP	OMOP	OMOP

### 3.0.5 2.1.5 OHDSI

## 3.0.6 2.1.6 OHDSI

OHDSI OMOP OHDSI  
CDM

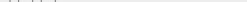
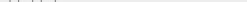
3.1

OHDSI  
OHDSI  
11 12 13 12 13 14  
**OHDSI**  
CDM OHDSI / OHDSI  
CDM OHDSI  
8 OMOP CDM OHDSI  
**OMOP CDM ETL/**

OMOP ETL  
CDM  
  
**OHDSI** R  
Render OHDSI/SqlRender GitHub  
  
**OHDSI**  
OHDSI Apache  
  
**OHDSI**  
  
**OHDSI** OHDSI

3.2



- OHDSI  **OHDSI** 
  - ETL  **OHDSI** 



# Chapter 4

— — — — —

4.1

2.0 (Wikipedia, 2019b) hacking Fisher 2019) et al., 2016) et al., 2018)

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4.2

1

4.3

OHDSI OMOP Chapter 5 Chapter 6 OHDSI (Garza et al., 2016)

4.4

OHDSI Methods Library Research Network Library ATLAS Athena GitHub OHDSI

4.5

//howoften.org http://data.ohdsi.org  
Research Network 21 OMOP  
CDM OHDSI ETL

4.6

OHDSI

## 4.7 OHDSI FAIR

#### 4.7.1

### 4.7.2

OMOP et al., 2019) IMI EHDEN

### 4.7.3

OMOP SQL  
CDM OMOP  
EHDEN OMOP

#### 4.7.4

OMOP OHDSI OM



## **Part II**

— —



# Chapter 5

□ □ □: *Clair Blacketer*

A horizontal row of 20 empty square boxes, intended for students to write their answers in a grid format.

CDM 6 CDM

CDM  5.1 

5.1

CDM



CDM

- **CDM**: **Contracting Out**
  - **CDM**: **Joint Venture**
  - **CDM**: **Partnership**

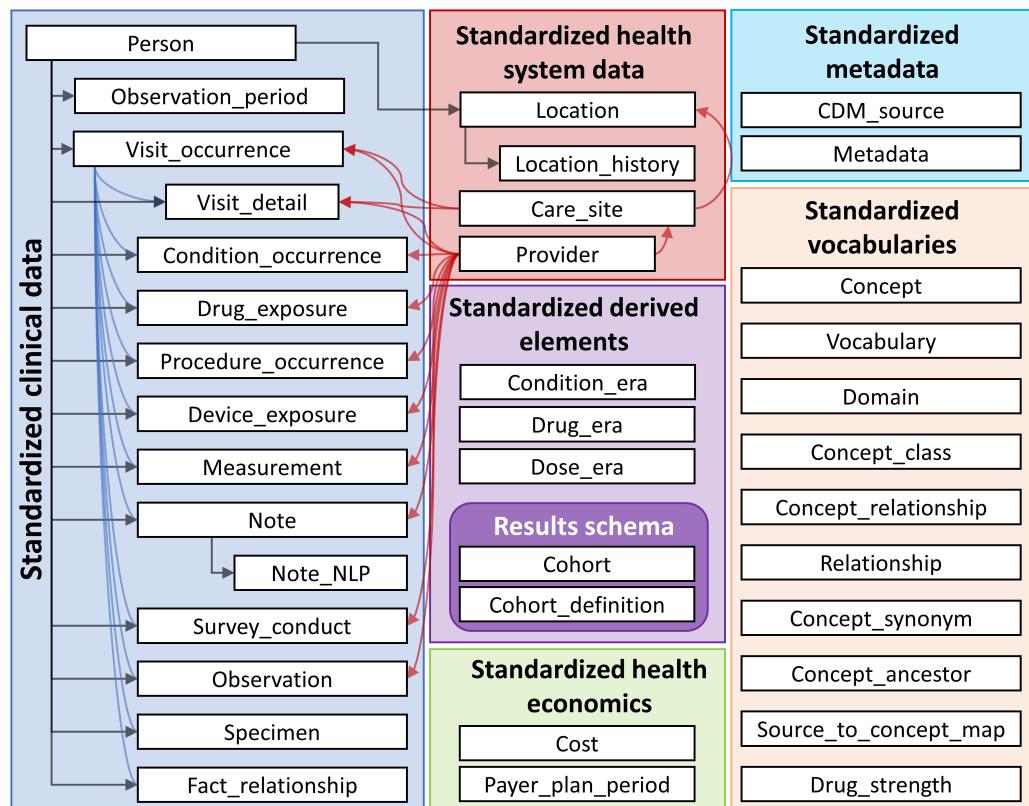


Figure 5.1: CDM 6.0

5.2

### 5.2.1

5.2.2

11 COHORT

5.2.3

524

A horizontal row of 20 empty square boxes, likely for grading student responses.

□□□□□□	□□□□ID	□□□□□□	□□□□ID
1731378	Drug	183	Route
477597	Device	180	Currency
257000	Procedure	158	Payer
163807	Condition	123	Visit

□□□□□□	□□□□ID	□□□□□□	□□□□ID
145898	Observation	51	Cost
89645	Measurement	50	Race
33759	Spec Anatomic Site	13	Plan Stop Reason
17302	Meas Value	11	Plan
1799	Specimen	6	Episode
1215	Provider Specialty	6	Sponsor
1046	Unit	5	Meas Value Operator
944	Metadata	3	Spec Disease Status
538	Revenue Code	2	Gender
336	Type Concept	2	Ethnicity
194	Relationship	1	Observation Type

## 5.2.5

5.2.6

□□□□□□□□□□□□□□□□□□□□□□□□1□□□□□□□□□□□□

5.2.7

- ICD9CM □ NDC □ Read □
  - CDM □

- CDM
- CDM

QA  
5.2 ICD9CM 011



The screenshot shows the ATHENA interface with the title "Pulmonary tuberculosis". Below it is a table titled "DETAILS" containing the following information:

Domain ID	Condition
Concept Class ID	3-dig nonbill code
Vocabulary ID	ICD9CM
Concept ID	44828631
Concept code	011
Invalid reason	Valid
Standard concept	Non-standard
Synonyms	Pulmonary tuberculosis
Valid start	12/31/1969
Valid end	12/30/2099

Figure 5.2: ICD9CM

011 UB04 A DRG  
TB Figure 5.3 OMOP  
253954 Read ICD10 CIEL MeSH

5.7

### 5.3 CDM

CDM 16 10 2 4  
CDM Wiki<sup>1</sup>

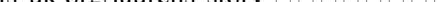
1 https://github.com/OHDSI/CommonDataModel/wiki

TERM CONNECTIONS (82)			
RELATIONSHIP	RELATES TO	CONCEPT ID	VOCABULARY
ICD-9-CM to MedDRA (MSSO)	Pulmonary tuberculosis	36110777	MedDRA
Non-standard to Standard map (OMOP)	Pulmonary tuberculosis	253954	SNOMED
Subsumes	Other specified pulmonary tuberculosis	44830894	ICD9CM
	Other specified pulmonary tuberculosis, bacteriological or histological examination not done	44836741	ICD9CM
	Other specified pulmonary tuberculosis, bacteriological or histological examination unknown (at present)	44836742	ICD9CM
	Other specified pulmonary tuberculosis, tubercle bacilli found (in sputum) by microscopy	44821641	ICD9CM
	Other specified pulmonary tuberculosis, tubercle bacilli not found (in sputum) by microscopy, but found by bacterial culture	44833188	ICD9CM

Figure 5.3: SNOMED

### 5.3.1 :



Lauren   
[//endometriosis-uk.org/laurens-story](http://endometriosis-uk.org/laurens-story) 

### 5.3.2 PERSON

- □□□36□□□□
  - □□□□□□□1982□3□12□□
  - □□□□□□□
  - □□□□□□□□□□□

PERSON

Table 5.3: PERSON□□□□□

PERSON_ID	1	PERSON_ID
GENDER_CONCEPT_ID	8532	ID 8532
YEAR_OF_BIRTH	1982	
MONTH_OF_BIRTH	3	
DAY_OF_BIRTH	12	
BIRTH_DATETIME	1982-03-12 00:00:00	
DEATH_DATETIME		
RACE_CONCEPT_ID	8527	ID 8527
ETHNICITY_CONCEPT_ID	38003564	US
LOCATION_ID		
PROVIDER_ID		
CARE_SITE		
PERSON_SOURCE_VALUE	1	
GENDER_SOURCE_VALUE	F	
GENDER_SOURCE_CONCEPT_ID	0	OHDSI
RACE_SOURCE_VALUE	white	PCORNet
RACE_SOURCE_CONCEPT_ID	0	
ETHNICITY_SOURCE_VALUE	english	GENDER_SOURCE_CONCEPT_ID
ETHNICITY_SOURCE_CONCEPT_ID	0	GENDER_SOURCE_CONCEPT_ID

### **5.3.3 OBSERVATION\_PERIOD**

Table 5.4: Lauren

□□□□□□□ID	□□□	□□□	□□□
70	2010-01-06	2010-01-06	□□□□

ID			
80	2011-01-06	2011-01-06	□□□□
90	2012-01-06	2012-01-06	□□□□
100	2013-01-07	2013-01-07	□□□□
101	2013-01-14	2013-01-14	□□□□
102	2013-01-17	2013-01-24	□□□□

## OBSERVATION\_PERIOD

#### 5.3.4 VISIT\_OCCURRENCE

## VISIT\_OCCURRENCE

VISIT\_OCCURRENCE

Table 5.6: VISIT\_OCCURRENCE

VISIT_OCCURRENCE_ID	514	
PERSON_ID	1	PERSON Laura
VISIT_CONCEPT_ID	9201	9201
VISIT_START_DATE	2013-01-17	
VISIT_START_DATETIME	2013-01-17 00:00:00	
VISIT_END_DATE	2013-01-24	
VISIT_END_DATETIME	2013-01-24 00:00:00	

-  wiki

### 5.3.5 CONDITION\_OCCURRENCE

## CONDITION OCCURRENCE

Table 5.7: CONDITION\_OCCURRENCE

EHR ID 32020 |  
|CONDITION\_STATUS\_CONCEPT\_ID|NULL|  
4203942 |STOP\_REASON|NULL|

5.4

5.5






5.6

10 of 10

CDM ATHENA<sup>3</sup> ATLA

<sup>2</sup><https://github.com/OHDSI/CommonDataModel/wiki>

<sup>3</sup><http://athena.ohdsi.org/>

<sup>4</sup><http://atlas-demo.ohdsi.org>

□□ 5.3. □□□□ 2019 □ 5 □ 1 □□□□□□□□□ 200 MG □□□□□ NDC □□□□ 76168009520 □ 30 □

10 of 10

□□□□3□□□□□□□□□@ref□installIR□□□□□□□□□R□R-Studio□□□□Java□□

```
install.packages(c("SqlRender", "DatabaseConnector", "remotes"))
remotes::install_github("ohdsi/Eunomia", ref = "v1.0.0")
```

Eunomia  R  CDM 

```
connectionDetails <- Eunomia::getEunomiaConnectionDetails()
```

CDM□□□□□□□□□□□ main□□□□□□ CONDITION OCCURRENCE□□□□□□

```
library(DatabaseConnector)
connection <- connect(connectionDetails)
sql <- "SELECT *
FROM @cdm.condition_occurrence
LIMIT 1;"
result <- renderTranslateQuerySql(connection, sql, cdm = "main")
```

□□ 5.6. SQL R □□□□□ PERSON ID 61 □□□□□□□□□□□□□□□□□□□□□□

□□□□□□□□□□@ref□Cdmanswers□□□□□□□



# Chapter 6

□□□□□: Christian Reich & Anna Ostropolets

OMOP  OHDSI 

6.1

of Mortality 6.1

OHDSI  OMOP CDM  CDM 



ОМОП CRM

6.1.1

CDM  OHDSI 

GitHub 

<sup>1</sup><https://github.com/OHDSI/Vocabulary-v5.0>

<sup>2</sup><https://forums.ohdsi.org>

<sup>3</sup><https://github.com/OHDSI/CommonDataModel/issues>

1669.

A General BILL for this present Year,

Ending the 11th Day of December 1660.

According to the Report made to the King's most excellent Majesty,  
By the Company of Parish Clerks of LONDON, &c.

## DISEASES and CASUALTIES.

A Bortive and Stillborn	421	Flox and Small Pox	1523	Palsy			17
Aged	909	Found dead in the Streets,	1523	Plague			36
Ague and Fever	2303	Fields, &c.	2	Plurify			12
Apoplexy and Suddenly	91	French Pox	51	Quinify and fore Throat			21
Blaisted and Planet	3	Gout	4	Rickets			44
Bleeding and bloody Issue	7	Grief	13	Rising of the Lights			210
Bloody Flux, Scowring, and Flux	346	Griping in the Guts	253	Rupture			12
Burnt and Scalded	6	Hanged and made away them-selves	11	Scurvy			82
Cancer, Gangrene and Fistula	63	Head-ach and Headmouldshot	35	Shot			7
Canker, fore Mouth and Thrush	73	Jaundies	102	Shingles			1
Childbed	226	Imposthume	105	Sores, Ulcers, broken and bruised Limbs			61
Chrisomes and Infants	858	Killed by several Accidents	55	Spleen			7
Cold, Cough and Hiccough	33	King's Evil	28	Spotted Fever and Purples			368
Colick and Wind	116	Lethargy	6	Starved			7
Consumption and Tiswick	2982	Livergrown	8	Strangury			22
Convulsion	742	Lunatick and Frenzy	14	Stopping of the Stomach			186
Cut of the Stone and Stone	46	Megrims	5	Surfeit			202
Dropfy and Tympany	646	Measles	6	Swine Pox			2
Drowned	57	Mother	1	Teeth and Worms			839
Executed	7	Murthered	7	Vomiting			8
Falling Sicknes	4	Overlaid and Starved at Nurse	46	Wen			1

Figure 6.1: 1660 62

### 6.1.2

Pallas ATHENA<sup>4</sup>

**OMOP CDM** zip

### 6.1.3 :

OHDSI Extension 6.6.9

6.2

## OMOP CDM

CONCEPT_ID	313217	Primary key
CONCEPT_NAME	Atrial fibrillation	English description
DOMAIN_ID	Condition	Domain
VOCABULARY_ID	SNOMED	Vocabulary
CONCEPT_CLASS_ID	Clinical Finding	Class in vocabulary
STANDARD_CONCEPT	S	Standard, Source of Classification
CONCEPT_CODE	49436004	Code in vocabulary
VALID_START_DATE	01-Jan-1970	Valid during time interval
VALID_END_DATE	31-Dec-2099	
INVALID_REASON		

Figure 6.2: OMOP CDM

621 ID

6.2.2

<sup>4</sup><http://athena.ohdsi.org>

### 6.2.3

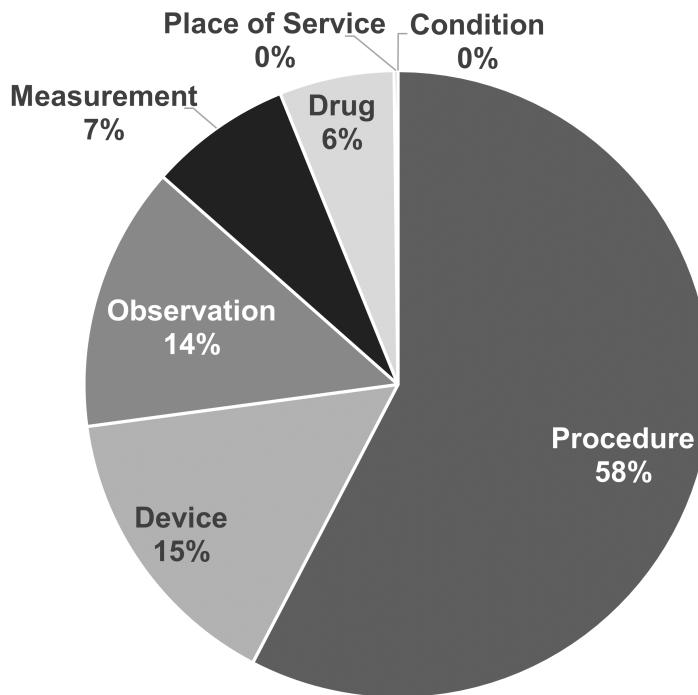
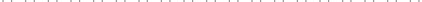
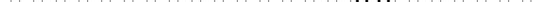


Figure 6.3: CPT4 HCPCS

5  **CDM**  6.6

6.2.4

## 6.2.5

Class ID

A horizontal row of 15 empty rectangular boxes, likely used for input fields or placeholder text in a form.

RxNox

## 6.2.6

## 6.2.7

6,2,8

- □ 6.4 □ □ □

6.2.9

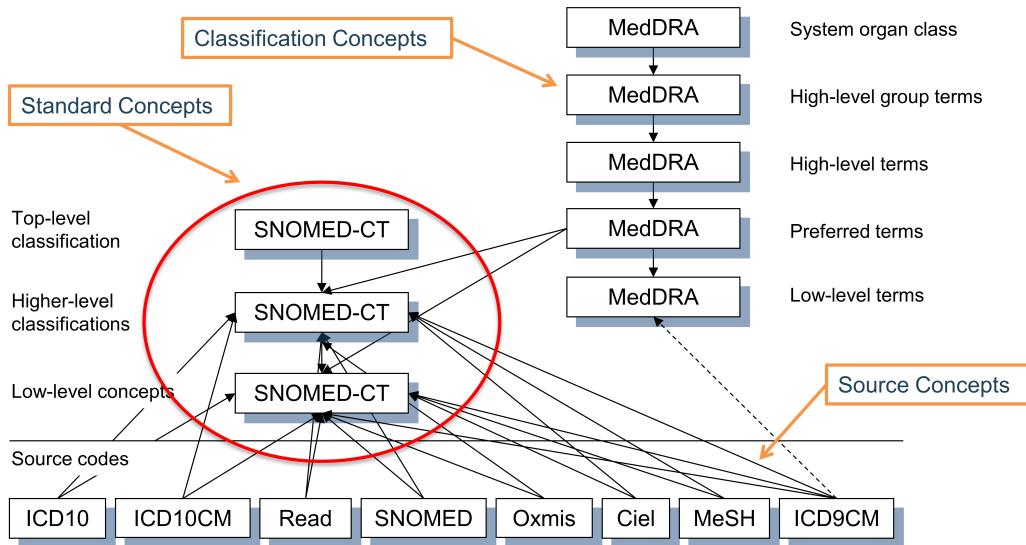


Figure 6.4: SNOMED

ID		ID		ID	
35803438	1001			HemOnc	
35942070	1001	AJCC TNM		NAACCR	NAACCR
		Clin T			
1036059	1001			RxNorm	
38003544	1001				
	-				
43228317	1001			BDPM	
45417187	1001			Multum	Multum
		mg/ml			
45912144	1001			CIEL	

## 6.2.10

CDM

- **VALID\_DATE**
  - **VALID\_DATE**: **DATE**
  - **VALID\_START\_DATE**: **DATE** 1-1
  - **VALID\_END\_DATE**: **DATE** 12-31
  - **INVALID\_REASON**: **NULL**
- **VALID\_DATE**
  - **VALID\_DATE**: **DATE** 6.2.6

- VALID\_START\_DATE:  1-10
  - VALID\_END\_DATE:
  - INVALID\_REASON: "D"
  - - 
    - VALID\_START\_DATE:  1-10
    - VALID\_END\_DATE:
    - INVALID\_REASON: "U"
  - - 
    - VALID\_START\_DATE:

6.3

to Mapped from

### 6.3.1

2 IP

Table 6.4:

□□ID□□ □□  
“Maps  
to”□□□”Mapped  
from”  
“Maps to  
value”□□□”Value  
mapped from”  
MEASUREMENT□□□OBSERVATION□□□□□VALUE\_AS\_CONCEPT\_1

W61.51 SNOMED  
217716004

Maps to value - EAV-OMOP CD-M

- 
- 
- 
- 

to Maps to value

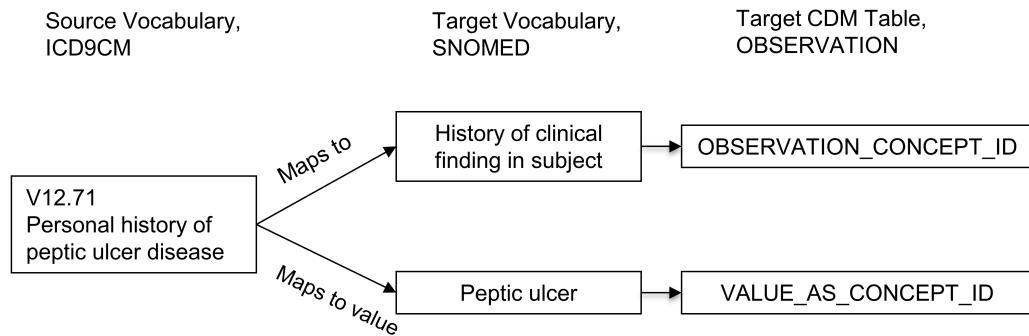


Figure 6.5: Maps to value

OMOP OHDSI Wiki<sup>5</sup>

### 6.3.2

Is a - Subsumes 49436004 SNOMED 17366009 Is a 49436004 SNOMED 40593004 Is a

### 6.3.3

Vocabulary A - Vocabulary B equivalent - RxNorm equivalent Maps to

### 6.3.4

OHDSI Wiki<sup>6</sup>

<sup>5</sup><https://www.ohdsi.org/web/wiki/doku.php?id=documentation:vocabulary:mapping>

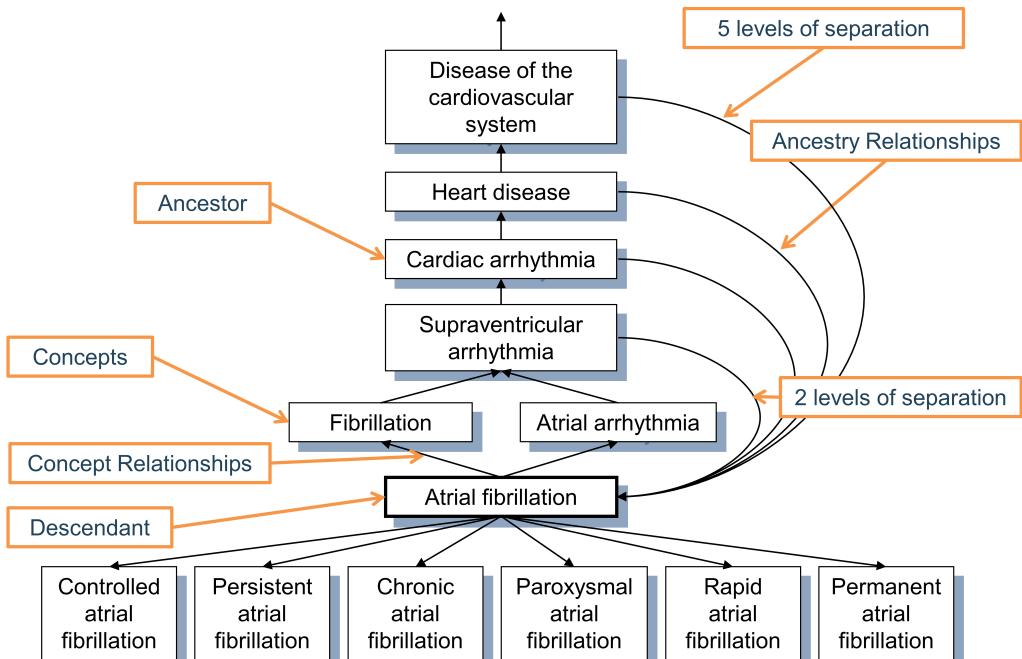
<sup>6</sup><https://www.ohdsi.org/web/wiki/doku.php?id=documentation:vocabulary>

site of 6.5

Table 6.5: Finding site

CONCEPT_ID_1	CONCEPT_ID_2
4000504 “Urethra part”	36713433 “□□□□□□□”
4000504 “Urethra part”	433583 “□□□□□□”
4000504 “Urethra part”	443533 “□□□□□□□”
4000504 “Urethra part”	4005956 “□□□□□□□”

CONCEPT\_ANCESTOR□□□□□ CONCEPT\_RELATIONSHIP□□□□□□□□□□□  
"Is a" - "Subsumes" □□□□ 6.6 □□□□□□□□□□□□□□□□□□□□□□□□□□□



MIN LEVELS OF SEPARATION MAX LEVELS OF SEPARATION

6.5

6.6

6.6.1

ОМОР CRM

### 6.6.2

6,6,3

OMOP

ICD-9 ICD-10

#### 6.6.4

HCPCS CPT4

6.6.5

6.6.6

6.6.7

6.6.8

6.6.9

RxNorm Extension

### 6.6.10 NULL

5 8507

6.7



- OMOP
  - OHD
  - CDM
  - Mapping
  - Maps to Maps to value
  - 
  - 
  -

6.8

1

ATHENA<sup>7</sup> ATLAS<sup>8</sup>

## 6.1. “ ” ID

**6.2. “ ” ICD-10CM 9CM**

### **6.3. “ ” MedDRA**

E.2

<sup>7</sup><http://athena.ohdsi.org/>

<sup>8</sup><http://atlas-demo.ohdsi.org>



# Chapter 7

□□□□□□□: *Clair Blacketer & Erica Voss*

7.1

□□□□/□□□□□ OMOP □□□□□ CDM □□□□□ ETL □

ETL 4

1. CDM ETL
  - 2.
  3. ETL
  - 4.

OHDSI

## □□□□ 1· ETL□□□

### 7.1.1 White Rabbit

Rabbit ETL White Rabbit OMOP CDM ETL Rabbit ETL

---

<sup>1</sup><https://github.com/OHDSI/WhiteRabbit>

5.1.1.1

□ □ □ □ □ □

A horizontal row of 20 empty square boxes, intended for students to write their answers in a handwriting practice exercise.



A horizontal row of nine empty square boxes, each with a thin black border, intended for children to draw or color in.

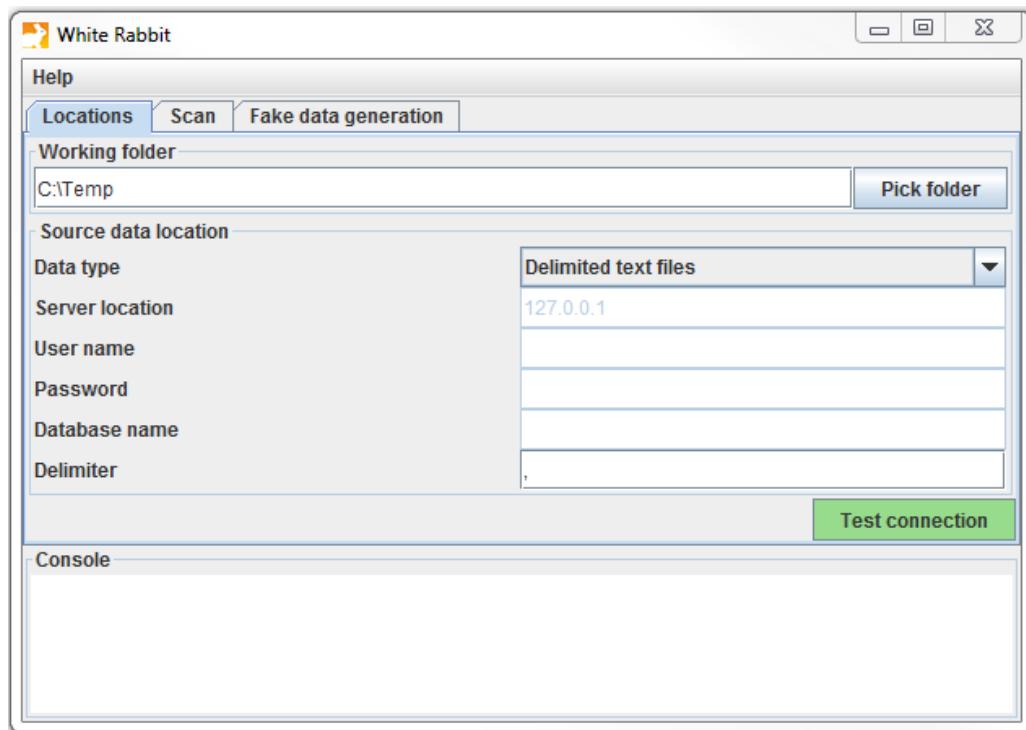


Figure 7.1: White Rabbit Pick  
Folder

## White Rabbit

A horizontal row of fifteen empty square boxes, intended for children to draw or write in.

ETL + all in DB

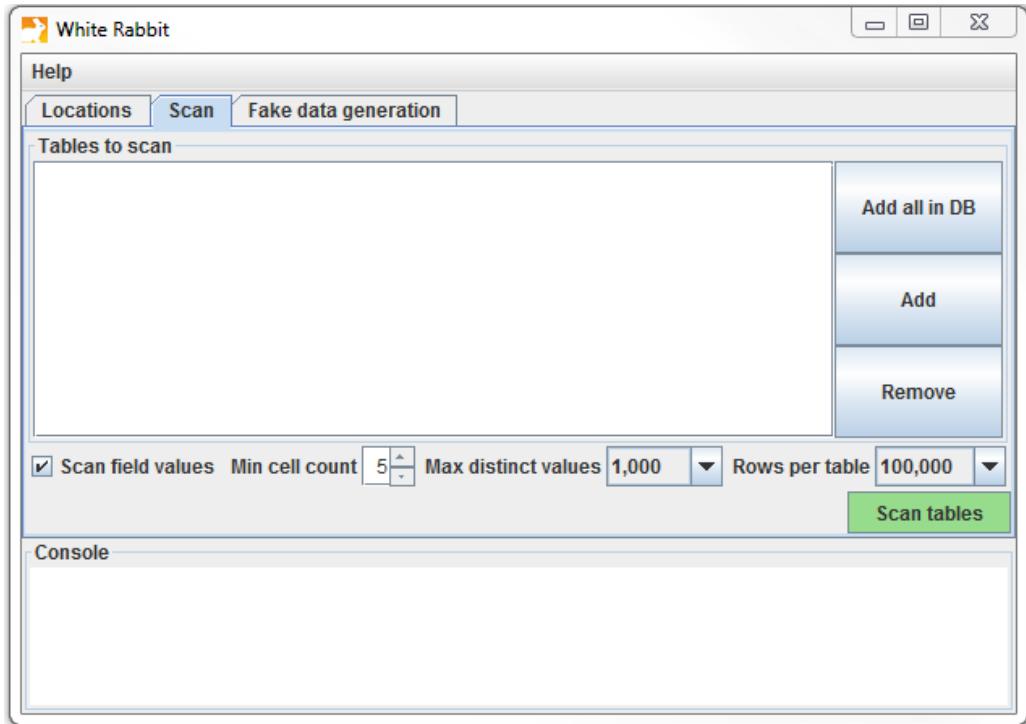


Figure 7.2: White Rabbit □□□□□□□□

A horizontal row of 20 empty square boxes, intended for children to write their names in, likely as part of a classroom activity or name recognition exercise.

- White Rabbit 5 White Rabbit 100,000

Excel

A	B	C	D	E	F	G
1 Table	Field	Type	Max length	N rows	N rows checked	Fraction empty
2 dbo.allergies	start	date	10	3184	3184	0
3 dbo.allergies	stop	date	10	3184	3184	0.725188442
4 dbo.allergies	patient	varchar	36	3184	3184	0
5 dbo.allergies	encounter	varchar	36	3184	3184	0
6 dbo.allergies	code	varchar	9	3184	3184	0
7 dbo.allergies	description	varchar	24	3184	3184	0
8						
9 dbo.careplans	id	varchar	36	30199	30199	0
10 dbo.careplans	start	date	10	30199	30199	0
11 dbo.careplans	stop	date	10	30199	30199	0.057849598
12 dbo.careplans	patient	varchar	36	30199	30199	0
13 dbo.careplans	encounter	varchar	36	30199	30199	0
14 dbo.careplans	code	varchar	15	30199	30199	0
15 dbo.careplans	description	varchar	62	30199	30199	0
16 dbo.careplans	reasoncode	varchar	9	30199	30199	0.050796384
17 dbo.careplans	reasondescription	varchar	56	30199	30199	0.050796384
18						

◀ ▶
Overview
dbo.allergies
dbo.carepla ...
+ :
◀ ▶
▼

Figure 7.3: □□□□□□□□□□□□□□□□□□

A	B
1 Sex	Frequency
2	2 61491
3	1 35401
4	List truncated...

Figure 7.4: □□□□□□□□□□

Rabbit 1 2 7.4

## 7.1.2 Rabbit-In-a-Hat

□ □ □ □ □

Rabbit-In-a-Hat White Rabbit  
Rabbit Rabbit-In-a-Hat  
In-a-Hat ETL ETL

ANSWER

ETL



## ETL

## ETL

## Person

Synthea patients 2076. patients CDM PERSON

?? Synthea patients CDM PERSON  
Field CDM Source  
field CDM patients  
& comments

<sup>2</sup>Syntheia™

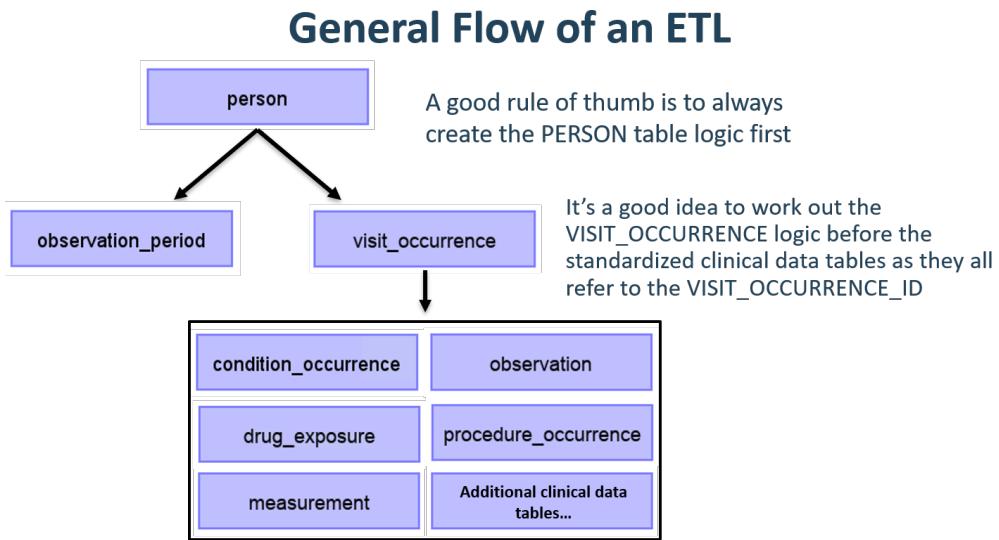


Figure 7.5: ETL

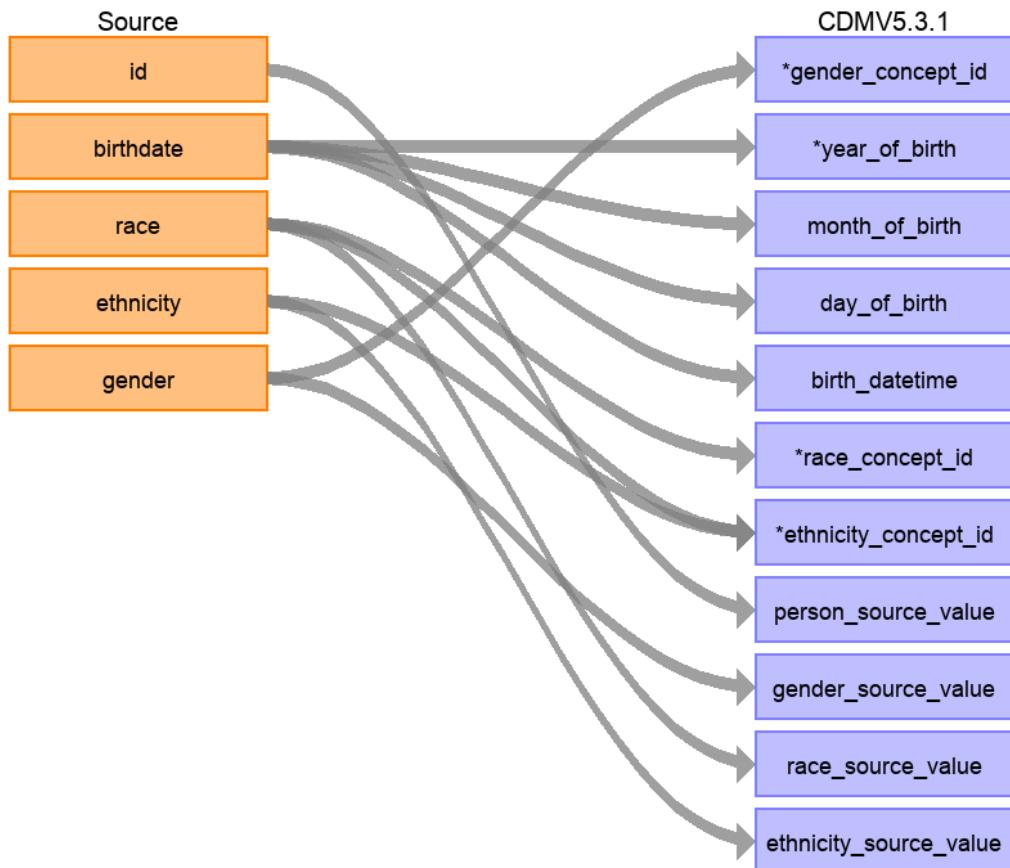


Figure 7.6: Synthea Patients → CDM PERSON

□□(□tab:syntheaEtlPerson) Synthea Patients □□□□□ CDM PERSON □□□□□□□□□ ETL

7.2 2:

```
SELECT concept_id_2 AS standard_concept_id
FROM concept_relationship
INNER JOIN concept AS source_concept
    ON concept_id = concept_id_1
WHERE concept_code = 'I21'
    AND vocabulary_id = 'ICD10CM'
    AND relationship_id = 'Maps to';
```

---

## STANDARD\_CONCEPT\_ID

---



### 7.2.1 Usagi

Usagi  
Translate<sup>3</sup> Usagi

1

□□□□□□□□□□□□□□□□□Usagi□□□□□□□□□□□□□□□□□

<sup>3</sup><https://translate.google.com/>

<sup>4</sup><https://github.com/OHDSI/Usagi>

# □□□□□□Usagi□□□□□

□□□□□□CSV□□□Excel (.xlsx)

## Translate

File → Import codes... 7.7

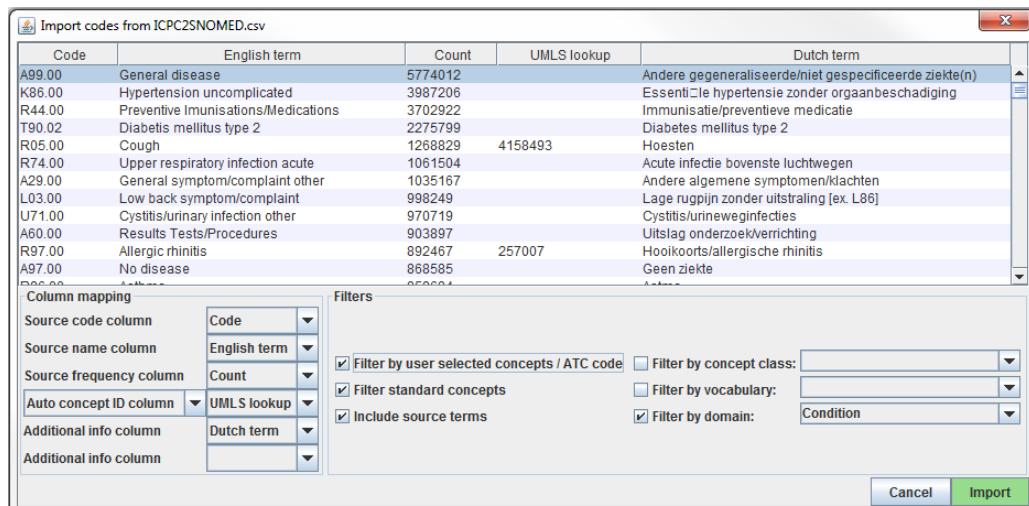


Figure 7.7: Usagi□□□□□□□□□

1. Column mapping “Auto concept ID column” ATC column
  2. Column mapping ATC AT

7.3 3: ETL

ETL



7.4 4:

- ETL 1
  - 1
  - 1
  - CDM 1
  - ETL 1

ETL OHDSI 16

## 7.5 ETL THEMIS

CPM ETL

<sup>5</sup><https://github.com/OHDSI/CommonDataModel/wiki>

WikiThemis

## 7.6 CDM ETL

ETL□□□□□□□□□□□□□□□ ETL□□□□□□□□□□□□□□□

# ОМОП

CDM ETL

7.7 ETL

## ETL

- 80/20
  - 
  - CDM ETL
  - OHDSI CDM

7.8



- ETL
  - \* CDM
  - \* OHDSI
  - \* ETL
  - \* OHDSI
  - OHDSI
  - ETL

7.9

7.1. ETL

- A) □□□□□□□CDM□□□□□□□□□ETL□□□  
B) □□□□ETL□□□

<sup>6</sup><https://github.com/OHDSI/Themis>

<sup>7</sup><http://forums.ohdsi.org/>

<sup>8</sup><https://forums.ohdsi.org/c/implementers>

- C) □□□□□□□□□□□□□□□□□  
D) □□□□□□□□□□□□□□□

## □□ 7.2. □□□□OHDSI□□□□□□□□□□ 7.2□□PERSON□□□□□□□□□□ 4□□□□□□□□□□

□: PERSON □□□□□

PERSON_ID	A123B456
GENDER_CONCEPT_ID	8532
YEAR_OF_BIRTH	NULL
MONTH_OF_BIRTH	NULL
DAY_OF_BIRTH	NULL
RACE_CONCEPT_ID	0
ETHNICITY_CONCEPT_ID	8527
PERSON_SOURCE_VALUE	A123B456
GENDER_SOURCE_VALUE	F
RACE_SOURCE_VALUE	WHITE
ETHNICITY_SOURCE_VALUE	□□□□□□□□□□

□□ 7.3. VISIT\_OCCURRENCE □□□□□□□□□□□□□□□ Synthea □□□□□□□  
PATIENT □ START □ END □□□□□□□□□□□□□ PERSON\_ID □□□□□□□ END □□□□

- MIN(START) □ VISIT\_START\_DATE □ □ □ □
  - MAX(END) □ VISIT\_END\_DATE □ □ □ □
  - “IP” □ PLACE\_OF\_SERVICE\_SOURCE\_VALUE □ □ □ □

□□□□□□□□□ 7.8 □□□□□□□□□□□□□□□ CDM □□□□□ VISIT OCCURRENCE

Data Output						Explain	Messages	Notifications	Query History
	id character varying (1000)	start date	stop date	patient character varying (1000)	encounterclass character varying (1000)				
1	12	2004-09-26	2004-09-27	11	inpatient				
2	13	2004-09-27	2004-09-30	11	inpatient				

Figure 7.8: □□□□□□□□□

E.3



## **Part III**

---



# Chapter 8

□□□□□□□□: *David Madigan*



Chapters 15 - 19

8.1

A horizontal row of fifteen empty rectangular boxes, evenly spaced, intended for the respondent to mark with an X or checkmark.

A horizontal row of 20 empty rectangular boxes, likely for students to write their answers in a worksheet.



- □□□□□□□...□
  - □□□□□□□□□□...□
  - □□□□□□□...□
  - □□□□□□□□□□□□□□□□□□□□□□...□
  - □□□HbA1c□□□□...□
  - □□□□□□□...□

- □□□□□□□□□□□□
  - □□□
  - □□□□□
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  - □□□□□□□□
  - □□□□□□□□
  - □□□□□□□□

8.2

A horizontal row of eight empty rectangular boxes, likely used for input fields or placeholder text in a form.

A horizontal row of 20 small, empty rectangular boxes, likely used for input fields or placeholder text in a form.



A horizontal row of 20 small, empty rectangular boxes, likely used for a grid or table structure.

- ...□□□□□
  - □□...□□□□□□□
  - □□□□□□□□□□□
  - Y□□□□X□□□□□
  - ...□□□□□□□□□

- - 
  - 
  - 
  - 
  - 
  - 
  - 
  -

8.3



A horizontal row of 20 empty square boxes, likely used for grading or marking student responses.

- 
  - 

A horizontal row of fifteen empty square boxes, intended for children to draw or color in.

- □□□□□
  - □□□□□
  - □□□□□/□□□□□□□□□□
  - □□□□□□



8.4

ACE

8.4.1

11 Chapter 12

## 8.4.2

Chapter 13 ACE

### 8.4.3

14 ACE 1

8.5

OHDSI 



## 8.5.1

8.5.2

OHDSI et al. (2017)

8.6



- □□□□□□□3□□□□□□□□□□□□□□□□□□
  - □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□
  - □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□
  - □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□
  - □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□
  - □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

8.7

## 8.1.

□□□□□□□□□ Appendix E.4 □□□□□



# Chapter 9

— — OHDSI

□□□□□ *Lead: Martijn Schuemie & Frank DeFalco*

# OHDSI

9.1

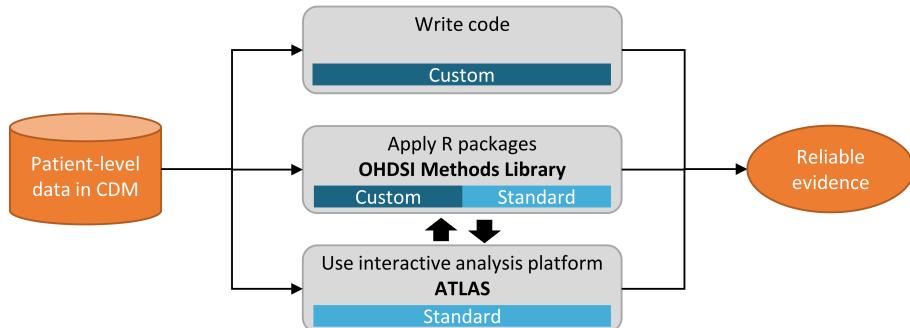
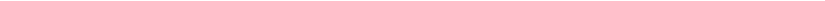


Figure 9.1: CDM

3 OHDSI  
R OHDSI Methods Library Ch  
10 SqlRend  
Library

30 ATLAS Libraries

Library  ATLAS

ATLAS Methods Library Similarly, cohorts used in the Methods Library are often designed in ATLAS.

9.2

# CDM Methods Library

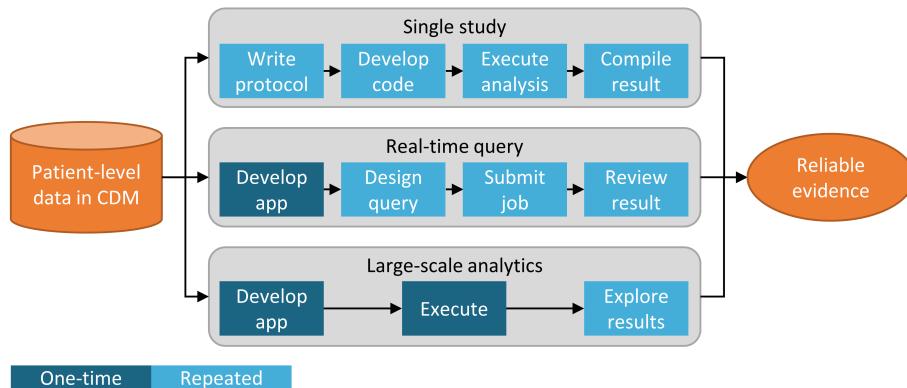


Figure 9.2: (□□□)□□□□□□□□□□□□□□□□□□□□□□□□□□

(Duke et al., 2017, ) OHDSI Methods Library OHDSI

(Schuemie et al., 2018b, ) //data.ohdsi.org/SystematicEvidence/

## 9.3 ATLAS

ATLAS CDM OHDSI  
WebAPI Apache Tom-  
cat CDM

ATLAS

<sup>1</sup><http://www.ohdsi.org/web/atlas>

The screenshot shows the ATLAS web application interface. On the left is a dark sidebar with navigation links: Home, Data Sources, Search, Concept Sets, Cohort Definitions (highlighted), Characterizations, Cohort Pathways, Incidence Rates, Profiles, Estimation, Prediction, Jobs, Configuration, and Feedback. Below the sidebar, it says "Apache 2.0 open source software" and "provided by OHDSI join the journey". The main content area has a header "Cohort #1770710" and a sub-header "New users of ACE inhibitors as first-line monotherapy for hypertension". Below this are tabs for Definition, Concept Sets, Generation, Reporting, Export, and Messages (with 3 notifications). A search bar says "enter a cohort definition description here". Under "Cohort Entry Events", there's a section for "Events having any of the following criteria:" with a dropdown set to "ACE inhibitors" and a button "+ Add attribute...". Below it is a red "Delete Criteria" button. A note says "with continuous observation of at least [365] days before and [0] days after event index date" and "Limit initial events to: earliest event per person". A green "Restrict initial events" button is at the bottom. Under "Inclusion Criteria", there's a "New inclusion criteria" section with two items: "1. has hypertension diagnosis in 1 yr prior to treatment" and "2. Has no prior antihypertensive drug exposures in medical".

Figure 9.3: ATLAS

ATLAS OMOP  
 ATLAS WebAPI  
 ATLAS Shiro  
 ATLAS WebAPI<sup>2</sup>

### 9.3.1

ATLAS WebAPI Shiro WebAPI

<sup>2</sup><https://github.com/OHDSI/WebAPI/wiki/Security-Configuration>

### 9.3.2

ATLAS  ATLAS GitHub  wiki  

### 9.3.3

ATLAS OHDSI WebAPI GitHub<sup>4</sup> WebAPI GitHub

9.4

OHDSI  9.4  R 

10 □ Chapter 12 □ Chapter 13 □ Chapter 14 □ CDM

9.4.1

Chapter  
19

942



---

<sup>3</sup><https://github.com/OHDSI/ATLAS/wiki>

<sup>4</sup><https://github.com/OHDSI/Atlas/wiki/Atlas-Setup-Guide>

Prediction and estimation methods	<p><b>Cohort Method</b> New-user cohort studies using large-scale regression for propensity and outcome models</p>	<p><b>Self-Controlled Case Series</b> Self-Controlled Case Series analysis using few or many predictors, includes splines for age and seasonality.</p>	<p><b>Self-Controlled Cohort</b> A self-controlled cohort design, where time preceding exposure is used as control.</p>
	<p><b>Patient Level Prediction</b> Build and evaluate predictive models for user-specified outcomes, using a wide array of machine learning algorithms.</p>	<p><b>Case-control</b> Case-control studies, matching controls on age, gender, provider, and visit date. Allows nesting of the study in another cohort.</p>	<p><b>Case-crossover</b> Case-crossover design including the option to adjust for time-trends in exposures (so-called case-time-control).</p>
Method characterization	<p><b>Empirical Calibration</b> Use negative control exposure-outcome pairs to profile and calibrate a particular analysis design.</p>	<p><b>Method Evaluation</b> Use real data and established reference sets as well as simulations injected in real data to evaluate the performance of methods.</p>	<p><b>Evidence Synthesis</b> Combining study diagnostics and results across multiple sites.</p>
Supporting packages	<p><b>Database Connector</b> Connect directly to a wide range of database platforms, including SQL Server, Oracle, and PostgreSQL.</p>	<p><b>Sql Render</b> Generate SQL on the fly for the various SQL dialects.</p>	<p><b>Cyclops</b> Highly efficient implementation of regularized logistic, Poisson and Cox regression.</p>
	<p><b>ParallelLogger</b> Support for parallel computation with logging to console, disk, or e-mail.</p>	<p><b>Feature Extraction</b> Automatically extract large sets of features for user-specified cohorts using data in the CDM.</p>	

Figure 9.4: OHDSI □□□□□□□□□□□□□□□□□

3. OHDSI Methods Library [Cyclops](https://ohdsi.github.io/MethodsLibrary)

### 9.4.3

R Library<sup>5</sup> GitHub CRAN R Library GitHub CRAN

### 9.4.4

CDM SQL Server Oracle Microsoft APS IBM Netezza Amazon Redshift Impala Hadoop Google Big-Query

Server R Library RStudio Java

### 9.4.5

OHDSI R

1. R
2. Rtools Windows R
3. RStudio R IDE

## 9.5

ATLAS Methods Library OHDSI Web Services (AWS)

### 9.5.1 Broadsea

Broadsea<sup>7</sup> Docker [dockerUrl] OHDSI Docker Windows MacOS Linux Broadsea

<sup>5</sup><https://ohdsi.github.io/MethodsLibrary>

<sup>6</sup><https://ohdsi.github.io/CohortMethod/articles/MultipleAnalyses.html>

<sup>7</sup><https://github.com/OHDSI/Broadsea>

Docker Methods Library ATLAS OHDSI

## 9.5.2 Amazon AWS

Amazon AWS in-a-Box<sup>8</sup> OHDSI on AWS [^ohdsiOnAwsUrl]

OHDSI-in-a-Box □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□ OHDSI □□□□□□□□□□□□□□□□□□□□□  
CDM □□□□□□□□□□□□□ ETL □□□ OHDSI-in-a-Box □□□□□□□□□□□□□ OHDSI-in-a-  
Box □□□□□□□□□□□ 9.5 □□□□□□□□□

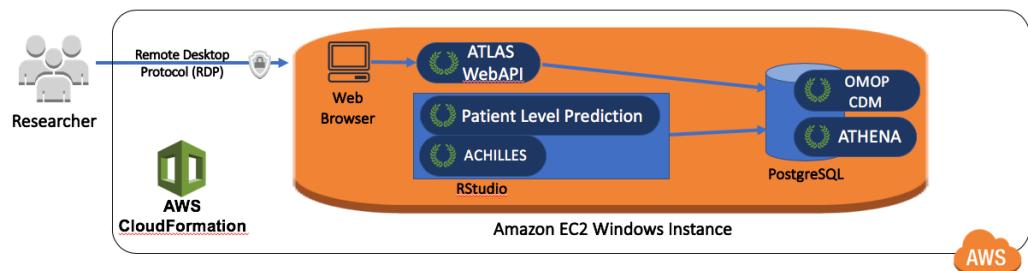


Figure 9.5: OHDSI-in-a-Box □ Amazon Web Services □ □ □ □ □ □ □

OHDSIonAWS  
Redshift  
Server

RStudio  
OHDSI Methods Library

OHDSI  
ATLAS

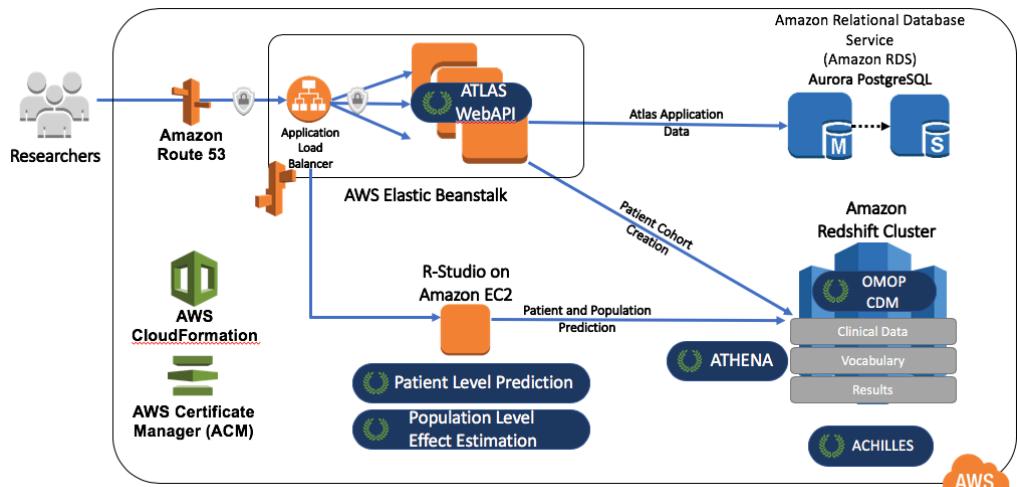


Figure 9.6: OHDSI on AWS □ Amazon Web Services □ □ □ □ □ □ □ □

<sup>8</sup><https://github.com/OHDSI/OHDSI-in-a-Box>

## 9.6



- CDM
  - \* OHDSI Methods Library R
  - \* OHDSI Methods Library ATLAS
- OHDSI
  - \* OHDSI
  - \* OHDSI
  - \* OHDSI
- OHDSI OHDSI
  - \* OHDSI ATLAS
  - \* OHDSI Methods Library R
- OHDSI OHDSI OHDSI OHDSI

# Chapter 10

# — — — SQL R

□□□□□□□: Martijn Schuemie & Peter Rijnbeek

CDM  
SQL Server  
Library  
OHDSI  
SQL  
Structured  
Query  
Language  
SQL  
Server  
PERSON  
10

```
SELECT TOP 10 * FROM person;
```

PostgreSQL

```
SELECT * FROM person LIMIT 10;
```

OHDSI - OHDSI SQL - SQL Server SQL - SQL OHDSI  
SQL

SOL

CPM QHPSI

SQL  
SqlRender  
Database  
SOL  
CDM  
OHDSI

## 10.1 SqIRender

SqlRender  CRAN Comprehensive R Archive Network

```
install.packages("SqlRender")
```

### 10.1.1 SQL

# SQL

a render

```
sql <- "SELECT * FROM concept WHERE concept_id = @a;"  
render(sql, a = 123)
```

```
## [1] "SELECT * FROM concept WHERE concept_id = 123;"
```

```
sql <- "SELECT * FROM @x WHERE person_id = @a;"  
render(sql, x = "observation", a = 123)
```

```
## [1] "SELECT * FROM observation WHERE person_id = 123;"
```

```
sql <- "SELECT * FROM concept WHERE concept_id IN (@a);"
render(sql, a = c(123, 234, 345))
```

```
## [1] "SELECT * FROM concept WHERE concept_id IN (123,234,345);"
```

### If-Then-Else

```
{Condition} ? {if true} : {if false}    condition  
true 1 iftrue iffase
```

```
sql <- "SELECT * FROM cohort {@x} ? {WHERE subject_id = 1}"
render(sql, x = FALSE)
```

```
## [1] "SELECT * FROM cohort "
```

```
render(sql, x = TRUE)
```

```
## [1] "SELECT * FROM cohort WHERE subject_id = 1"
```

```
□□□□□□□□□□□□□□□□□
```

```
sql <- "SELECT * FROM cohort {@x == 1} ? {WHERE subject_id = 1};"
render(sql, x = 1)
```

```
## [1] "SELECT * FROM cohort WHERE subject_id = 1;"
```

```
render(sql, x = 2)
```

```
## [1] "SELECT * FROM cohort ;"
```

```
IN □□□□□□□□□□□□□□□
```

```
sql <- "SELECT * FROM cohort {@x IN (1,2,3)} ? {WHERE subject_id = 1};"
render(sql, x = 2)
```

```
## [1] "SELECT * FROM cohort WHERE subject_id = 1;"
```

## 10.1.2 SQL

SqlRender □□□□□□□□□□□□□□□ OHDSI SQL □□□□ SQL □□□□□□□□□□□□□□

```
sql <- "SELECT TOP 10 * FROM person;"
translate(sql, targetDialect = "postgresql")
```

```
## [1] "SELECT * FROM person LIMIT 10;"
## attr("sqlDialect")
## [1] "postgresql"
```

```
targetDialect □□□□□□□□□□□□□□□ oracle, postgresql, pdw, redshift, impala, netezza, bigquery, sqlite, sql server □
```

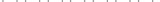
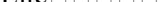


□□□□SQL□□□□  
SQL□□□□□SQL□□□  
Server□□□□□□□□□

OHDSI  
SQL

supported by translate

ABS	EXP	RAND
ACOS	FLOOR	RANK
ASIN	GETDATE	RIGHT
ATAN	HASHBYTES*	ROUND
Avg	ISNULL	ROW_NUMBER
CAST	ISNUMERIC	RTRIM
CEILING	LEFT	SIN
CHARINDEX	LEN	SQRT
CONCAT	LOG	SQUARE
COS	LOG10	STDEV
COUNT	LOWER	SUM
COUNT_BIG	LTRIM	TAN
DATEADD	MAX	UPPER
DATEDIFF	MIN	VAR
DATEFROMPARTS	MONTH	YEAR
DATETIMEFROMPARTS	NEWID	
DAY	PI	
EOMONTH	POWER	

\* Oracle  SQLite 

## SQL

```
-- Simple selects:  
SELECT * FROM table;  
  
-- Selects with joins:  
SELECT * FROM table_1 INNER JOIN table_2 ON a = b;  
  
-- Nested queries:
```

```
SELECT * FROM (SELECT * FROM table_1) tmp WHERE a = b;

-- Limiting to top rows:
SELECT TOP 10 * FROM table;

-- Selecting into a new table:
SELECT * INTO new_table FROM table;

-- Creating tables:
CREATE TABLE table (field INT);

-- Inserting verbatim values:
INSERT INTO other_table (field_1) VALUES (1);

-- Inserting from SELECT:
INSERT INTO other_table (field_1) SELECT value FROM table;

-- Simple drop commands:
DROP TABLE table;

-- Drop table if it exists:
IF OBJECT_ID('ACHILLES_analysis', 'U') IS NOT NULL
    DROP TABLE ACHILLES_analysis;

-- Drop temp table if it exists:
IF OBJECT_ID('tempdb..#cohorts', 'U') IS NOT NULL
    DROP TABLE #cohorts;

-- Common table expressions:
WITH cte AS (SELECT * FROM table) SELECT * FROM cte;

-- OVER clauses:
SELECT ROW_NUMBER() OVER (PARTITION BY a ORDER BY b)
    AS "Row Number" FROM table;

-- CASE WHEN clauses:
SELECT CASE WHEN a=1 THEN a ELSE 0 END AS value FROM table;

-- UNIONs:
SELECT * FROM a UNION SELECT * FROM b;

-- INTERSECTIONS:
SELECT * FROM a INTERSECT SELECT * FROM b;

-- EXCEPT:
SELECT * FROM a EXCEPT SELECT * FROM b;
```

□ □ □ □ □ □

```
SQL Server SELECT first_name + ' ' + last_name AS full_name FROM table
PostgreSQL   Oracle   SELECT first_name || ' ' || last_name AS full_name FROM table
SqlRender    SELECT first_name + last_name AS full_name FROM table
SqlRender    last_name + CAST(age AS VARCHAR(3)) AS full_name FROM table
```

□ □ □ □ □ □ □ □ □ □ AS □ □ □ □ □

SQL AS Server PostgreSQL Redshift

```
-- Using AS keyword
SELECT *
FROM my_table AS table_1
INNER JOIN (
    SELECT * FROM other_table
) AS table_2
ON table_1.person_id = table_2.person_id;

-- Not using AS keyword
SELECT *
FROM my_table table_1
INNER JOIN (
    SELECT * FROM other_table
) table_2
ON table_1.person_id = table_2.person_id;
```

Oracle AS SqlRend AS

6 / 6



10 of 10

```
sql <- "SELECT * FROM #children;"  
translate(sql, targetDialect = "oracle", oracleTempSchema = "temp_schema")
```

```
## Warning: The 'oracleTempSchema' argument is deprecated. Use 'tempEmulation'
## This warning is displayed once every 8 hours.

## [1] "SELECT * FROM temp_schema.ui6r639fchildren ;"
## attr(,"sqlDialect")
## [1] "oracle"

temp_schema Oracle30Oracle22
Oracle Oracle
TRUNCATE DROP Oracle

SQL Server SQL Server
```

```
CREATE TABLE #temp (txt VARCHAR);

INSERT INTO #temp
SELECT '1';

SELECT * FROM #temp WHERE txt = 1;
```

```
SELECT * FROM #temp WHERE txt = CAST(1 AS VARCHAR);
```

1

```
SELECT * FROM #temp WHERE CAST(txt AS INT) = 1;
```

A horizontal row of fifteen small, uniform white squares, likely representing a sequence or a set of data points.

## SQL Server DBMS

```
SELECT * FROM concept WHERE concept_class_id = 'Clinical Finding'
```

A horizontal row of 20 empty square boxes, intended for students to write their answers in a handwriting practice exercise.

```
SELECT * FROM concept WHERE LOWER(concept_class_id) = 'clinical finding'
```

SQL Server cdm\_data.dbo.person  
Server 1.dbo  
PostgreSQL SQL Server

SQL Server @databaseSchema SQL

```
SELECT * FROM @databaseSchema.person
```

```
SQL Server databaseSchema  
= "cdm_data.dbo"  
= "cdm_data"
```

```
USE cdm_data.dbo; GO
```

# SQL

## SQL

SQL Shiny  
SqlRender

```
launchSqlRenderDeveloper()
```

```
“`{r sqlDeveloper, fig.cap=“The SqlDeveloper”}
```

## 10.2 DatabaseConnector

DatabaseConnector Java JDBC R Archive Network

```
install.packages("DatabaseConnector")
```

## 10.2.1

```
conn <- connect(dbms = "postgresql",
                 server = "localhost/postgres",
                 user = "joe",
                 password = "secret",
                 schema = "cdm")
```

```
## Connecting using PostgreSQL driver
```

`disconnect(conn)`

JDBC

```
connString <- "jdbc:postgresql://localhost:5432/postgres"  
conn <- connect(dbms = "postgresql",  
                 connectionString = connString,  
                 user = "joe",  
                 password = "secret",  
                 schema = "cdm")
```

```
## Connecting using PostgreSQL driver
```

```
details <- createConnectionDetails(dbms = "postgresql",
                                    server = "localhost/postgres",
                                    user = "joe",
                                    password = "secret",
                                    schema = "cdm")
conn <- connect(details)
```

```
## Connecting using PostgreSQL driver
```

## 10.2.2

querySql executeSql querySql

```
querySql(conn, "SELECT TOP 3 * FROM person")
```

```
##   person_id gender_concept_id year_of_birth
## 1           1             8507      1975
## 2           2             8507      1976
## 3           3             8507      1977
```

```
executeSql(conn, "TRUNCATE TABLE foo; DROP TABLE foo;")
```

### 10.2.3 ffdff

```
x <- querySql.ffdf(conn, "SELECT * FROM person")
```

x□□□ffdf□□□□□□□□□□

## 10.2.4 SQL

```
SqlRender → render → translate → renderTransl...
```

```
x <- renderTranslateQuerySql(conn,
                           sql = "SELECT TOP 10 * FROM @schema.person",
                           schema = "cdm_synput")
```

SQL Server TOP 10 PostgreSQL LIMIT 10 SQL @schema

## 10.2.5

executeSql SQL

```
data(mtcars)
insertTable(conn, "mtcars", mtcars, createTable = TRUE)
```

mtcars

10.3 CDM

OHDSI SOL CDM Q

A horizontal row of 24 small, identical rectangular boxes arranged in a single line.

```
SELECT COUNT(*) AS person_count FROM @cdm.person;
```

---

**PERSON\_COUNT**

---

A horizontal row of 20 empty square boxes, each with a thin black border, intended for children to practice writing their names.

```
SELECT AVG(DATEDIFF(DAY,
                      observation_period_start_date,
                      observation_period_end_date) / 365.25) AS num_years
FROM @cdm.observation_period;
```

---

**NUM\_YEARS**  
1.980803

```
SELECT MAX(YEAR(observation_period_end_date) -  
          year_of_birth) AS max_age  
FROM @cdm.person  
INNER JOIN @cdm.observation_period  
  ON person.person_id = observation_period.person_id;
```

---

**MAX\_AGE**

---

... AS “ages” ages < .50 \* n

```
WITH ages
AS (
    SELECT age,
           ROW_NUMBER() OVER (
               ORDER BY age
           ) order_nr
    FROM (
        SELECT YEAR(observation_period_start_date) - year_of_birth AS age
        FROM @cdm.person
```

```

    INNER JOIN @cdm.observation_period
        ON person.person_id = observation_period.person_id
    ) age_computed
)
SELECT MIN(age) AS min_age,
MIN(CASE
    WHEN order_nr < .25 * n
        THEN 9999
    ELSE age
    END) AS q25_age,
MIN(CASE
    WHEN order_nr < .50 * n
        THEN 9999
    ELSE age
    END) AS median_age,
MIN(CASE
    WHEN order_nr < .75 * n
        THEN 9999
    ELSE age
    END) AS q75_age,
MAX(age) AS max_age
FROM ages
CROSS JOIN (
    SELECT COUNT(*) AS n
    FROM ages
) population_size;

```

MIN_AGE	Q25_AGE	MEDIAN_AGE	Q75_AGE	MAX_AGE
0	6	17	34	90

```
sql <- "SELECT YEAR(observation_period_start_date) -  
        year_of_birth AS age  
FROM @cdm.person  
INNER JOIN @cdm.observation_period  
    ON person.person_id = observation_period.person_id;  
age <- renderTranslateQuerySql(conn, sql, cdm = "cdm")  
quantile(age[, 1], c(0, 0.25, 0.5, 0.75, 1))
```

##	0%	25%	50%	75%	100%
##	0	6	17	34	90

```
SELECT TOP 10 condition_source_value,
       COUNT(*) AS code_count
  FROM @cdm.condition_occurrence
 GROUP BY condition_source_value
 ORDER BY -COUNT(*);
```

CONDITION_SOURCE_VALUE	CODE_COUNT
4019	49094668
25000	36149139
78099	28908399
319	25798284
31401	22547122
317	22453999
311	19626574
496	19570098
I10	19453451
3180	18973883

□□□□□ CONDITION OCCURRENCE □□□□□ CONDITION SOURCE VALUE □□□□□

10.4

CDM SQL

```
SELECT COUNT(*) AS subject_count,
       concept_name
  FROM @cdm.person
 INNER JOIN @cdm.concept
    ON person.gender_concept_id = concept.concept_id
 GROUP BY concept.name;
```

SUBJECT_COUNT	CONCEPT_NAME
14927548	FEMALE
11371453	MALE

```
    ON drug_concept_id = descendant_concept_id
INNER JOIN @cdm.concept ingredient
    ON ancestor_concept_id = ingredient.concept_id
WHERE LOWER(ingredient.concept_name) = 'ibuprofen'
    AND ingredient.concept_class_id = 'Ingredient'
    AND ingredient.standard_concept = 'S';
```

PRESCRIPTION\_COUNT

---

26871214

## 10.5 QueryLibrary

QueryLibrary CDM SQL 1

Select a query	
Column visibility	Show 10 entries
Search:	<input type="text"/>
Group	Name
[ "drug exposure" ]	All
drug exposure	DEX01 Counts of persons with any number of exposures to a certain drug
drug exposure	DEX02 Counts of persons taking a drug, by age, gender, and year of exposure
drug exposure	DEX03 Distribution of age, stratified by drug
drug exposure	DEX04 Distribution of gender in persons taking a drug
drug exposure	DEX05 Counts of drug records for a particular drug
drug exposure	DEX06 Counts of distinct drugs in the database
drug exposure	DEX07 Maximum number of drug exposure events per person over some time period

## Query Description

### DEX01: Counts of persons with any number of exposures to a certain drug

### Description

This query is used to count the persons with at least one exposures to a certain drug (drug\_concept\_id). See vocabulary queries for obtaining valid drug\_concept\_id values. The input to the query is a value (or a comma-separated list of values) of a drug\_concept\_id. If the input is omitted, all drugs in the data table are summarized.

### Query

The following is a sample run of the query. The input parameters are highlighted in blue.

```
SELECT
    c.concept_name,
    drug_concept_id,
    COUNT(person_id) AS num_persons
FROM cdm.drug_exposure
INNER JOIN cdm.concept c
ON drug_concept_id = c.concept_id
WHERE drug_concept_id = 'Pain'
```

Figure 10.1: □□□□□□□□□ CDM □□□□□ SQL □□□□□□□□□

CDM

<sup>1</sup><http://data.ohdsi.org/QueryLibrary>

<sup>2</sup><https://github.com/OHDSI/QueryLibrary>

10.6

## 10.6.1

□□□□□□□□□□□□□□□□□ 1 □□□□□□□□□□□□□□□□□

## 10.6.2

10.6.3

10.6.4

```
library(DatabaseConnector)
conn <- connect(dbms = "postgresql",
                 server = "localhost/postgres",
                 user = "joe",
                 password = "secret")
cdmDbSchema <- "cdm"
cohortDbSchema <- "scratch"
cohortTable <- "my_cohorts"

sql <- "
CREATE TABLE @cohort_db_schema.@cohort_table (
  cohort_definition_id INT,
  cohort_start_date DATE,
  cohort_end_date DATE,
  subject_id BIGINT
);
"
renderTranslateExecuteSql(conn, sql,
                         cohort_db_schema = cohortDbSchema,
                         cohort_table = cohortTable)
```

10.6.5

□ □ □ □ □ □ □ □ □ □ □ □ COHORT □ □ □ □ □ □ □ □ □

CDM DRUG ERA DRUG EXPOSURE

## 10.6.6

□□□□□ CONDITION OCCURRENCE □□□□□ CONCEPT ANCESTOR □□□□□ □□□□

10.6.7

A horizontal row of 24 small white squares arranged in a single line.

```
sql <- "
WITH tar AS (
    SELECT concept_name AS gender,
           FLOOR((YEAR(cohort_start_date) -
                   year_of_birth) / 10) AS age,
           subject_id,
           cohort_start_date,
           CASE WHEN DATEADD(DAY, 7, cohort_start_date) >
                 observation period end date

```

```

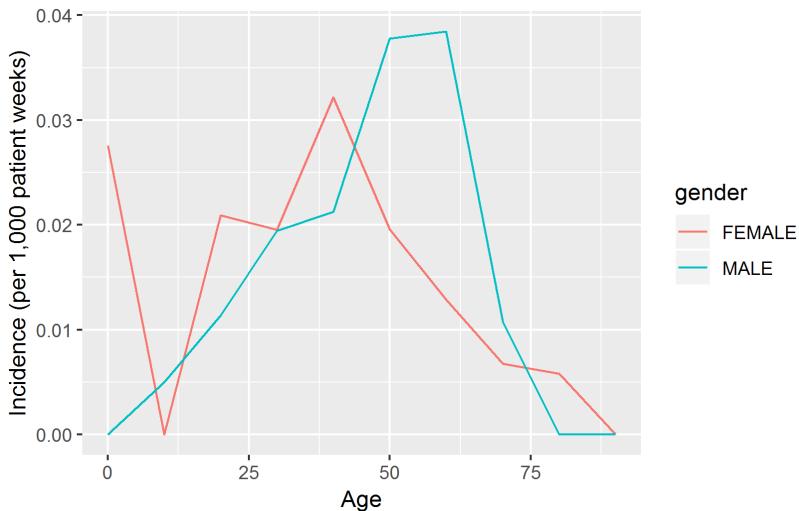
THEN observation_period_end_date
ELSE DATEADD(DAY, 7, cohort_start_date)
END AS cohort_end_date
FROM @cohort_db_schema.@cohort_table
INNER JOIN @cdm_db_schema.observation_period
    ON subject_id = observation_period.person_id
        AND observation_period_start_date < cohort_start_date
        AND observation_period_end_date > cohort_start_date
INNER JOIN @cdm_db_schema.person
    ON subject_id = person.person_id
INNER JOIN @cdm_db_schema.concept
    ON gender_concept_id = concept_id
WHERE cohort_definition_id = 1 --
)
SELECT days.gender,
    days.age,
    days,
    CASE WHEN events IS NULL THEN 0 ELSE events END AS events
FROM (
    SELECT gender,
        age,
        SUM(DATEDIFF(DAY, cohort_start_date,
            cohort_end_date)) AS days
    FROM tar
    GROUP BY gender,
        age
) days
LEFT JOIN (
    SELECT gender,
        age,
        COUNT(*) AS events
    FROM tar
    INNER JOIN @cohort_db_schema.@cohort_table angioedema
        ON tar.subject_id = angioedema.subject_id
            AND tar.cohort_start_date <= angioedema.cohort_start_date
            AND tar.cohort_end_date >= angioedema.cohort_start_date
    WHERE cohort_definition_id = 2 --
    GROUP BY gender,
        age
) events
ON days.gender = events.gender
    AND days.age = events.age;
"
results <- renderTranslateQuerySql(conn, sql,
    cohort_db_schema = cohortDbSchema,
    cohort_table = cohortTable,
    cdm_db_schema = cdmDbSchema,
    snakeCaseToCamelCase = TRUE)

```

```
#      IR
results$ir <- 1000 * results$events / results$days / 7

#
results$age <- results$age * 10

library(ggplot2)
ggplot(results, aes(x = age, y = ir, group = gender, color = gender)) +
  geom_line() +
  xlab(" ") +
  ylab(" 1,000      ")
```



## 10.6.8

A horizontal row of 30 empty square boxes, intended for students to write their answers in a grid format.

```
sql <- "
TRUNCATE TABLE @cohort_db_schema.@cohort_table;
DROP TABLE @cohort_db_schema.@cohort_table;
"
renderTranslateExecuteSql(conn, sql,
                         cohort_db_schema = cohortDbSchema,
                         cohort_table = cohortTable)

disconnect(conn)
```

10.6.9

10.7



- **SQL** Structured Query Language Common Data Model CDM
  - SQL Database Connector CDM
  - **SqlRender** Database Connector CDM
  - R SQL OHDSI
  - **QueryLibrary** CDM SQL

10.8

10

```
install.packages(c("SqlRender", "DatabaseConnector", "remotes"))
remotes::install_github("ohdsi/Eunomia", ref = "v1.0.0")
```

Eunomia CDM R

```
connectionDetails <- Eunomia::getEunomiaConnectionDetails()
```

CDM [REDACTED] main [REDACTED]

10.1 SQL R

□□□□□□□□□□ ?? □□□□□□

# Chapter 11

□□□□□: *Kristin Kostka!*

ATLAS SQL

11.1

OHDSI






A horizontal row of 20 empty rectangular boxes, likely used for grading or marking student responses.



Page 10 of 10

- CDM
  - 
  - DRUG\_EXPOSURE DAYS\_SUPPORT
  - 365

11.1



**Figure 11.1:** □□□□□□□□□□□

A horizontal row of 20 empty square boxes, likely used for grading student responses.



11.2

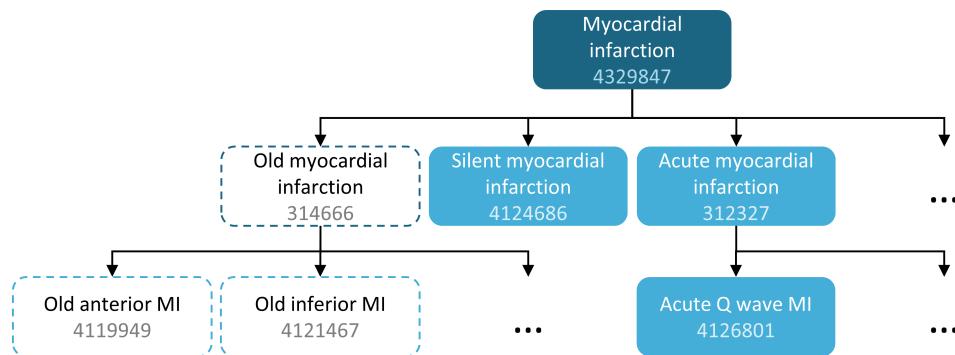


## □□□#tab:conceptSetExpression□□□□□□□□□□□

ID					
4329847	□□□□	□□□	□□	□□□□□□□□	
314666	□□□□□□□	□□	□□	□□□	

# □□□□□□□□□ 11.2 □□□□□

## 9□□□ ICD-10 □□□□□□□□□□



11.3

<sup>1</sup><https://github.com/OHDSI/Aphrodite> ## □□□□□□□□□□

## Chapter 17

### 11.3.1 OHDSI

OHDSI APHRODITE (Banda et al., 2017) PheValuator (Swerdel et al., 2019) OHDSI eMERGE Phenotype Library (Hripcak et al., 2019)

## 11.4

11.3

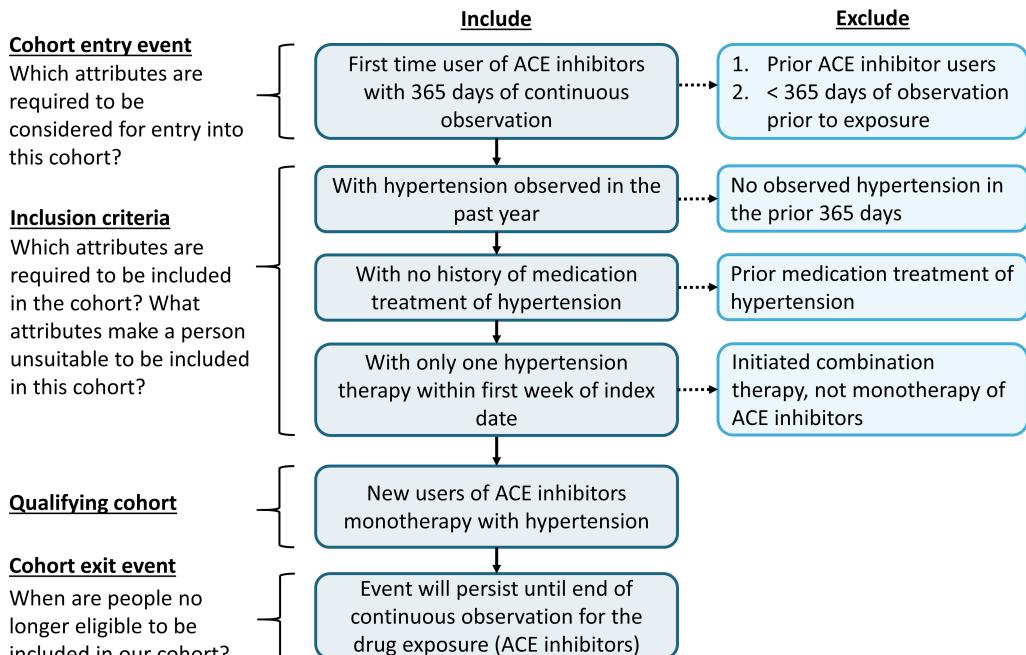


Figure 11.3:

ATLAS CDM  
## ATLAS

ATLAS Cohort Definitions cohort

<sup>2</sup><https://www.ohdsi.org/web/wiki/doku.php?id=projects:workgroups:gold-library-wg>



## New Cohort Definition

New Cohort Definition Save  Cancel 

- [Definition !\[\]\(dae4756483289d49145c6870d7a99152\_img.jpg\)](#)
- [Concept Sets](#)
- [Generation](#)
- [Reporting](#)
- [Export](#)

enter a cohort definition description here

### Cohort Entry Events

Events having any of the following criteria:

with continuous observation of at least  days before and  days after event index date

Limit initial events to:  per person.

[+ Add Initial Event !\[\]\(8fbd992114ef07ab2d4fee7edda363ab\_img.jpg\)](#)

[Restrict initial events](#)

### Inclusion Criteria

[New inclusion criteria](#)

Limit qualifying events to:  per person.

Figure 11.4: □□□□□□□□□□

#### New Cohort Definition



ATLAS

A horizontal row of 24 small, empty square boxes for writing answers, followed by a green box containing a blue floppy disk icon.

## 11.4.1

## 11.4.2

ACE

**1:** □□□□□□□□□□□□□□□□□□□□□□□□□□

set New Concept Set Un  
Concept Set ACE

**Cohort #1771427**

EXAMPLE: new users of ACE inhibitors as first-line mono-therapy for hypertension

Definition Concept Sets Generation Reporting Export

enter a cohort definition description here

**Cohort Entry Events**

Events having any of the following criteria:

with continuous observation of at least  days before and  days after event index date.

Limit initial events to: earliest event per person.

**+ Add Initial Event**

**Restrict initial events**

**Inclusion Criteria**

**New inclusion criteria**

Limit qualifying events to: earliest event per person.

**Cohort Exit**

Add Condition Era  
Find patients with specific diagnosis era.

Add Condition Occurrence  
Find patients with specific diagnoses.

Add Death  
Find patients based on death.

Add Device Exposure  
Find patients based on device exposure.

Add Dose Era  
Find patients with dose eras.

Add Drug Era  
Find patients with exposure to drugs over time.

Add Drug Exposure  
Find patients with exposure to specific drugs or drug classes.

Figure 11.5: □□□□□□□□□

**Cohort Entry Events**

Events having any of the following criteria:

a drug exposure of **Any Drug**

**+ Add attribute...**  **Delete Criteria**

with continuous observation of at least  days before and  days after event index date.

Limit initial events to: earliest event per person.

**Restrict initial events**

Figure 11.6: □□□□□□□

Figure 11.7: □□□□□ - ACE□□□

11.7

11.8 ACE

ATLAS Import Concept Set  
inhibitors pt Any  
Drug

### 11.4.3

ACE attribute Add first exposure criteria

**11.10**



ATLASXNo

event 11.10

Concept Set Expression   Included Concepts 21536   Included Source Codes   Export   Import

Name: ACE Inhibitors

Show 25 ▾ entries   Search:

Showing 1 to 15 of 15 entries   Previous 1 Next

	Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	Exclude	Descendants	Mapped
1335471	18867	benazepril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
1340128	1998	Captopril	Drug	Standard	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
19050216	21102	Cilazapril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
1341927	3827	Enalapril	Drug	Standard	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
1342001	3829	Enalaprilat	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
1363749	50166	Fosinopril	Drug	Standard	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
19122327	60245	imidapril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
1308216	29046	Lisinopril	Drug	Standard	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
1310756	30131	moexipril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
1373225	54552	Perindopril	Drug	Standard	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
1331235	35208	quinapril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
1334456	35296	Ramipril	Drug	Standard	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
19040051	36908	spirapril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
1342439	38454	trandolapril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	
19102107	39990	zofenopril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	

■ Classification   ■ Non-Standard   ■ Standard

Figure 11.8: ACE

Import Concept Set From Repository...

New Concept Set

Show 10 ▾ entries   Filter Repository Concept Sets: ace inhibitors

ID	Title	Created	Modified	Author
1794480	[OHDSI EU 2019] Excluded concepts of ACE inhibitors or Thiazide diuretics	03/28/2019 11:04 AM	03/28/2019 11:04 AM	anonymous
963	ACE Inhibitors			anonymous
3268	COPY OF: ACE Inhibitors			anonymous
99283	Ace Inhibitors			anonymous
142965	PheKB ACE-I ACE inhibitors			anonymous

Showing 1 to 5 of 5 entries (filtered from 11,667 total entries)   Previous 1 Next

Figure 11.9: ATLAS

Cohort Entry Events

Events having any of the following criteria:

a drug exposure of ACE inhibitors

for the first time in the person's history

with continuous observation of at least  days before and  days after event index date  
 Limit initial events to:  per person.

Figure 11.10: □□□□□□□□□□□□□□□□□□□□□□□□□□

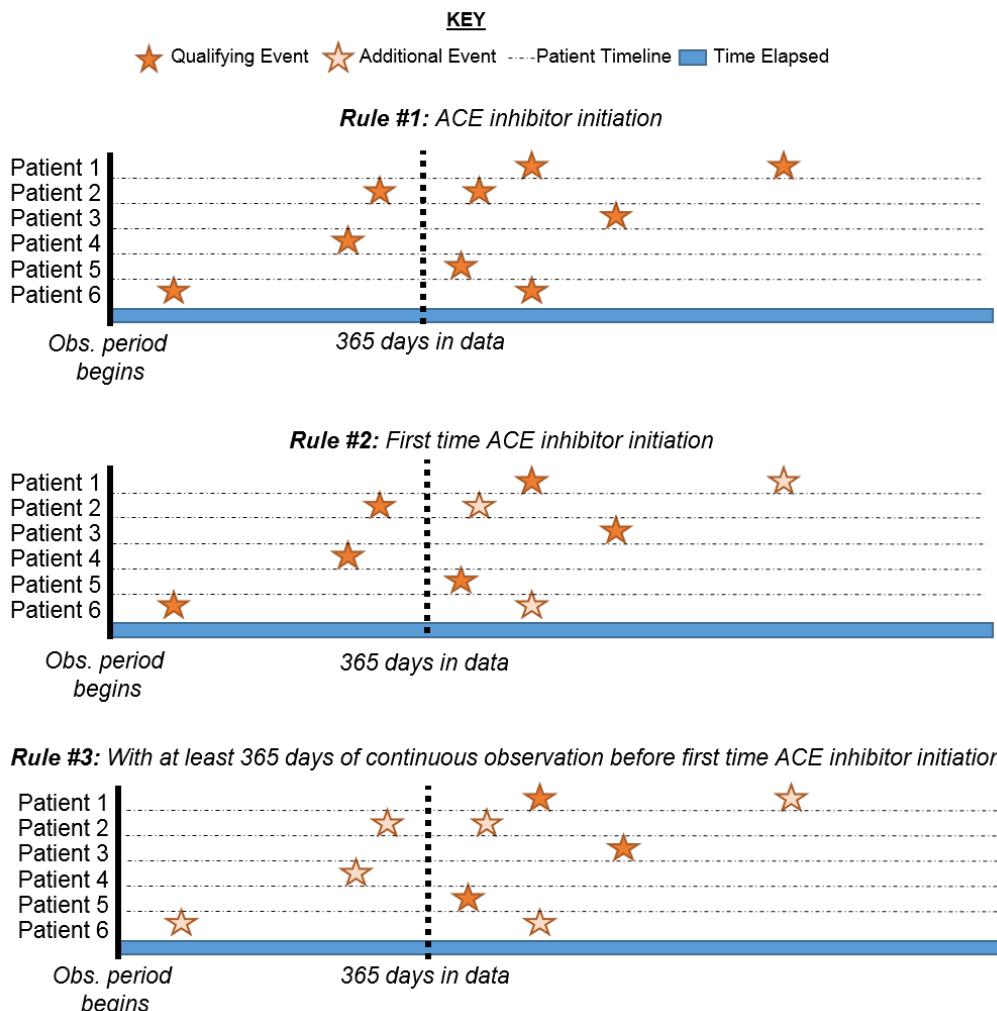


Figure 11.11: □□□□□□□□□□□□□□□□□□□□

#### 11.4.4

New inclusion criteria  
inclusion criteria

+Add criteria to group  
Initial Event  
criteria to group  
condition occ ## SQL

SQL R 10 OHDSI

## 11.4.5

```
library(CohortMethod)
connDetails <- createConnectionDetails(dbms = "postgresql",
                                         server = "localhost/ohdsi",
                                         user = "joe",
                                         password = "supersecret")

cdmDbSchema <- "my_cdm_data"
cohortDbSchema <- "scratch"
cohortTable <- "my_cohorts"
```

1146

□□□□□□□□□□□□□□□□□□ID□R□□□□□□□□□□□□□□□□□□SQL□□□□□□□□

```
aceI <- c(1308216, 1310756, 1331235, 1334456, 1335471, 1340128, 1341927,  
        1342439, 1363749, 1373225)  
  
hypertension <- 316866  
  
allHtDrugs <- c(904542, 907013, 932745, 942350, 956874, 970250, 974166,
```

```
978555, 991382, 1305447, 1307046, 1307863, 1308216,
1308842, 1309068, 1309799, 1310756, 1313200, 1314002,
1314577, 1317640, 1317967, 1318137, 1318853, 1319880,
1319998, 1322081, 1326012, 1327978, 1328165, 1331235,
1332418, 1334456, 1335471, 1338005, 1340128, 1341238,
1341927, 1342439, 1344965, 1345858, 1346686, 1346823,
1347384, 1350489, 1351557, 1353766, 1353776, 1363053,
1363749, 1367500, 1373225, 1373928, 1386957, 1395058,
1398937, 40226742, 40235485)
```

## 11.4.7

□□□□□□ACE□□□□□□□□□□□□

```
conn <- connect(connDetails)

sql <- "SELECT person_id AS subject_id,
    MIN(drug_exposure_start_date) AS cohort_start_date
INTO #first_use
FROM @cdm_db_schema.drug_exposure
INNER JOIN @cdm_db_schema.concept_ancestor
    ON descendant_concept_id = drug_concept_id
WHERE ancestor_concept_id IN (@ace_i)
GROUP BY person_id;"

renderTranslateExecuteSql(conn,
    sql,
    cdm_db_schema = cdmDbSchema,
    ace_i = aceI)
```

DRUG\_EXPOSURE□□□□□CONCEPT\_ANCESTOR□□□□□□□□□□□ACE□□□□□

## 11.4.8 365

□□□OBSERVATION\_PERIOD□□□□□□□□365□□□□□□□□□□□□□

```
sql <- "SELECT subject_id,
    cohort_start_date
INTO #has_prior_obs
FROM #first_use
INNER JOIN @cdm_db_schema.observation_period
    ON subject_id = person_id
        AND observation_period_start_date <= cohort_start_date
        AND observation_period_end_date >= cohort_start_date
WHERE DATEADD(DAY, 365, observation_period_start_date) < cohort_start_date;"
```

```
renderTranslateExecuteSql(conn, sql, cdm_db_schema = cdmDbSchema)
```

11.4.9

**365**□□□□□□□□□□□□□□□□□□□□□□

```
sql <- "SELECT DISTINCT subject_id,
  cohort_start_date
INTO #has_ht
FROM #has_prior_obs
INNER JOIN @cdm_db_schema.condition_occurrence
  ON subject_id = person_id
    AND condition_start_date <= cohort_start_date
    AND condition_start_date >= DATEADD(DAY, -365, cohort_start_date)
INNER JOIN @cdm_db_schema.concept_ancestor
  ON descendant_concept_id = condition_concept_id
WHERE ancestor_concept_id = @hypertension;"
```

```
renderTranslateExecuteSql(conn,
  sql,
  cdm_db_schema = cdmDbSchema,
  hypertension = hypertension)
```

```
SELECT  
DISTINCT
```

#### 11.4.10

A horizontal row of 20 empty square boxes, intended for students to write their answers in a handwriting practice exercise.

```
sql <- "SELECT subject_id,
  cohort_start_date
INTO #no_prior_ht_drugs
FROM #has_ht
LEFT JOIN (
  SELECT *
  FROM @cdm_db_schema.drug_exposure
  INNER JOIN @cdm_db_schema.concept_ancestor
    ON descendant_concept_id = drug_concept_id
  WHERE ancestor_concept_id IN (@all_ht_drugs)
) ht_drugs
  ON subject_id = person_id
  AND drug_exposure_start_date < cohort_start_date
WHERE person_id IS NULL;"
```

```
renderTranslateExecuteSql(conn,
```

```
sql,  
cdm_db_schema = cdmDbSchema,  
all_ht_drugs = allHtDrugs)
```

## 11.4.11

```

sql <- "SELECT subject_id,
  cohort_start_date
INTO #monotherapy
FROM #no_prior_ht_drugs
INNER JOIN @cdm_db_schema.drug_exposure
  ON subject_id = person_id
    AND drug_exposure_start_date >= cohort_start_date
    AND drug_exposure_start_date <= DATEADD(DAY, 7, cohort_start_date)
INNER JOIN @cdm_db_schema.concept_ancestor
  ON descendant_concept_id = drug_concept_id
WHERE ancestor_concept_id IN (@all_ht_drugs)
GROUP BY subject_id,
  cohort_start_date
HAVING COUNT(*) = 1;"

renderTranslateExecuteSql(conn,
  sql,
  cdm_db_schema = cdmDbSchema,
  all_ht_drugs = allHtDrugs)

```

11.4.12

```
sql <- "
SELECT person_id,
       CAST(1 AS INT) AS concept_id,
       drug_exposure_start_date AS exposure_start_date,
       drug_exposure_end_date AS exposure_end_date
  INTO #exposure
  FROM @cdm_db_schema.drug_exposure
  INNER JOIN @cdm_db_schema.concept_ancestor
    ON descendant_concept_id = drug_concept_id
   WHERE ancestor_concept_id IN (@ace_i);"
renderTranslateExecuteSql(conn,
                         sql,
```

```
cdm_db_schema = cdmDbSchema,
ace_i = aceI)
```

```
□□□□□□□□□□□□□□□□□□□□□□□□□
```

```
sql <- "
SELECT ends.person_id AS subject_id,
    ends.concept_id AS cohort_definition_id,
    MIN(exposure_start_date) AS cohort_start_date,
    ends.era_end_date AS cohort_end_date
INTO #exposure_era
FROM (
    SELECT exposure.person_id,
        exposure.concept_id,
        exposure.exposure_start_date,
        MIN(events.end_date) AS era_end_date
    FROM #exposure exposure
    JOIN (
--cteEndDates
        SELECT person_id,
            concept_id,
            DATEADD(DAY, - 1 * @max_gap, event_date) AS end_date
        FROM (
            SELECT person_id,
                concept_id,
                event_date,
                event_type,
                MAX(start_ordinal) OVER (
                    PARTITION BY person_id ,concept_id ORDER BY event_date,
                    event_type ROWS UNBOUNDED PRECEDING
                ) AS start_ordinal,
                ROW_NUMBER() OVER (
                    PARTITION BY person_id, concept_id ORDER BY event_date,
                    event_type
                ) AS overall_ord
            FROM (
-- select the start dates, assigning a row number to each
            SELECT person_id,
                concept_id,
                exposure_start_date AS event_date,
                0 AS event_type,
                ROW_NUMBER() OVER (
                    PARTITION BY person_id, concept_id ORDER BY exposure_start_date
                ) AS start_ordinal
            FROM #exposure exposure

            UNION ALL
-- add the end dates with NULL as the row number, padding the end dates by
```



11.4.13

```
sql <- "TRUNCATE TABLE #first_use;  
DROP TABLE #first_use;  
  
TRUNCATE TABLE #has_prior_obs;  
DROP TABLE #has_prior_obs;  
  
TRUNCATE TABLE #has_ht;  
DROP TABLE #has_ht;  
  
TRUNCATE TABLE #no_prior_ht_drugs;  
DROP TABLE #no_prior_ht_drugs;  
  
TRUNCATE TABLE #monotherapy;  
DROP TABLE #monotherapy;  
  
TRUNCATE TABLE #exposure;  
DROP TABLE #exposure;  
  
TRUNCATE TABLE #exposure_era;  
DROP TABLE #exposure_era;"  
  
renderTranslateExecuteSql(conn, sq  
  
disconnect(conn)
```

11.5






116

100

ATLAS <http://atlas-demo.ohdsi.org>

11.1. ATLAS

- □□□□□□□□□□□□□□□
  - 16□□□
  - □□□□□□□□□□□365□□□□□□□□□□□□□□□
  - □□□NSAID□□□□□□□□□□□□□□□□□□□□□□□□□
  - □□□□□□□□□□□□□
  - □□□□□□□□□□□□□30□□□□□□□□□□□□□□□□□□□□□□□□□

□ □ □ □

R R-Studio Java  
9.4.5 SqlRender DatabaseConnector Eunomia

```
install.packages(c("SqlRender", "DatabaseConnector", "remotes"))
remotes::install_github("ohdsi/Eunomia", ref = "v1.0.0")
```

Eunomia□□□□□□CDM□□□□□R□

```
connectionDetails <- Eunomia::getEunomiaConnectionDetails()
```

# CDM main

□□ 11.2. SQL□□ R□□□□□□□ COHORT □□□□□□□□□ AMI □□□□□□□□



□□□□□□□□□ Appendix E.5 □□□□□□□□□



# Chapter 12

□□□□: *Anthony Sena & Daniel Prieto-Alhambra*



12.1



16

12.2

the Reporting of Observation Studies in Epidemiology STROBE (

Elm et al., 2008, )

12.3

et al. (2016) OHDSI 2

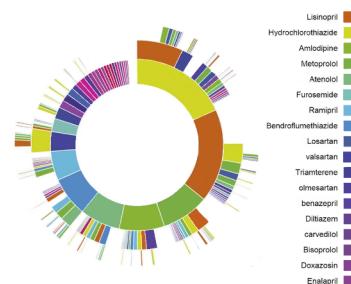
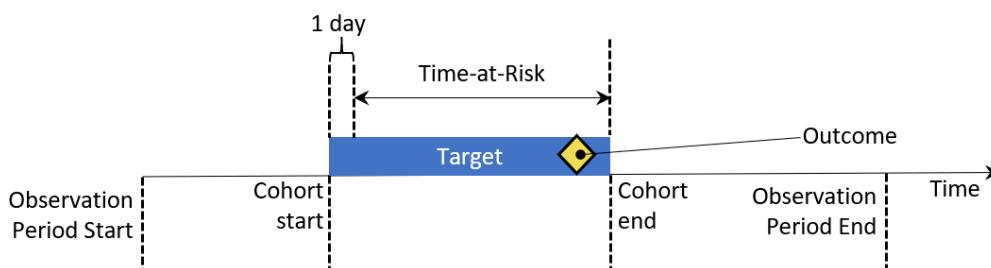


Figure 12.1: OHDSI

124



= #

= #

**PUS** 

12.5

WHO (Who, 2013, )

WHO

## 12.6 ATLAS

 Data Sources  ATLAS

## Occurrence

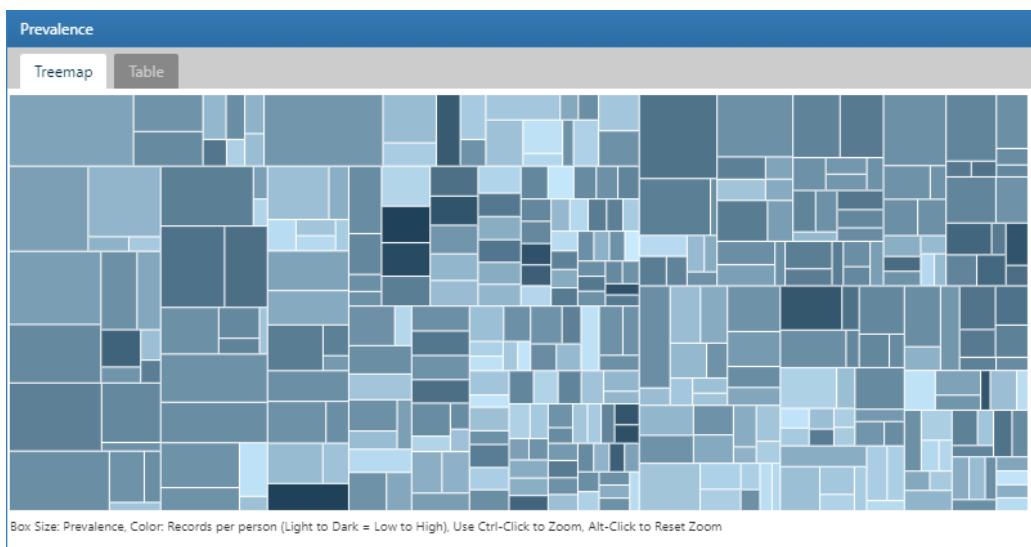


Figure 12.3: Atlas Data Sources: Condition Occurrence Treemap

Prevalence				
	Treemap	Table		
	Column visibility	Copy	CSV	Show 15 ▾ entries
				Filter: hypertension
Showing 1 to 15 of 47 entries (filtered from 15,907 total entries)				Previous 1 2 3 4 Next
Concept	Id	Name	Person Count	Records per person
320128		Essential hypertension	17,814,076	12.30% 5.80
312648		Benign essential hypertension	11,014,877	7.61% 4.35
317898		Malignant essential hypertension	1,021,441	0.70% 2.22
381290		Ocular hypertension	521,264	0.36% 2.40
441922		Transient hypertension of pregnancy	209,317	0.14% 2.45
44782429		Chronic kidney disease due to hypertension	170,534	0.12% 3.60
137940		Transient hypertension of pregnancy - delivered	153,806	0.11% 1.07
321080		Hypertension complicating pregnancy, childbirth and the puerperium	148,728	0.10% 2.15
314423		Benign essential hypertension complicating pregnancy, childbirth and the puerperium - not delivered	132,245	0.09% 3.94
44782690		Chronic kidney disease stage 5 due to hypertension	119,375	0.08% 5.20
44783618		Heritable pulmonary arterial hypertension	104,737	0.07% 3.61
319826		Secondary hypertension	96,356	0.07% 2.14
4167493		Pregnancy-induced hypertension	91,675	0.06% 2.60
321074		Pre-existing hypertension complicating pregnancy, childbirth and puerperium	74,311	0.05% 2.99
192680		Portal hypertension	71,240	0.05% 3.11

Showing 1 to 15 of 47 entries  
(filtered from 15,907 total entries)

Previous 1 2 3 4 Next

Figure 12.4: ATLAS□□□□□□: □□□□□□□□□□□□□□□□□□

essential hypertension

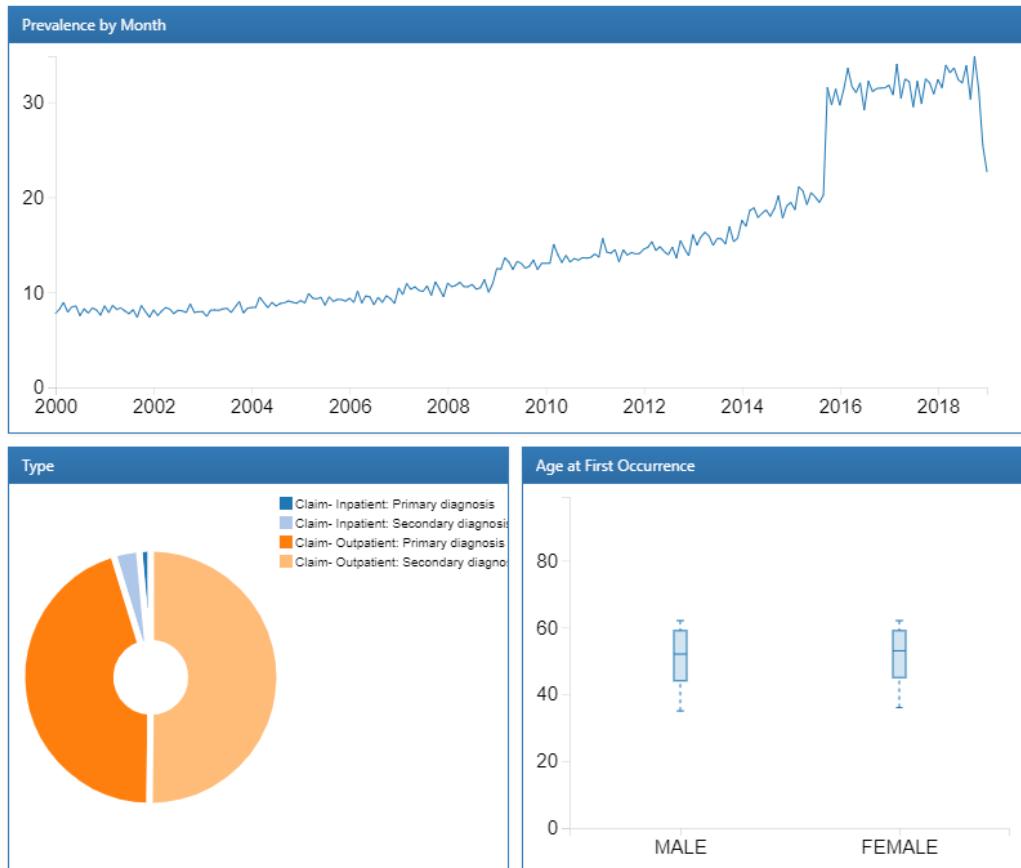


Figure 12.5: ATLAS: Essential Hypertension

Era

## 12.7 ATLAS

ATLAS

### 12.7.1

1 1 2 B.6 2 1 3 B.7

## Design

**Cohort characterization** is defined as the process of generating cohort level descriptive summary statistics from person level covariate data. Summary statistics of these person level covariates may be count, mean, sd, var, min, max, median, range, and quantiles. In addition, covariates during a period may be stratified into temporal units of time for time-series analysis such as fixed intervals of time relative to cohort\_start\_date (e.g. every 7 days, every 30 days etc.), or in absolute calendar intervals such as calendar-week, calendar-month, calendar-quarter, calendar-year.

## Cohort definitions

**Import**

ID	Name	Edit cohort	Remove
10447	Patients initiating first-line therapy for hypertension with >1 yr follow-up	<a href="#">Edit cohort</a>	<a href="#">Remove</a>
10448	Patients initiating first-line therapy for hypertension with >3 yr follow-up	<a href="#">Edit cohort</a>	<a href="#">Remove</a>

Figure 12.6: □□□□□ - □□□□□□□□□

6 / 6

ATLAS Chapter 11 Import 12.6

ATLAS OMOP CDM Feature Extraction R

Import 

Demographics



□ □ □ □ □ □ □

## Feature analyses

**Import**

Show 25 ▾ entries Search:

ID	Name	Description	Actions
43	Drug Era Short Term	One covariate per drug in the drug_era table overlapping with any part of the short window.	<a href="#">Remove</a>
49	Charlson Index	The Charlson comorbidity index (Romano adaptation) using all conditions prior to the window end.	<a href="#">Remove</a>
67	Condition Occurrence Long Term	One covariate per condition in the condition_occurrence table starting in the long term window.	<a href="#">Remove</a>
71	Demographics Age Group	Age of the subject on the index date (in 5 year age groups)	<a href="#">Remove</a>
72	Demographics Race	Race of the subject.	<a href="#">Remove</a>
73	Demographics Prior Observation Time	Number of continuous days of observation time preceding the index date.	<a href="#">Remove</a>
74	Demographics Gender	Gender of the subject.	<a href="#">Remove</a>
76	Condition Occurrence Medium Term	One covariate per condition in the condition_occurrence table starting in the medium term window.	<a href="#">Remove</a>
77	Demographics Age	Age of the subject on the index date (in years).	<a href="#">Remove</a>
79	Demographics Time In Cohort	Number of days of observation time during cohort period.	<a href="#">Remove</a>
80	Demographics Index Year	Year of the index date.	<a href="#">Remove</a>
81	Demographics Post Observation Time	Number of continuous days of observation time following the index date.	<a href="#">Remove</a>
87	Procedure Occurrence Any Time Prior	One covariate per procedure in the procedure_occurrence table any time prior to index.	<a href="#">Remove</a>
103	Visit Count Long Term	The number of visits observed in the long term window.	<a href="#">Remove</a>

Figure 12.7: □□□□□□ - □□□□

## Subgroup analyses

**New subgroup**

Calculate subgroup analyses only

having **all** of the following criteria:

**+ Add criteria to group...**

with the following event criteria:

**+ Add attribute...**

**X** with a gender of: **FEMALE** **Add** **Import** **Delete Criteria**

Figure 12.8: □□□□ - □□□□□□□□□□□



ATLAS

12.7.2

Design	Executions	Utilities
<h2>Executions</h2>		
SYNPUF 1K	Generate	View latest result
SYNPUF 5%	Generate	View latest result

Figure 12.9: □□□□□□ - CDM □□□□□□

All Executions View  
Reports View latest result

12.7.3

CONDITION / Condition Occurrence Long Term / stratified by Female												
<a href="#">Export</a> <a href="#">Export comparison</a>			Show 10 ▾ entries					Search: <input type="text" value="cardi"/>				
Covariate	Explore	Concept ID	Patients initiating first-line therapy for hypertension with > 1 yr follow-up				Patients initiating first-line therapy for hypertension with > 3 yr follow-up				Std diff ▾	
			Count		Pct		Female		Count			
			Count		Pct		Count		Pct			
Tachycardia	<a href="#">Explore ▾</a>	<a href="#">444070</a>	17,322	1.04%	9,042	1.18%	6,547	0.78%	3,530	0.90%	-0.0193	
Cardiomegaly	<a href="#">Explore ▾</a>	<a href="#">314658</a>	20,958	1.26%	8,007	1.04%	9,016	1.08%	3,465	0.89%	-0.0121	
Cardiac arrhythmia	<a href="#">Explore ▾</a>	<a href="#">44784217</a>	30,474	1.83%	13,221	1.72%	14,540	1.74%	6,318	1.62%	-0.0052	

Figure 12.10: □□□□ - □□1□□□□□□□□

12.10 365

A horizontal row of 20 empty square boxes for writing responses.

[@cerning.atlasCharacterizationResultsExplore](#)          

@ref{fig:atlasCharacterizationResultsContra}

Exploring condition_occurrence during day -365 through 0 days relative to index: Cardiac arrhythmia							X
Cohort: Patients initiating first-line therapy for hypertension with >1 yr follow-up							
Export	Show 10 ▾ entries					Search: <input type="text"/>	
Relationship type	Distance	Concept name		All stratas		Female	
		Count	Pct	Count	Pct	Count	Pct
Explore Ancestor	4	Disorder by body site		32	0.00%	17	0.00%
Explore Ancestor	4	Finding of trunk structure		991	0.06%	605	0.08%
Explore Ancestor	3	Disorder of trunk		23	0.00%	14	0.00%
Explore Ancestor	3	Disorder of thorax		241	0.01%	104	0.01%
Explore Ancestor	3	Disorder of body system		4,135	0.25%	1,992	0.26%
Explore Ancestor	2	Disorder of cardiovascular system		12,979	0.78%	6,073	0.79%
Explore Ancestor	2	Disorder of mediastinum		138	0.01%	62	0.01%
Explore Ancestor	2	Disorder of body cavity		24	0.00%	10	0.00%
Explore Ancestor	1	Heart disease		4,691	0.28%	1,869	0.24%
Explore Selected	0	Cardiac arrhythmia		30,474	1.83%	13,221	1.72%

Showing 1 to 10 of 62 entries

Previous 1 2 3 4 5 6 7 Next

Figure 12.11: □□□□ - □□□□□□□□

## CONDITION / Condition Occurrence Long Term / stratified by Female

Export	Export comparison	Show 10 ▾ entries	Patients initiating first-line therapy for hypertension with >1 yr follow-up				Patients initiating first-line therapy for hypertension with >3 yr follow-up				Std diff ▾	
Covariate	Explore	Concept ID	Female				Female					
			Count	Pct	Count	Pct	Count	Pct	Count	Pct		
Edema	Explore ▾	433595	32,243	1.94%	20,200	2.63%	15,173	1.81%	9,684	2.48%	-0.0066	

Showing 1 to 1 of 1 entries (filtered from 206 total entries)

Previous 1 Next

Figure 12.12: □□□□ - □□□□□□□□

explore

Exploring condition_occurrence during day -365 through 0 days relative to index: Edema							
Cohort: Patients initiating first-line therapy for hypertension with >1 yr follow-up							
Export		Show 10 ▾ entries		Search: and			
Relationship type	Distance	Concept name	All stratas		Female		Pct
			Count	Pct	Count	Pct	
Explore Descendant	-2	Angioedema	2,605	0.16%	1,506	0.20%	

Figure 12.13: □□□□ - □□□□□□□□□□□□

1

## DEMOGRAPHICS / Demographics Age

Strata		Patients initiating first-line therapy for hypertension with > 1 yr follow-up				Patients initiating first-line therapy for hypertension with > 3 yr follow-up				Std diff ▼
		Count	Avg	Std Dev	Median	Count	Avg	Std Dev	Median	
Female	768,180	49.39	9.78	51.00	390,693	49.01	9.03	51.00	-0.0291	
All stratas	1,661,604	48.96	10.00	50.00	837,459	48.64	9.26	50.00	-0.0232	

Figure 12.14: □□□□□□□□□□□□□□□□□□□□□□□□

12.7.4

ATLAS Analysis New Feature Analysis 

██████████ ACE ██████████

## Analysis

128 R

R Feature Extraction OHDSI

**Design**

**Criteria** Custom

Analysis type:

Prevalence

Add Criteria feature

Ace inhibitor exposure after index

having all of the following criteria:

+ Add criteria to group... □

with at least 1 using all occurrences of:

+ Add attribute... □

a drug era of ACE inhibitors □

where event starts between 1 days After and All days After □ index start date add additional constraint □

allow events from outside observation period □

Delete Criteria

Figure 12.15: ATLAS□□□□□□□□□□□

## DRUG / Ace inhibitor exposure after index / stratified by Female

			Patients initiating first-line therapy for hypertension with >1 yr follow-up				Patients initiating first-line therapy for hypertension with >3 yr follow-up				Std diff	
Covariate	Explore	Concept ID	Count	Pct	Female		Count	Pct	Female			
					Count	Pct			Count	Pct		
Ace inhibitor exposure after index	Explore ▾	0	686,034	41.29%	289,215	17.41%	426,280	50.90%	182,219	21.76%	0.1001	

Showing 1 to 1 of 1 entries

Previous 1 Next

Figure 12.16: □□□□□□□□□□□

- □□□□□□□□□□□□□□□□
  - □□□□□□□□□□□□□□□□□□
  - □□□□□□□□□□□□□□□□

## Feature Extraction

## 12.8.1

Chapter  
11 B.6 Appendix B scratch.my\_cohorts

## 12.8.2

```
library(FeatureExtraction)
connDetails <- createConnectionDetails(dbms = "postgresql",
                                         server = "localhost/ohdsi",
                                         user = "joe",
                                         password = "supersecret")

cdmDbSchema <- "my_cdm_data"
cohortsDbSchema <- "scratch"
cohortsDbTable <- "my_cohorts"
cdmVersion <- "5"
```

```
cdmDbSchema cohortsDbSchema cohortsDbTable CDM  
SQL Server  
<- "my_cdm_data.dbo"
```

### 12.8.3

```
createCovariateSettings
```

```
settings <- createCovariateSettings(  
  useDemographicsGender = TRUE,  
  useDemographicsAgeGroup = TRUE,  
  useConditionOccurrenceAnyTimePrior = TRUE)
```

```
settings <- createCovariateSettings(useConditionEraLongTerm = TRUE,  
                                     useConditionEraShortTerm = TRUE,  
                                     useDrugEraLongTerm = TRUE,  
                                     useDrugEraShortTerm = TRUE,  
                                     longTermStartDays = -180,  
                                     shortTermStartDays = -14,  
                                     endDays = -1)
```

180 14  
ID

```
settings <- createCovariateSettings(useConditionEraLongTerm = TRUE,  
                                     useConditionEraShortTerm = TRUE,  
                                     useDrugEraLongTerm = TRUE,  
                                     useDrugEraShortTerm = TRUE,  
                                     longTermStartDays = -180,  
                                     shortTermStartDays = -14,  
                                     endDays = -1,  
                                     excludedCovariateConceptIds = 1124300,  
                                     addDescendantsToExclude = TRUE,  
                                     aggregated = TRUE)
```



aggregated = TRUE Feature Extraction

## 12.8.4

```
covariateSettings <- createDefaultCovariateSettings()  
  
covariateData2 <-getDbCovariateData(  
  connectionDetails = connectionDetails,  
  cdmDatabaseSchema = cdmDatabaseSchema,  
  cohortDatabaseSchema = resultsDatabaseSchema,  
  cohortTable = "cohorts_of_interest",  
  cohortId = 1,  
  covariateSettings = covariateSettings,  
  aggregated = TRUE)  
  
summary(covariateData2)
```

```
## CovariateData Object Summary  
##  
## Number of Covariates: 41330  
## Number of Non-Zero Covariate Values: 41330
```

## 12.8.5

```
covariateData2$covariates  
covariateData2$covariatesContinuous
```

12.8.6

Feature Extraction //ohdsi.github.io/FeatureExtraction/

## 12.9 ATLAS

Hripcak7329 ATLAS Cohort  
Pathways



**Figure 12.17:** □□□□□□□□□□□□□□□□□□□□□□□□

ATLAS Cohort Pathways

12.9.1



## Event Cohorts

Each Event Cohort defines the step in a pathway that may occur for a person in the Target Cohort.

## Import

Show 10 ▾ entries

Search:

ID	Name		
9174	<a href="#">ACE inhibitor use</a>	Edit cohort	Remove
9175	<a href="#">Angiotensin receptor blocker (ARB) use</a>	Edit cohort	Remove
9176	<a href="#">Thiazide or thiazide-like diuretic use</a>	Edit cohort	Remove
9177	<a href="#">dihydropyridine Calcium Channel Blocker (dCCB) use</a>	Edit cohort	Remove
9178	<a href="#">non-dihydropyridine Calcium Channel Blocker (ndCCB) use</a>	Edit cohort	Remove
9179	<a href="#">beta blocker use</a>	Edit cohort	Remove
9180	<a href="#">Diuretic-loop use</a>	Edit cohort	Remove
9181	<a href="#">Diuretic-potassium sparing use</a>	Edit cohort	Remove
9182	<a href="#">alpha-1 blocker use</a>	Edit cohort	Remove

Showing 1 to 9 of 9 entries

Previous 1 Next

Figure 12.19: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

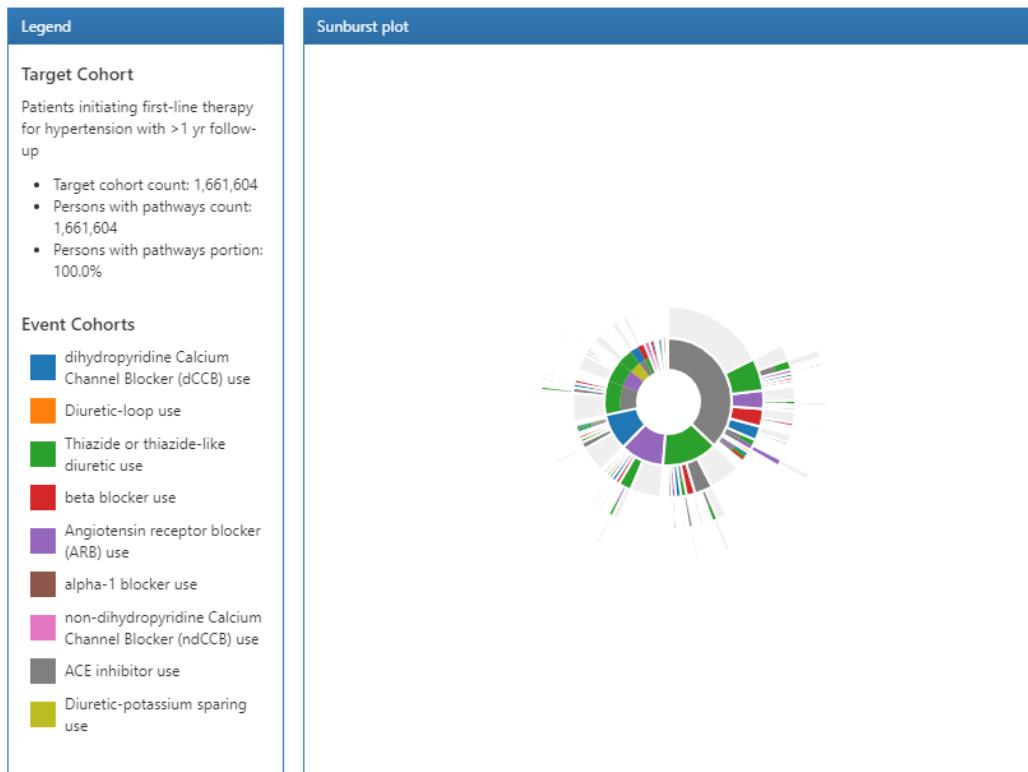
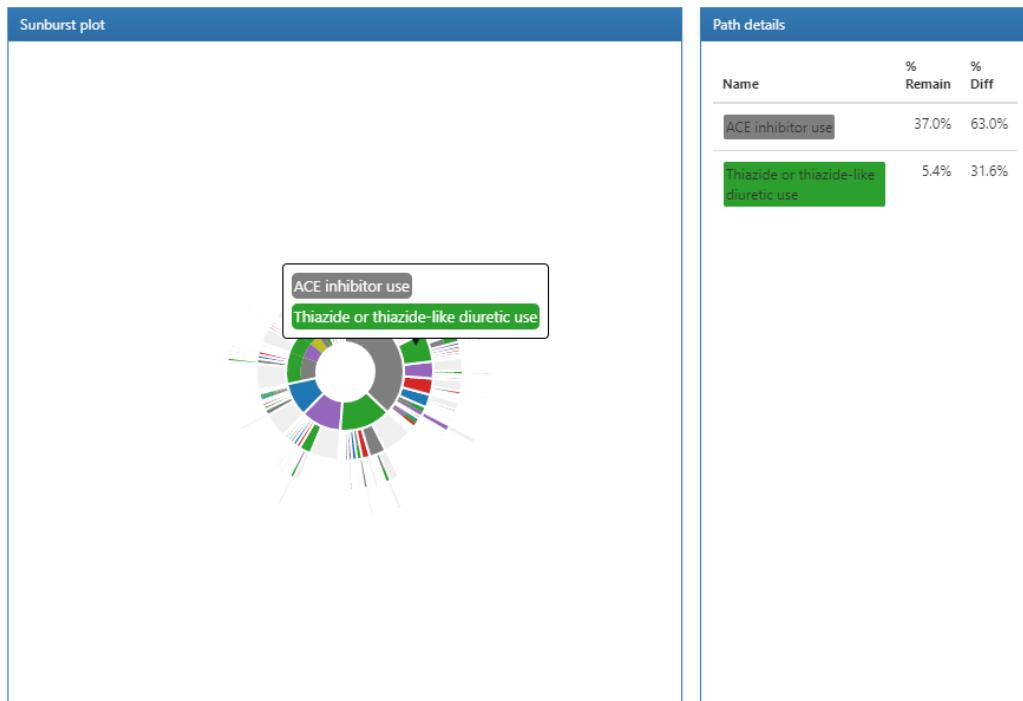


Figure 12.20: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□



**Figure 12.21:** □□□□□□□□□□□□□□□□

ATLAS Incidence Rates

## 12.10.1

□□□□□□□□□□□□□□□□□ ATLAS □□□□□□□□□□□□□□□ (Chapter 11  
□□□)□□□□□□□□□□□□□□□ (Appendix B.2, B.5) □□□□□□□□□□□□□  
(Appendix B.4, B.3, B.9) □□□□□□□□□□□□□□□

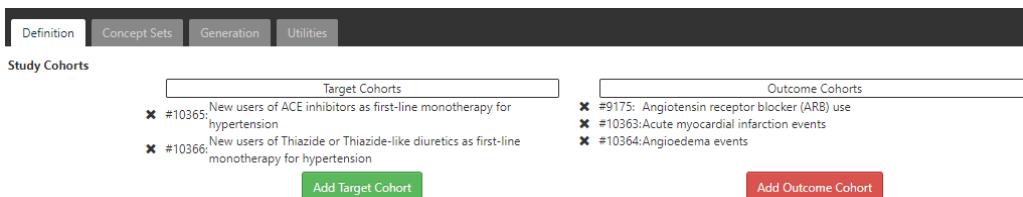


Figure 12.22: □□□□□□□□□□□□□□□□□□□□□□□□

## Time At Risk

Time at risk defines the time window relative to the cohort start or end date with an offset to consider the person 'at risk' of the outcome.

- Time at risk starts with **start date** ▾ plus **1** ▾ days.
  - Time at risk ends with **end date** ▾ plus **0** ▾ days.

No study window defined.

### Add Study Window

Figure 12.23: □□□□□□□□□□□□□□□□□□□□□□□□

# THz

ATLAS :

**Stratify Criteria:** You can provide optional stratification criteria to the analysis that will divide the population into unique groups based on their satisfied criteria.

New stratify criteria	Gender = Female	Copy	Delete
1. Gender = Female			
enter an inclusion rule description			
having <input style="border: 1px solid #ccc; padding: 2px 10px;" type="button" value="all"/> of the following criteria: <span style="float: right;"><input style="border: 1px solid #0078D4; background-color: #0078D4; color: white; padding: 2px 10px;" type="button" value="+"/> Add criteria to group...</span>			
with the following event criteria: <span style="float: right;"><input style="border: 1px solid red; background-color: red; color: white; padding: 2px 10px;" type="button" value="Delete Criteria"/></span>			
<span style="border: 1px solid #0078D4; padding: 2px 10px; display: inline-block;"><input style="border: none; font-size: small; margin-right: 5px;" type="button" value="+"/> Add attribute...</span>			
<span style="color: red; font-weight: bold;">✖</span> with a gender of: <span style="border: 1px solid #ccc; padding: 2px 10px; display: inline-block;">✖ FEMALE</span> <span style="margin-left: 10px;"><input style="border: 1px solid #ccc; padding: 2px 10px;" type="button" value="Add"/></span> <span style="margin-left: 10px;"><input style="border: 1px solid #ccc; padding: 2px 10px;" type="button" value="Import"/></span>			

Figure 12.24: □□□□□□□□□□□□□□

Chapter 11

12.10.2

[ ] **► Generate** [ ]

12.10.3

**TAR**

**1000** □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □



## Select sources

Filter:

## Name



SYNPUF 1K



SYNPUF 5%

Previous 1 Next

Figure 12.25: □□□□□□□□

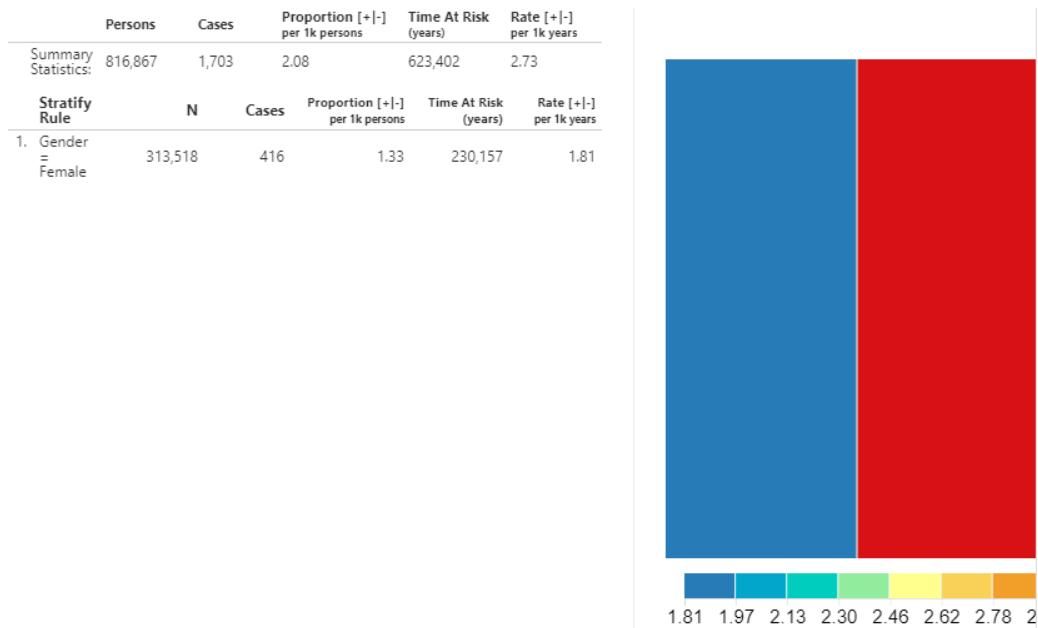


Figure 12.26: □□□□□□□ - AMI □□□□□□□□□□ ACEi □□□□

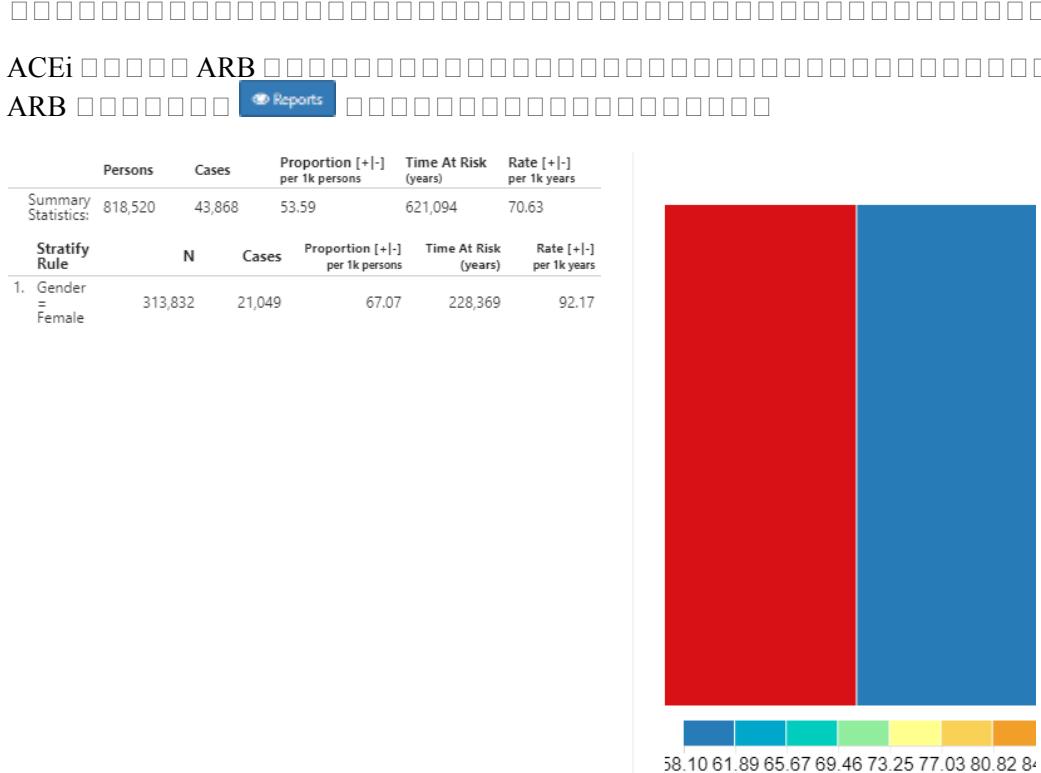


Figure 12.27: □□□ - ACEi □□□□ ARB □□□□□□□□□□ ACEi □□□□

12.11



- OHDSI
  - 
  - ATLAS OHDSI
  - ATLAS

12.12



ATLAS <http://atlas-demo.ohdsi.org>

## **12.1.**

**12.3.** (GI) GI (“”) GI GI

□□□□□□□ Appendix E.6 □□□□□□□□□□

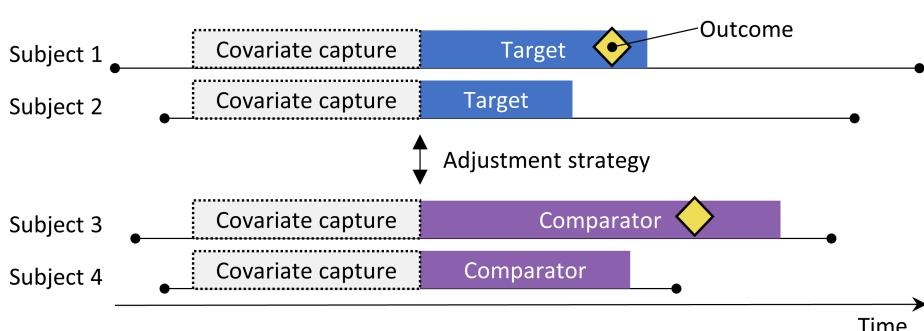
# Chapter 13

   <img alt="Open Access icon" data-bbox="10236 88



□□□□□□□ QUDSI Methods Library □

131



□: □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □



### 13.1.1

PS (Rosenbaum and Rubin, 1983, ) 0.5 et al., 2012, )

PS Jan 0.4  
Jun

### 13.1.2

PS OHDSI et al., 2018, ) et al., 2013, ) Cyclops



13.1.3

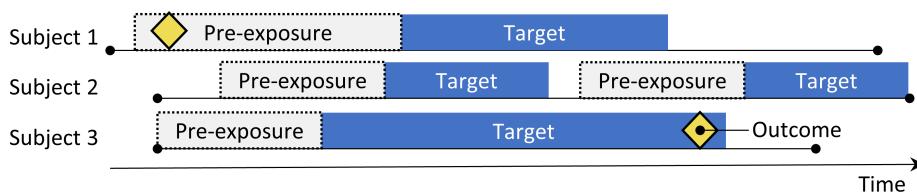
### 13.1.4

$$\ln\left(\frac{F}{1-F}\right) = \ln\left(\frac{S}{1-S}\right) - \ln\left(\frac{P}{1-P}\right)$$

13.1.5

PS 13.11.2001, )

13.2



□□□□□□□□□SCC□□□□□(Ryan et al., 2013, ) □□□□  
13.2 □□□4□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

13,3

Table 13.2: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

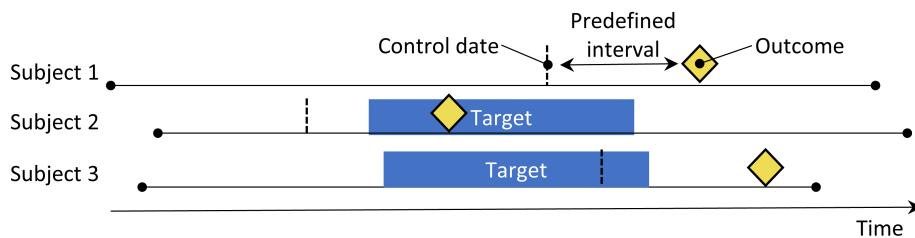


13.4

13.5

SCCS (Farrington, 1995; Whitaker et al., 2006,  
SCCS

**Table 13.3:** □□□□□□□□□□□□□□□□□□



**Table 13.4:** □□□□□□□□□□□□□□□□□□□□□□

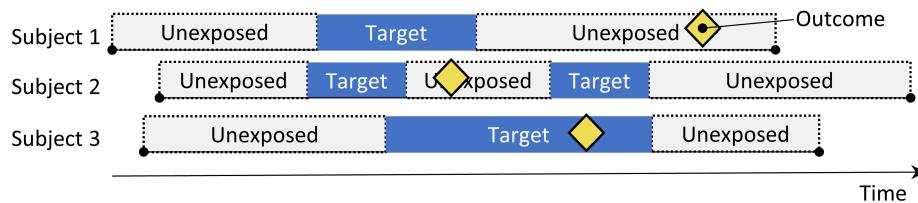
SCCS et al., 2013, )

13.6

13.6.1

ACE (Huang et al., 2002, ) and Black, 1997, ) et al., 2012, ) ACEi  $\beta$  (Mak et al., 2010; Toh et al., 2012, )  $\beta$  (Whelton et al., 2018, ) THZ

Table 13.5: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□



□□□□□□□□□□□□□□□□□□@ref(CohortMethod)□□□□□□□□□□□□□□□□

13.6.2

ACEi THZ

### 13.6.3

ER 7

13.6.4

13.6.5

13.6.6

A decorative horizontal bar consisting of a series of small, evenly spaced rectangular boxes, likely a placeholder for an image or a decorative element.

### 13.6.7

MethodValidity  
@ref

## 13.7 ATLAS

ATLAS  
Estimation  
G

3

### 13.7.1

1  
2  
11 Appendix B.2 Appendix @ref(ThiazidesMono  
@ref(Angioedema  
Appendix @ref(Ami  
13.6

Comparison  
Add or update the target, comparator, outcome(s) cohorts and negative control outcomes

Choose your target cohort:

New users of ACE inhibitors as first-line monotherapy for hypertension

Choose your comparator cohort:

New users of Thiazide-like diuretics as first-line monotherapy for hypertension

Choose your outcome cohorts:

**Add Outcome**

Show 10 entries		Search: <input type="text"/>
ID	Name	
1770712	Angioedema outcome	<a href="#">Edit cohort</a> <a href="#">Remove</a>
1770713	Acute myocardial infarction outcome	<a href="#">Edit cohort</a> <a href="#">Remove</a>

Showing 1 to 2 of 2 entries

Previous 1 Next

Figure 13.6: ATLAS

MethodValidity  
@ref

MethodValidity  
@ref

19

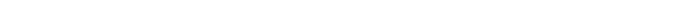
13.7 

Negative controls for ACEi and THZ									Optimize							
Concept Set Expression		Included Concepts <span style="border: 1px solid black; padding: 2px;">75</span>		Included Source Codes		Explore Evidence		Export		Compare						
Show <span style="border: 1px solid black; padding: 2px;">25 ▾</span> entries						Search: <input type="text"/>										
Showing 1 to 25 of 75 entries																
	Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	Previous <span style="border: 1px solid black; padding: 2px;">1</span>	2	3	Next				
	72748	74779009	Strain of rotator cuff capsule	Condition	Standard	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>								
	73241	197210001	Anal and rectal polyp	Condition	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>								
	73560	55260003	Calcaneal spur	Condition	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>								
	75911	65358001	Acquired hallux valgus	Condition	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>								
	76786	63643000	Derangement of knee	Condition	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>								

**Figure 13.7:** □□□□□□□□□□□□□□□□□□

5.5.1.1

□ □ □ □ □ □

13.8 

Concept Set #1798551																
Concepts to exclude for ACEi and THZ										Optimize						
Concept Set Expression			Included Concepts <span style="border: 1px solid black; padding: 2px;">38225</span>		Included Source Codes		Explore Evidence		Export	Compare						
Show <span style="border: 1px solid black; padding: 2px;">25</span> entries										Search: <input type="text"/>						
Showing 1 to 14 of 14 entries										Previous <span style="border: 1px solid black; padding: 2px;">1</span> Next						
Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	<input checked="" type="checkbox"/> Exclude	<input checked="" type="checkbox"/> Descendants	<input type="checkbox"/> Mapped									
1342439	38454	trandolapril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>									
1334456	35296	Ramipril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>									
1331235	35208	quinapril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>									
1373225	54552	Perindopril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>									
1310756	30131	moexipril	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>									

Figure 13.8: □□□□□□□□□□□□□□□□

Figure 13.9

Negative control concept set:

Negative controls for ACEi and THZ



Covariate selection

Concepts to **include** when constructing the covariates to be used in this study. (Leave blank if you want to include every concept).\*



\* Concepts defined here are combined with those defined in the Analysis settings section.

Concepts to **exclude** when constructing the covariates to be used in this study.\*

Concepts to exclude for ACEi and THZ



\* Concepts defined here are combined with those defined in the Analysis settings section.

Figure 13.9: R

### 13.7.2

R

R

R

R

R

### 13.8 R

R ATLAS R

CohortMethod CohortMethod CDM

13.8.6 AMI

#### 13.8.1

(1) (B.2) (B.5)

(B.4) ACEi THZ

1 2 3 scratch.my\_cohorts

#### 13.8.2

R CohortMethod DatabaseConnector

```
library(CohortMethod)
connDetails <- createConnectionDetails(dbms = "postgresql",
                                         server = "localhost/ohdsi",
                                         user = "joe",
                                         password = "supersecret")

cdmDbSchema <- "my_cdm_data"
cohortDbSchema <- "scratch"
cohortTable <- "my_cohorts"
cdmVersion <- "5"
```

```
cdmDbSchema cohortDbSchema cohortTable CDM  
SQL Server cdm  
<- "my_cdm_data.dbo"
```

## CohortMethod

```
## CohortMethodData  
##  
##      ID 1  
##      ID 2  
##      ID(s) 3
```

CohortMethod createDefault  
(13.1) 2

cohortsMethodData  
(9.4.2)□□□□□□□

summary()

```
summary(cmData)
```

```
## CohortMethodData
##
##      ID 1
##      ID 2
##      ID(s) 3
##
##      67166
##      35333
##
##      980          891
##  3
##
##      58349
##      24484665
```

**cohortMethodData** ━━━━━━

```
loadCohortMethodData()
```

□ □ □ □ □ □ □ □ □

1. `getStudyPopulation`
  2. `getDbCohortMethodData`
  3. `createStudyPopulation`

## CohortMethod

### 13.8.3

```
getAttritionTable(studyPop)
```

```
##                                     ...
## 1                               67212      35379 ...
## 2           67166      35333 ...
## 3                               67061      35238 ...
## 4     1       66780      35086 ...
```

13.8.4

```
getDbcohortMethodData()
```

```
ps <- createPs(cohortMethodData = cmData, population = studyPop)
```

```
matchedPop <- matchOnPs(population = ps, caliper = 0.2,  
                        caliperScale = "standardized logit", maxRatio = 100)
```

□□□□□PS□trimByPs□trimByPsToEquipoise□□□□stratifyByPs□□□□□□□□□□□□□

### 13.8.5

```
outcomeModel <- fitOutcomeModel(population = matchedPop,
                                  modelType = "cox",
                                  stratified = TRUE)

outcomeModel

##      cox
##    TRUE
##   FALSE
##   FALSE
##     OK
##
##                95%  95% logRr seLogRr
## 4.3203    2.4531  8.0771 1.4633  0.304
```

13.8.6

```
#  
ois <- c(3, 4) #      AMI  
  
#  
ncs <- c(434165, 436409, 199192, 4088290, 4092879, 44783954, 75911, 137951, 77965,  
       376707, 4103640, 73241, 133655, 73560, 434327, 4213540, 140842, 81378,  
       432303, 4201390, 46269889, 134438, 78619, 201606, 76786, 4115402,  
       45757370, 433111, 433527, 4170770, 4092896, 259995, 40481632, 4166231,  
       433577, 4231770, 440329, 4012570, 4012934, 441788, 4201717, 374375,  
       4344500, 139099, 444132, 196168, 432593, 434203, 438329, 195873, 4083487,  
       4103703, 4209423, 377572, 40480893, 136368, 140648, 438130, 4091513,  
       4202045, 373478, 46286594, 439790, 81634, 380706, 141932, 36713918,  
       443172, 81151, 72748, 378427, 437264, 194083, 140641, 440193, 4115367)  
  
tcos <- create  
  
##      {#studyOutputs}
```

ATLAS R R

###

(ROC) AUC AUC 1

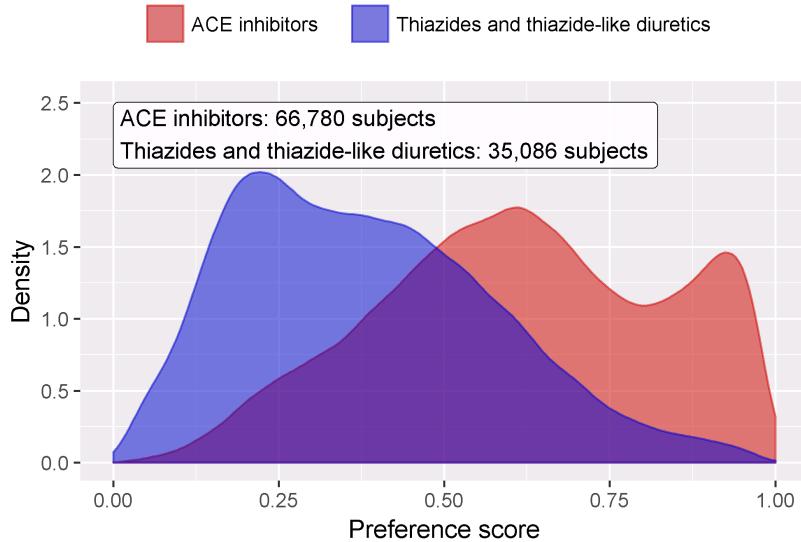


Figure 13.10: □□□□□□□□

0.68         : 05-09

0.64    □□□□□-365□□□0□□□□□□□□□: □□

0.55    □□□□□-365□□□0□□□□□□□□□□□□: □□□□

0.52 (□□)

0.50      □ □ :

---

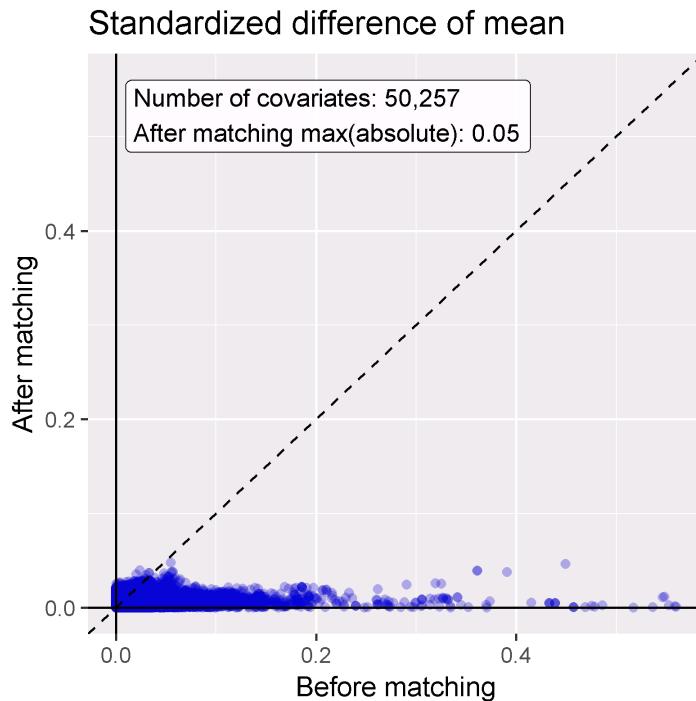
www.IBM.com

13.8.7

PS 2 13.11 0.1

13.8.8

13.12 drawAttritionDiagram



13.9



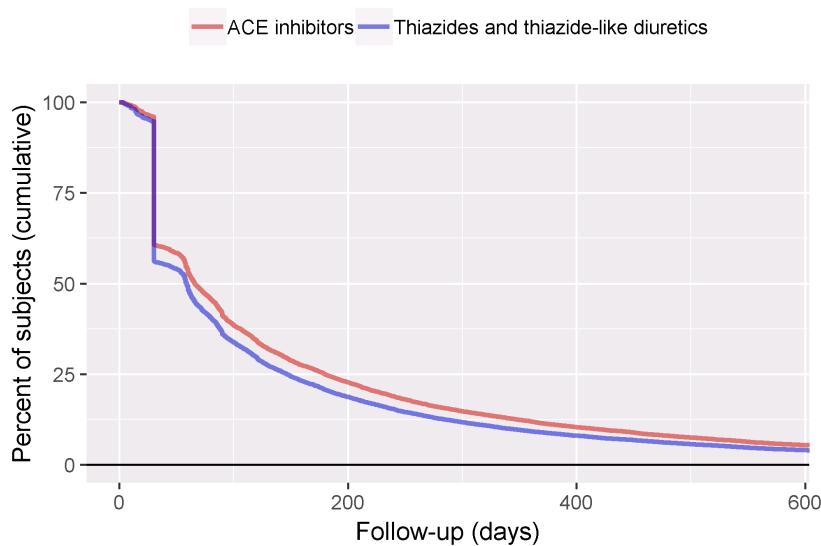
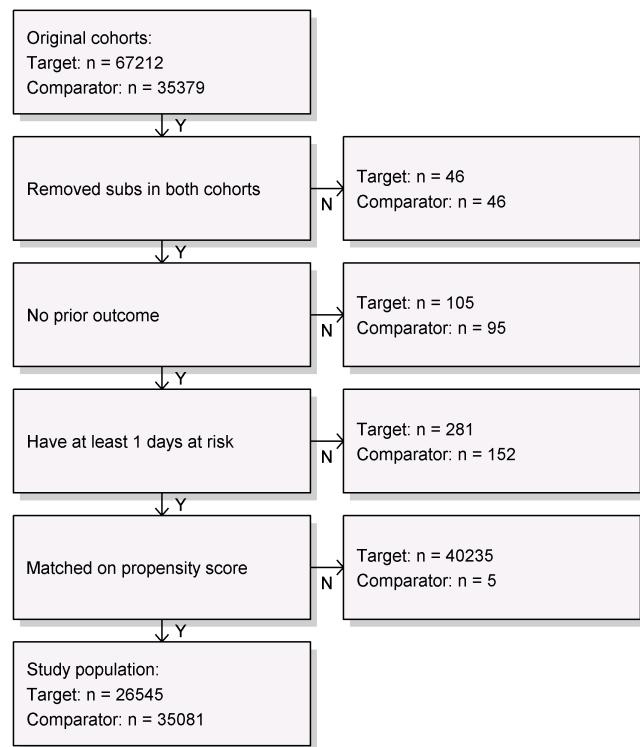



13.10

10 of 10

□□□□□□□□□□□□□ R R-Studio □□□□ Java 9.4.5 □□□□□□□□□□□□□

```
install.packages(c("SqlRender", "DatabaseConnector", "remotes"))
remotes::install_github("ohdsi/Eunomia", ref = "v1.0.0")
```



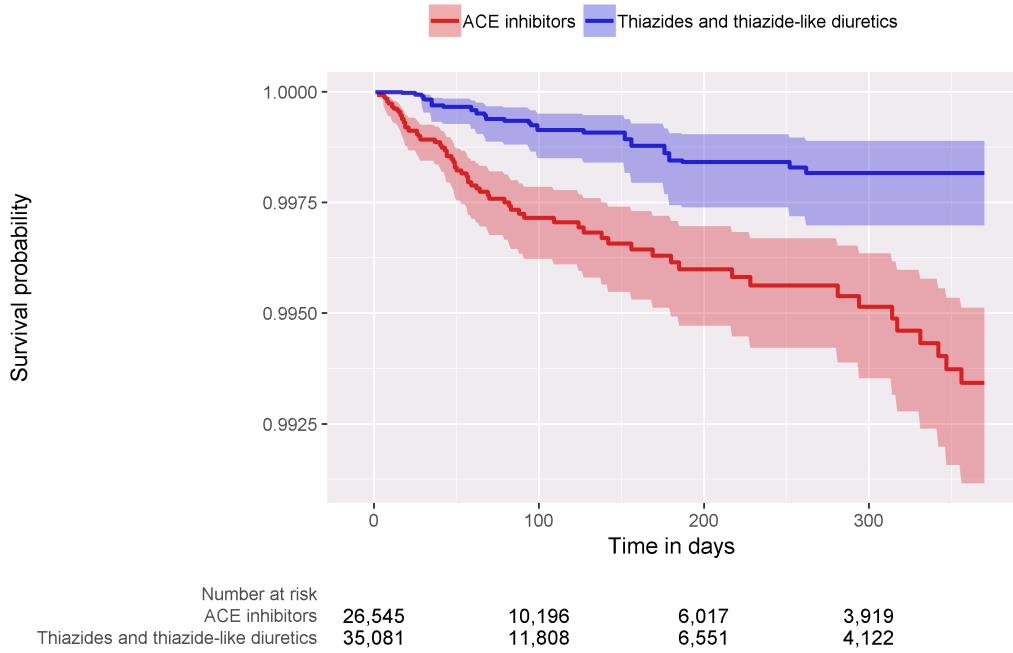


Figure 13.14: □□□□—□□□□□□□□□□□

```
remotes::install_github("ohdsi/CohortMethod")
```

Eunomia R CDM

```
connectionDetails <- Eunomia::getEunomiaConnectionDetails()
```

CDM main Eu

```
Eunomia:::createCohorts(connectionDetails)
```

1

□□□□□□□□□□□□□□□□□ COHORT DEFINITION ID □1□□□□□□□□□□□□□□□□□

□ 13.1. CohortMethod R CDM □ CohortM

### **13.3.**

□ □ 13.5. 5 □ □ □ □ □ □ PS □

E.7

# Chapter 14

Lead: Peter Rijnbeek & Jenna Reps

OHDSI  CDM

14.1

[□□□□□□□□□□] J □□□□□□□□ [□□□□□□□□] J □□□□□□□□□□□

□: □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □

<sup>1</sup><https://www.equator-network.org/reporting-guidelines/tripod-statement/>

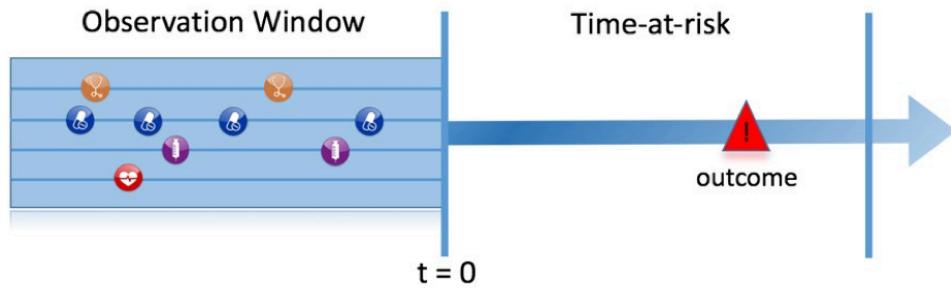


Figure 14.1: □□□□□



A horizontal row of 20 empty square boxes, intended for children to write their names in, likely as part of a classroom activity or name recognition exercise.



14.2

14.2.1

COHORT_DEFINITION_ID	SUBJECT_ID	COHORT_START_DATE
1	1	2000-06-01
1	2	2001-06-01
2	2	2001-07-01

PERSON_ID	CONDITION_CONCEPT_ID	CONDITION_START_DATE
1	320128	2000-10-01
2	320128	2001-05-01

## 14.2.2 vs

14.3

14.2

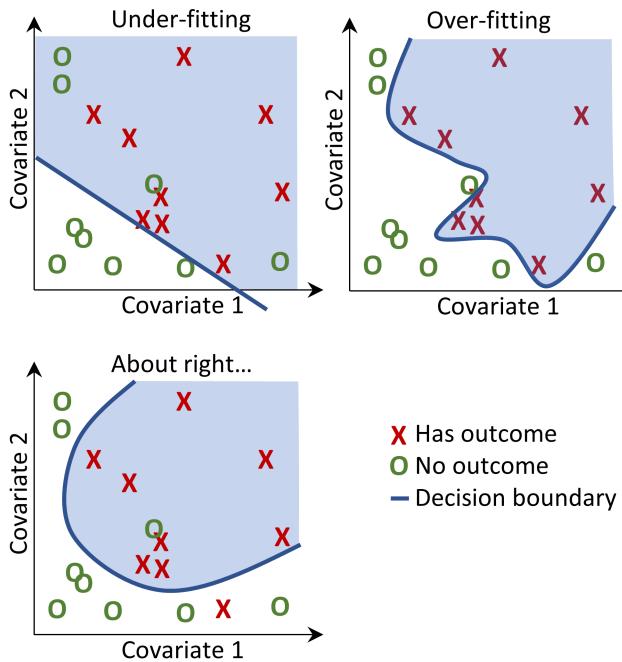


Figure 14.2: Decision boundary.

### 14.3.1

The diagram consists of two horizontal rows of empty rectangular boxes. The top row contains four boxes on the left, two empty boxes in the center, and four empty boxes on the right. The bottom row contains three boxes on the left, followed by a group of seven empty boxes, and the number '0.1' on the right.

14.3.2

R

Table 14.5: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

□□□□□ □□ □□□□□  
earlyStopRound □□□□□□□□□□□□□□ 25

□□□□□	□□	□□□□□
learningRate	□□□□□□□□□□□□	0.005,0.01,0.1
maxDepth	□□□□□□	4,6,17
minRows	□□□□□□□□□□□□□□	2
ntrees	□□□	100,1000

### **14.3.3**

**Table 14.6:** □□□□□□□□□□□□□□□□□□□□□□□□

<input type="checkbox"/>					
maxDepth	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	4,10,17
mtries	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	-1 = <input type="checkbox"/> <input type="checkbox"/> <input type="checkbox"/> <input type="checkbox"/> <input type="checkbox"/> <input type="checkbox"/> ,5,20
ntrees	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	500	

#### 14.3.4 K-

## K-nearest neighbors (KNN)

Table 14.7: K-□□□□□□□□□□□□□□□□

<input type="checkbox"/>						
k					1000	

14.3.5

### 14.3.6 AdaBoost

**AdaBoost**   
**AdaboostClassifier**

Table 14.8: AdaBoost

□□□□□	□□	□□□□□
nEstimators	□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□4□□□	
learningRate	□□□□□□□□□□□□□□learning_rate□□□□□□□□□□□□□□□□learningRate□□E	

---

□□□□□ □□ □□□□□

---

### **14.3.7**

DecisionTreeClassifier

Table 14.9: □□□□□□□□□□□□□□

classWeight	“Balance”	”None”
maxDepth	10	10
minImpuritySplit	10	10
minSamplesLeaf	10	10
minSamplesSplit	2	2

14.3.8

Table 14.10: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□

□□□□□	□□	□□□□□
alpha	I2□□□	0.00001
size	□□□□□□□	4

14.3.9

14.3.10

14.4

## 14.4.1



14.3

A horizontal row of 20 empty square boxes, intended for students to draw or write in.



□□□□□□□□□□□□□□□□**2**□□□□□□□□



## 14.4.2

100

0 1  
14.11 0.5 137100.5

**Table 14.11:** □□□□□□□□□□□□□□□□

ID	0.5	1	1	TP
1	0.8	1	1	TP
2	0.1	0	0	TN
3	0.7	1	0	FP
4	0	0	0	TN
5	0.05	0	0	TN
6	0.1	0	0	TN
7	0.9	1	1	TP
8	0.2	0	1	FN
9	0.3	0	0	TN
10	0.5	1	0	FP

TP

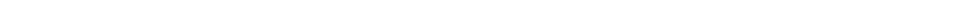
A horizontal row of 20 empty square boxes, likely used for grading or marking student responses.

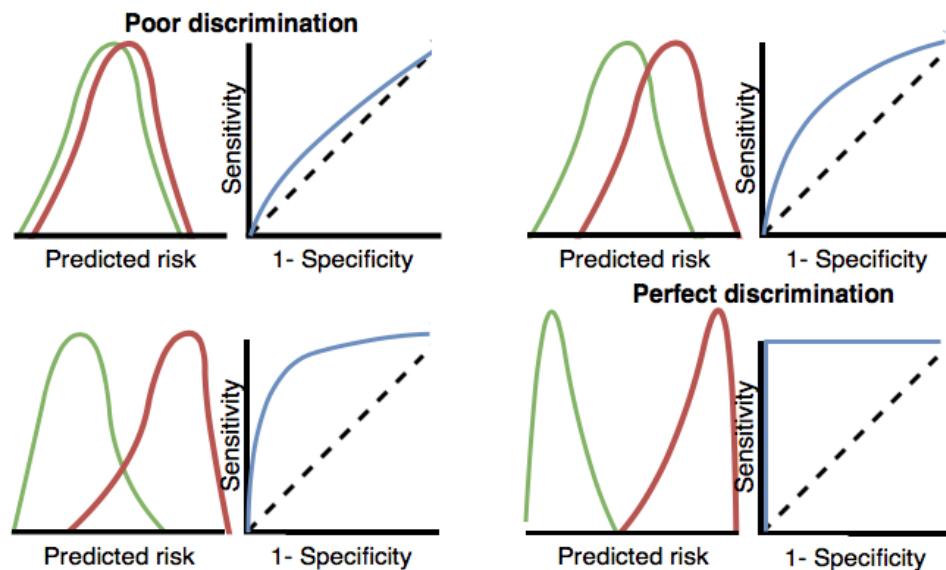
- $\square\square$ :  $(TP + TN) / (TP + TN + FP + FN)$

- $\square\Box$ :  $TP/(TP + FN)$
  - $\square\Box\Box$ :  $TN/(TN + FP)$
  - $\square\Box\Box\Box$ :  $TP/(TP + FP)$

三

14.11 AUC 0.5

AUC  14.3



AUC AUPRC AUPRC x

# Calibration Curves

14.5

14.1

## 14.5.1

ACE

14.5.2



14.5.3

14.3 

Boosting Machines □ GBM □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □

14.5.4

## 14.5.5

□□□□□ 14.5.5□□

□ : □ □ □ □ □ □ □ □ □ □ □ □ □

14.6 ATLAS

ATLAS Prediction 4

14.6.1

ATLAS  
Appendix B.1 Appendix  
Add  
Add  
Add  
Add  
14.4

14.6.2

□ □ □ □ □

Model Settings ATLAS

Prediction Problem Settings	
<p><b>Target Cohorts</b></p> <p>Show <select>10</select> entries</p> <p><a href="#">Remove</a> <a href="#">Name</a></p> <p> New users of ACE inhibitors as first-line monotherapy for hypertension</p>	<a href="#">+ Add Target Cohort</a>
Showing 1 to 1 of 1 entries	Previous <a href="#">1</a> Next
<p><b>Outcome Cohorts</b></p> <p>Show <select>10</select> entries</p> <p><a href="#">Remove</a> <a href="#">Name</a></p> <p> Angioedema outcome</p>	<a href="#">+ Add Outcome Cohort</a>
Showing 1 to 1 of 1 entries	Previous <a href="#">1</a> Next

Figure 14.4: □□□□□□.

5.5.5.5

Add Covariate Settings

**concepts do you want to include in baseline covariates in the patient-level prediction**

**model? (Leave blank if you want to include everything)**  

**descendant concepts be added to the list of included concepts?**       yes

**concepts do you want to exclude in baseline covariates in the patient-level prediction**

**comma delimited list of covariate IDs that should be restricted to**  ID  ID

14.6□□□□□□□□□

- □□: □□□□□□□□□□□□□□□□□
  - □□: □□□□□□□□
  - □□□□□□: 5□□□□□□□□□□□□□0-4□5-9□...□95+□
  - □□: □□□□□□□□□□
  - □□: □□□□□□□□□□
  - □□□□□□□□: □□□□□□□□□□□□□□□□□□
  - □□□□□□□□: □□□□□□□□□□□□□□□□□□
  - □□□□□□□□: [□□□□□□□□□□□□] □□□□□□□□□□
  - □□□□□□□□: [□□□□□□□□□□□□] □□□□□□□□□□
  - □□□□□□: □□□□□□□□□□□
  - □□□□□□□□□□: [□□□□□□□□□□] □□□□□□□□□□□□□□

**Gradient Boosting Machine Model Settings**  
Use the options below to edit the model settings

The boosting learn rate (default = 0.01,0.1):

Boosting learn rate	Action
0.001	<a href="#">Remove</a>
0.01	<a href="#">Remove</a>
0.1	<a href="#">Remove</a>
0.9	<a href="#">Remove</a>

[Add](#) [Reset to default](#)

Maximum number of interactions - a large value will lead to slow model training (default = 4,6,17):

Maximum number of interactions	Action
4	<a href="#">Remove</a>
7	<a href="#">Remove</a>
10	<a href="#">Remove</a>

[Add](#) [Reset to default](#)

The minimum number of rows required at each end node of the tree (default = 20):

Minimum number of rows	Action
20	<a href="#">Remove</a>

[Add](#) Using default

The number of trees to build (default = 10,100):

Trees to build	Action
5000	<a href="#">Remove</a>

[Add](#) [Reset to default](#)

The number of computer threads to use (how many cores do you have?) (default = 20):

20	Using default
----	---------------

Figure 14.5: □□□□□□□□□□□□

What concepts do you want to include in baseline covariates in the propensity score model? (Leave blank if you want to include everything)




Should descendant concepts be added to the list of included concepts?

 No ▾

What concepts do you want to exclude in baseline covariates in the propensity score model? (Leave blank if you want to include everything)




Should descendant concepts be added to the list of excluded concepts?

 No ▾

A comma delimited list of covariate IDs that should be restricted to:

Figure 14.6: Configuration of baseline covariates.

14.7 Configuration of baseline covariates.

### Select Covariates

	Gender	Age	Age Groups	Race	Ethnicity	Index Year	Index Month	Prior Observation Time	Post Observation Time	Time In Cohort	Index Year & Month
Demographics	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

Figure 14.7: Configuration of baseline covariates.

14.8 Configuration of baseline covariates.

- Age: [-365, 0]
- [ -365, -180 ]
- [ -365, -30 ]

14.8 Configuration of time bound covariates.

### Time bound covariates

Set the time windows for the time bound covariates in days relative to the cohort index

	Any Time Prior	Long Term	Medium Term	Short Term	End Days
Time Windows	All Time	-365	-180	-30	0

Figure 14.8: Configuration of time bound covariates.

14.9 Configuration of time bound covariates.

- Condition: ID
- Condition group: ID
- Drug: ID
- Drug group: ID

□□□□□□□□□□□□□□□□□□□□□□□ 14.9 □□□□□□□□□□□□□

Set the time bound era covariates

Domain	Any Time Prior	Long Term (-365 days)	Medium Term (-180 days)	Short Term (-30 days)	Overlapping	Era Start		
						Long Term (-365 days)	Medium Term (-180 days)	Short Term (-30 days)
Condition	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Condition Group	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Drug	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Drug Group	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

Figure 14.9: □□□□□□□

- Condition:      ID
  - Condition Primary Inpatient:
  - Drug:
  - Procedure:
  - Measurement:

14.7 R

ATLAS R Patient  
CDM

14.7.1

11 Appendix (Appendix B.1) (Appendix B.4) ACE ID 1 ID 2

14.7.2

PatientLevelPrediction DatabaseConnector

```
library(PatientLevelPrediction)
connDetails <- createConnectionDetails(dbms = "postgresql",
                                         server = "localhost/ohdsi",
                                         user = "joe",
                                         password = "supersecret")

cdmDbSchema <- "my_cdm_data"
cohortsDbSchema <- "scratch"
```

```
cohortsDbTable <- "my_cohorts"  
cdmVersion <- "5"
```

```
4 cdmDbSchema cohortsDbSchema cohortsDbTable CDM  
SQL Server  
<- "my_cdm_data.dbo"
```

```
sql <- paste("SELECT cohort_definition_id, COUNT(*) AS count",  
"FROM @cohortsDbSchema.cohortsDbTable",  
"GROUP BY cohort_definition_id")  
conn <- connect(connDetails)  
renderTranslateQuerySql(connection = conn,  
                         sql = sql,  
                         cohortsDbSchema = cohortsDbSchema,  
                         cohortsDbTable = cohortsDbTable)
```

```
##   cohort_definition_id   count  
## 1                      1 527616  
## 2                      2  3201
```

PatientLevelPrediction Features

```
covariateSettings <- createCovariateSettings(  
  useDemographicsGender = TRUE,  
  useDemographicsAge = TRUE,  
  useConditionGroupEraLongTerm = TRUE,  
  useConditionGroupEraAnyTimePrior = TRUE,  
  useDrugGroupEraLongTerm = TRUE,  
  useDrugGroupEraAnyTimePrior = TRUE,  
  useVisitConceptCountLongTerm = TRUE,  
  longTermStartDays = -365,  
  endDays = -1)
```

```
getPlpData
```

```
plpData <- getPlpData(connectionDetails = connDetails,  
                        cdmDatabaseSchema = cdmDbSchema,  
                        cohortDatabaseSchema = cohortsDbSchema,  
                        cohortTable = cohortsDbSchema,  
                        cohortId = 1,  
                        covariateSettings = covariateSettings,  
                        outcomeDatabaseSchema = cohortsDbSchema,  
                        outcomeTable = cohortsDbSchema,  
                        outcomeIds = 2,
```

```
    sampleSize = 10000  
)
```

getPlpData PatientLevelPrediction  
plpData

```
savePlpData(plpData, "angio_in_ace_data")
```

loadPlpData()

### 14.7.3

```
2  
= 30  
riskWindowEnd = 365  
= TRUE
```

14,7,4

Gradient Boosting Machine ntr

```
= c(100,200), maxDepth = 4, nntrees = 100, maxDepth = 4, nntrees = 200, maxDepth = 4
```

```
runPlP plpData  
25% testSplit / testFraction
```

```
gbmResults <- runPlp(population = population,
                      plpData = plpData,
                      modelSettings = gbmModel,
                      testSplit = 'person',
                      testFraction = 0.25,
                      nfold = 2,
                      splitSeed = 1234)
```

runPlp() plpData() plpResults() plpPlots() evaluation()  
()

```
savePlpModel(gbmResults$model, dirPath = "model")
```

```
plpModel <- loadPlpModel("model")
```

```
savePlpResult(gbmResults, location = "gbmResults")
```

```
gbmResults <- loadPlpResult("gbmResults")
```

14.7.5

runPlp  
= gbmResults) Shiny  
14.10

A horizontal row of 20 empty square boxes for writing responses.

```
plotPlp(gbmResults, "plots")
```

Section 14.4.2

14.7.6

viewPlp R runPLp ATLAS

F

runPLp

ATLAS

1

```
plpResult <- loadPlpResult(file.path(outputFolder,  
                                      'Analysis_1',  
                                      'plpResult'))
```

Analysis 1

□□□□□□□□□ Shiny □□□□□□□□□□□□□□□□

`viewPlp(plpResult)`

14.10 AUC 0.78 0.74

14.13  40 

□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□attrition□□□□□□□□□□

14.7.7

ATLAS

The screenshot shows a Shiny web application interface. At the top, there's a header bar with a logo, the URL "http://127.0.0.1:4888", an "Open in Browser" button, and a "Publish" dropdown. Below the header, the title "PatientLevelPrediction Explorer" is followed by tabs for "Internal Validation" and "External Validation". A navigation bar below the tabs includes "Evaluation Summary" (which is active, indicated by a blue border), "Characterization", "ROC", "Calibration", "Demographics", "Preference", "Box Plot", and "Settings".

The main content area is titled "Evaluation Summary". It features a table with 11 rows of data. The table has three columns: "Metric" (leftmost), "test" (middle), and "train" (rightmost). The "Metric" column lists various performance metrics. The "test" and "train" columns show their respective values.

Metric	test	train
1 AUC	0.72130	0.75348
2 AUC_lb95ci	0.70057	0.74215
3 AUC_ub95ci	0.74203	0.76482
4 AUPRC	0.10971	0.13571
5 BrierScaled	0.03755	0.04902
6 BrierScore	0.03355	0.03304
7 CalibrationIntercept.Intercept	-0.00089	-0.00813
8 CalibrationSlope.Gradient	1.02041	1.22457
9 outcomeCount	601.00000	1802.00000
10 populationSize	16685.00000	50054.00000
11 Incidence	3.60204	3.60011

At the bottom left, it says "Showing 1 to 11 of 11 entries". On the right, there are buttons for "Previous", a page number "1", and "Next".

Figure 14.10: Shiny

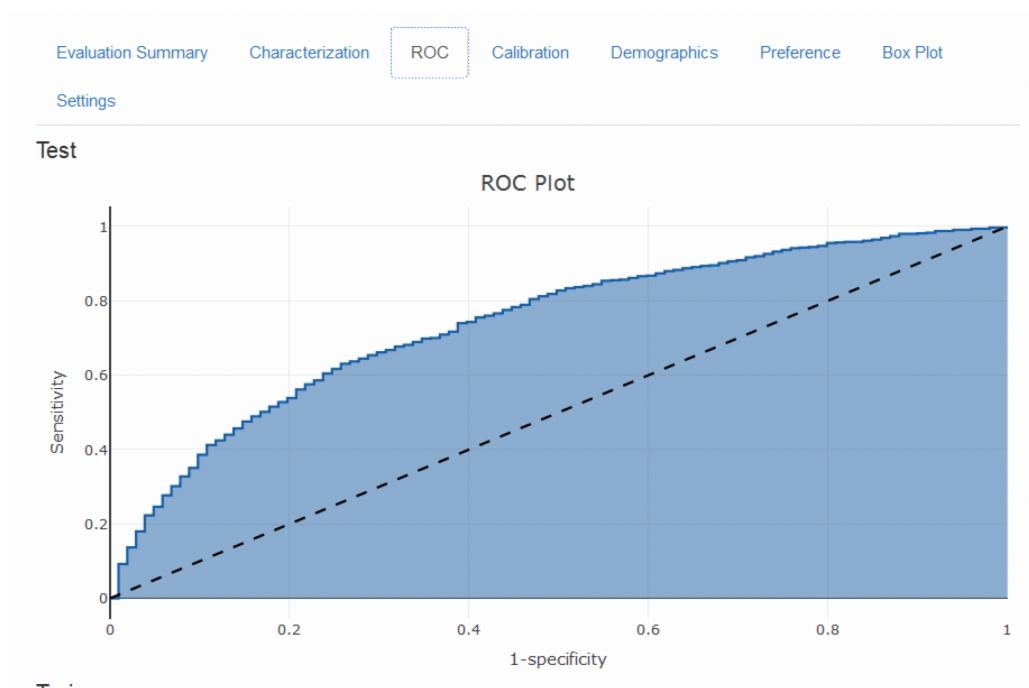


Figure 14.11: ROC

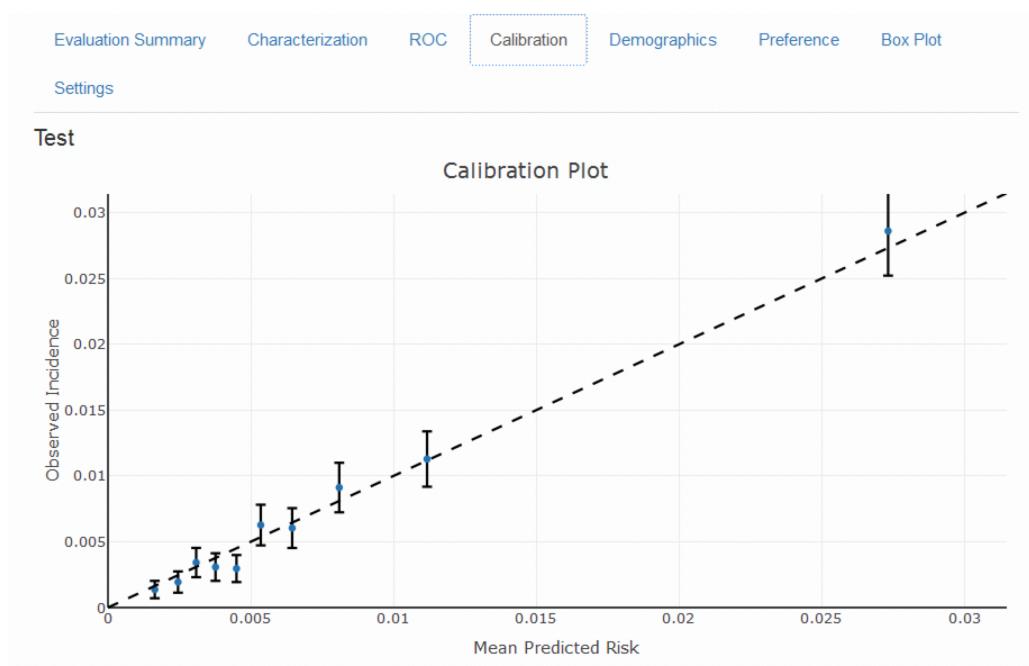


Figure 14.12:

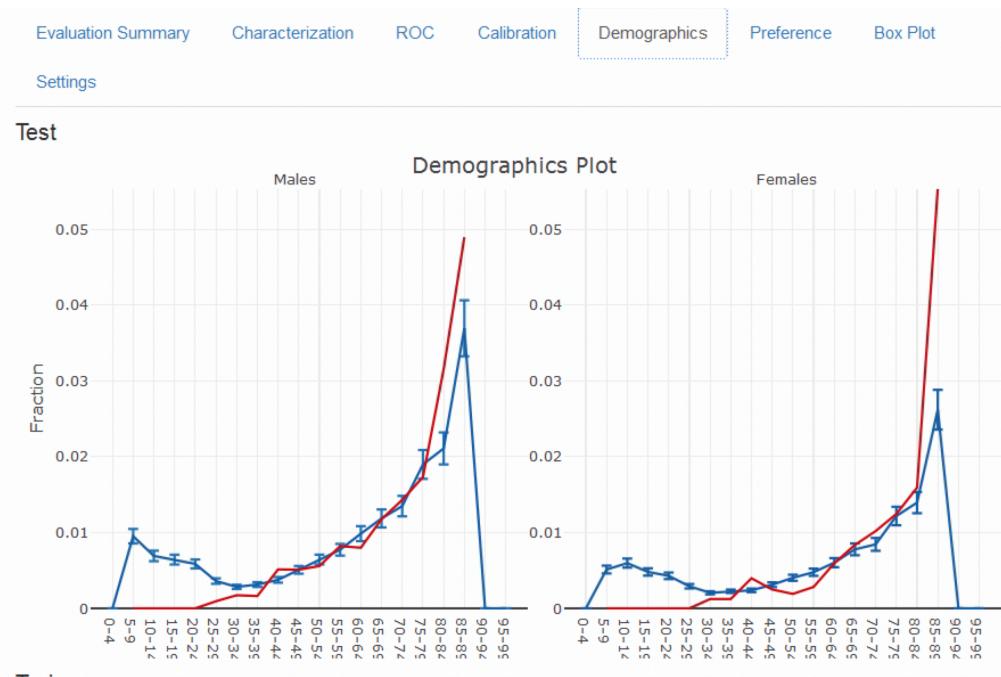


Figure 14.13: □□□□□□□□□□□□□□□□□□

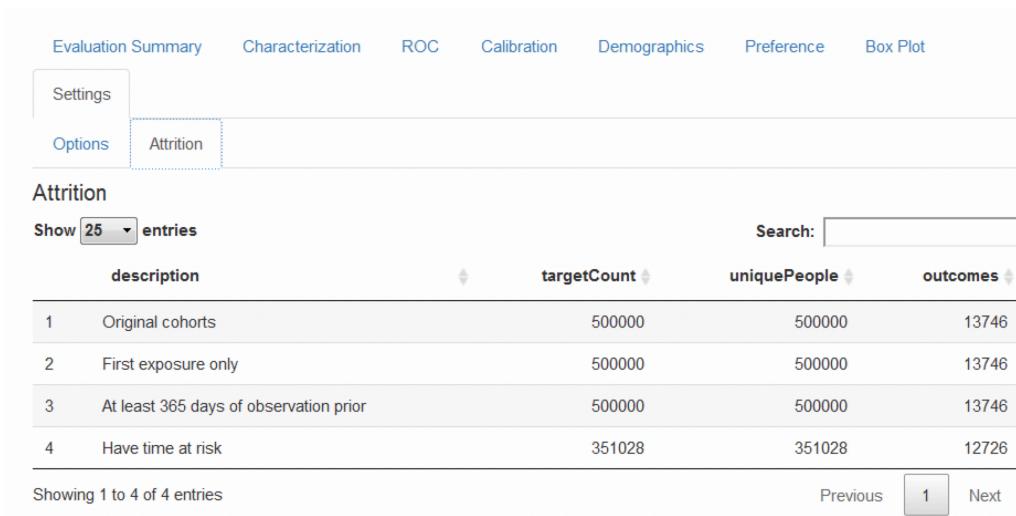


Figure 14.14: attrition

Shiny 14.15

Filters	Results	Model Settings	Population Settings	Covariate Settings									
	Show 10 entries											Search	
Development Database	Analysis	Dev	Val	T	O	Model	TAR start	TAR end	AUC	AUPRC	T Size	O Count	O Incidence (%)
	Analysis_1	Optum claims	Optum claims	New users of ACE inhibitors as first-line monotherapy for hypertension	Acute myocardial infarction events	Lasso Logistic Regression	1	365	0.74466	0.03094	87757	650	0.74066
Validation Database	Analysis_3	Optum claims	Optum claims	New users of ACE inhibitors as first-line monotherapy for hypertension	Angioedema events	Lasso Logistic Regression	1	365	0.60523	0.00254	87615	148	0.16892
Target Cohort	Analysis_5	Optum claims	Optum claims	New users of ACE inhibitors as first-line monotherapy for hypertension	Acute myocardial infarction events	Random forest	1	365	0.71667	0.03102	87757	650	0.74066
	Analysis_7	Optum claims	Optum claims	New users of ACE inhibitors as first-line monotherapy for hypertension	Angioedema events	Random forest	1	365	0.64163	0.02447	87615	148	0.16892

Figure 14.15: Shiny

A horizontal row of 20 empty square boxes, intended for children to practice writing their names.

- - 
  - AUC AUPRC

A horizontal row of 20 empty square boxes, intended for children to practice writing their names.

## Settings

Results Model Settings Population Settings Covariate Settings

## Model Settings: help

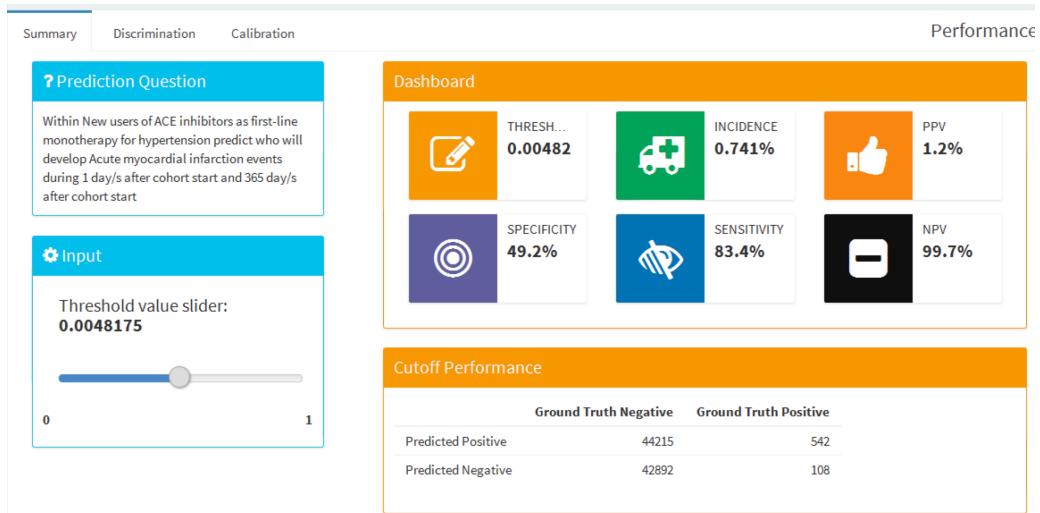
Show 10 entries

	Setting	Value
1	Model	lr_lasso
2	variance	0.01
3	seed	50975614

Showing 1 to 3 of 3 entries

Figure 14.16: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

14 17  Performance



**Figure 14.17:** □□□□□□□□□□□□□□□□□□□□□□□□□□

14.4.2 PPV 14.17

Discrimination ROC -  
14.18 ROC - ROC

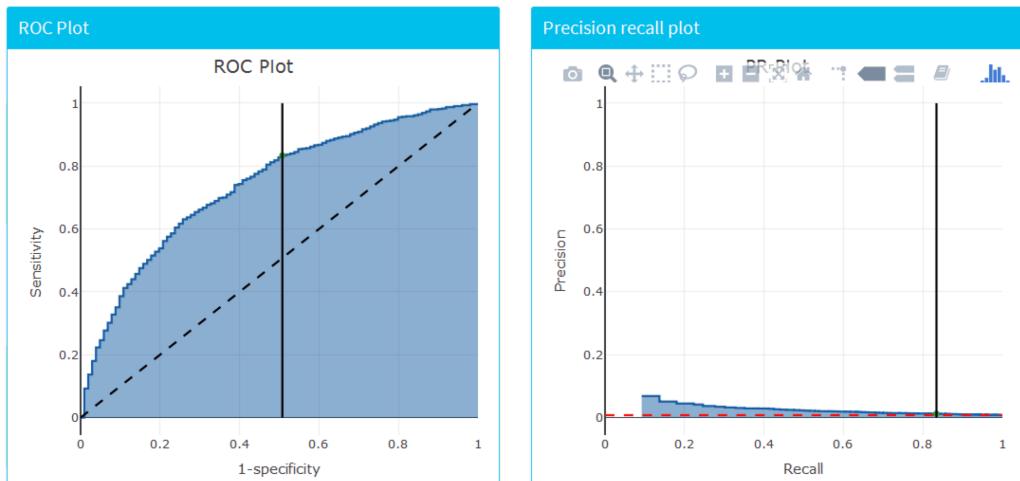


Figure 14.18: ROC curve - Confusion matrix

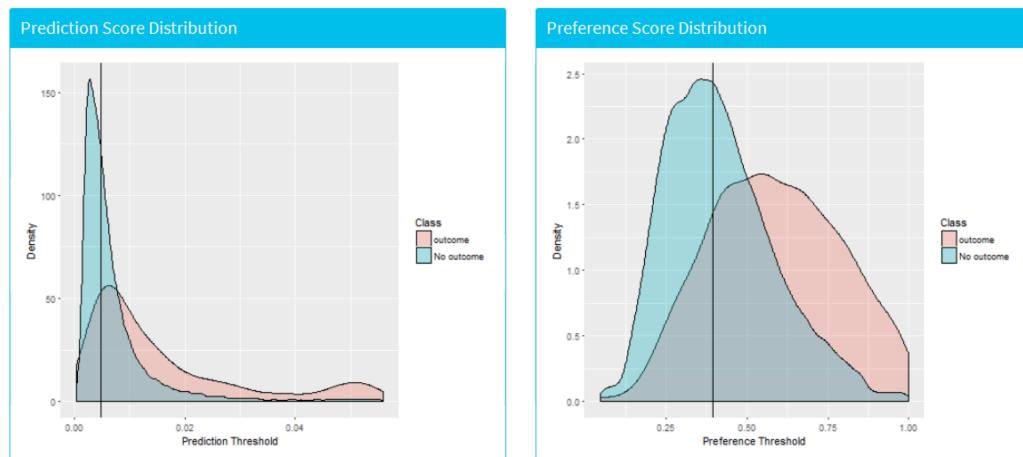
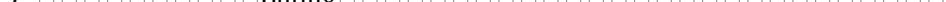
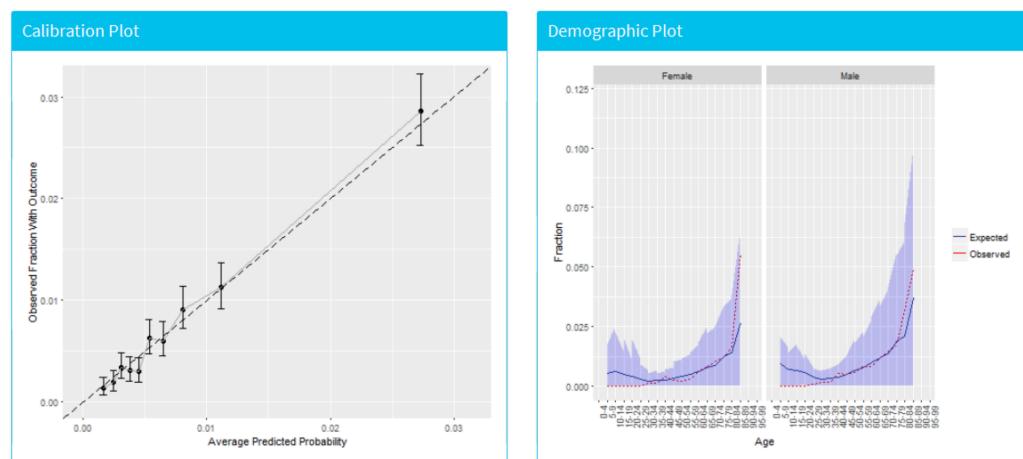


Figure 14.19: outing



□ □ □ □ □



14.21  14.22 

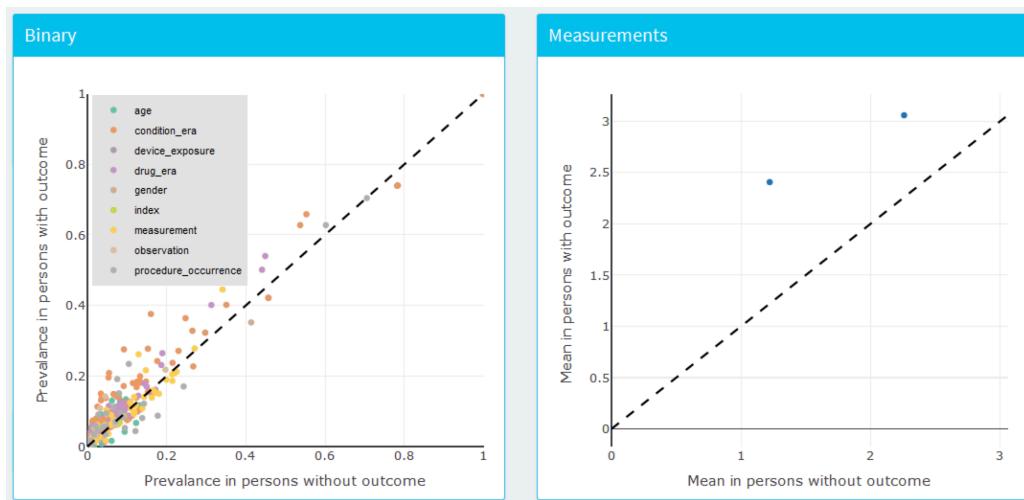


Figure 14.21: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

Model Table				
<a href="#">Download Model</a>				
Show	10	entries	Search:	
Covariate Name	Value	Outcome Mean	Non-outcome Mean	
1 age group: 00-04	0	0.0004	0.0001	
2 age group: 05-09	0	0	0.0003	
3 index month: 1	0	0.1307	0.1096	
4 observation during day -365 through 0 days relative to index: Domain	0	0.1188	0.0514	
5 Charlson index - Romano adaptation	0	2.4783	1.3817	
6 Diabetes Comorbidity Severity Index (DCSI)	0.1478	2.4056	1.2207	
7 CHADS2VASc	0.9279	3.0573	2.2576	
8 visit_occurrence concept count during day -365 through 0 concept_count relative to index	0	19.5263	13.8837	
9 age group: 10-14	0	0	0.001	
10 index month: 2	0	0.0934	0.0909	

**Figure 14.22:** □□□□□□□□□□



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14.22□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

## 14.8

### 14.8.1

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```
createPlpJournalDocument(plpResult = <your plp results>,
                         plpValidation = <your validation results>,
                         plpData = <your plp data>,
                         targetName = "<target population>",
                         outcomeName = "<outcome>",
                         table1 = F,
                         connectionDetails = NULL,
                         includeTrain = FALSE,
                         includeTest = TRUE,
                         includePredictionPicture = TRUE,
                         includeAttritionPlot = TRUE,
                         outputLocation = "<your location>")
```

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## 14.9



- □□
- □□
- PatientLevelPrediction□□□□□□□OMOP-CDM□□□□□□□□□□□□□□□□□
- □□
- OHDSI□□

## 14.10

□□□□

□□□□□□□□□□□□□□□□□ 9.4.5□□□□□□□□□□□ R□R-Studio□□□□Java□□□□□□□□□□□

```
install.packages(c("SqlRender", "DatabaseConnector", "remotes"))
remotes::install_github("ohdsi/Eunomia", ref = "v1.0.0")
remotes::install_github("ohdsi/PatientLevelPrediction")
```

Eunomia CDM R

```
connectionDetails <- Eunomia::getEunomiaConnectionDetails()
```

CDM main Eunomia

```
Eunomia::createCohorts(connectionDetails)
```

NSAIDs GI

NSAID COHORT\_DEFINITION\_ID 4 GI COHORT\_D

14.1. PatientLevelPrediction R CDM PL

14.2. createStudyPopulation

14.3. LASSO Shiny

E.8



## **Part IV**

---



# Chapter 15

□□□□□□□: *Patrick Ryan & Jon Duke*

15.1

Desired attribute	Question	Researcher	Data	Analysis	Result
<b>Repeatable</b>	Identical	Identical	Identical	Identical	= Identical
<b>Reproducible</b>	Identical	Different	Identical	Identical	= Identical
<b>Replicable</b>	Identical	Same or different	Similar	Identical	= Similar
<b>Generalizable</b>	Identical	Same or different	Different	Identical	= Similar
<b>Robust</b>	Identical	Same or different	Same or different	Different	= Similar
<b>Calibrated</b>	Similar (controls)	Identical	Identical	Identical	= Statistically consistent

Figure 15.1: □□□□□□□□□□□□□□□□

Austin Bradford Hill (1965) et al., 2013) et al., 2016, 2018a,b)

15.2

(Bots et al., 2010; Hersh et al., 2013; Sherman et al., 2016) et al., 2012; Liaw et al., 2013; Weiskopf and Weng, 2013) OHDSI CDM OHDSI (Huser et al., 2016; Kahn et al., 2015; Callahan et al., 2017; Yoon et al., 2016)

A horizontal row of 30 small, empty rectangular boxes, likely used for input fields or placeholder text in a form.

□: □ □ □ □ □ 4 □ □ □ □ □ □ □ □ □

A horizontal row of 15 empty square boxes, intended for children to draw or color in.

A horizontal row of 20 empty square boxes, intended for students to write their answers in a handwriting practice exercise.

15.3

##



# Chapter 16

□□□□□: Martijn Schuemie, Vojtech Huser & Clair Blacketer

A horizontal row of 20 empty square boxes, intended for children to write their names in, likely as part of a classroom activity or name recognition exercise.

□□□□□ DQ□□□□□□□□□□□ (Roebuck, 2012):

□□□□□□□□ DQ □□□□□□□□□□□□□□□□□□□□□□□□ DQ □□□□□□□□

16.1

## Chapter 15 and Johnson (2003) DQ



16.1.2 DQ (Defalco et al., 2013; Makadia and Ryan, 2014; Matcho et al., 2014; Voss et al., 2015a,b; Hripcak et al., 2018) CDM

3 OHDSI DQ  
17 Chapter 18 Chapter 19 #

Kahn et al.  
(2016) DQ 3



□ □ □ □ □ □ □ □ □ **2** □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □



## 16.1.1

ACHILLES (Huser et al., 2018) ACHILLES (Huser et al., 2016) ACHILLES ATLAS

ATLAS 

(DQD) ACHILLES CDM CDM  
1,500 FAIL 16.1.1

			□ □	□ □
0.99	MEASUREMENT □□□□□ measurement_source_value □□□□□□□□□□□□□□□□□□□□□□□□□ 0 □□□□□□□□□□□	0.30		FAIL
0.09	DRUG_ERA □□□□□ drug_concept_id □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□	0.10		PASS
0.02	DRUG_EXPOSURE □□□□□ DRUG_EXPOSURE_END_DATE □□□□□□□ DRUG_EXPOSURE_START_DATE □□□□□□□□□□□□□□□□□□□□□□□□□	0.05		PASS
0.00	PROCEDURE_OCCURRENCE □□□□□ procedure_occurrence_id □□□□□□□□□□□□□□□□□□□□□□□□□	0.00		PASS

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**CDM** □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□  
**CDM** □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□  
**ID** □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□  
**ID** □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□



ACHILLES □ DQD □ CDM □□□□□□□□□□□□□□□□□□□□  
 CDM □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

### 16.1.2 ETL

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**CDM** □□□□□ ETL (Extract-Transform-Load) □□□□□□□□□□□□  
 □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□ ETL  
  
 ETL □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□  
 7 □□□□□ Rabbit-in-a-Hat □□□□□□□□□□□□□□□□□□□□□□  
 □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□ R  
 CDM □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

```
source("Framework.R")
declareTest(101, "Person gender mappings")
add_enrollment(member_id = "M000000102", gender_of_member = "male")
add_enrollment(member_id = "M000000103", gender_of_member = "female")
expect_person(PERSON_ID = 102, GENDER_CONCEPT_ID = 8507)
expect_person(PERSON_ID = 103, GENDER_CONCEPT_ID = 8532)
```

Rabbit-in-a-Hat gender mappings ENROLLMENT  
Rabbit-in-a-Hat add\_enrollment MEMBER\_ID  
GENDER\_OF\_MEMBER 2 ETL  
PERSON 2

ENROLLMENT White Rabbit

ETL SQL ETL SQL

```
insertSql <- generateInsertSql(databaseSchema = "source_schema")
testSql <- generateTestSql(databaseSchema = "cdm_test_schema")
```

□□□□□□□□□ 16.1 □□□□□□□□□

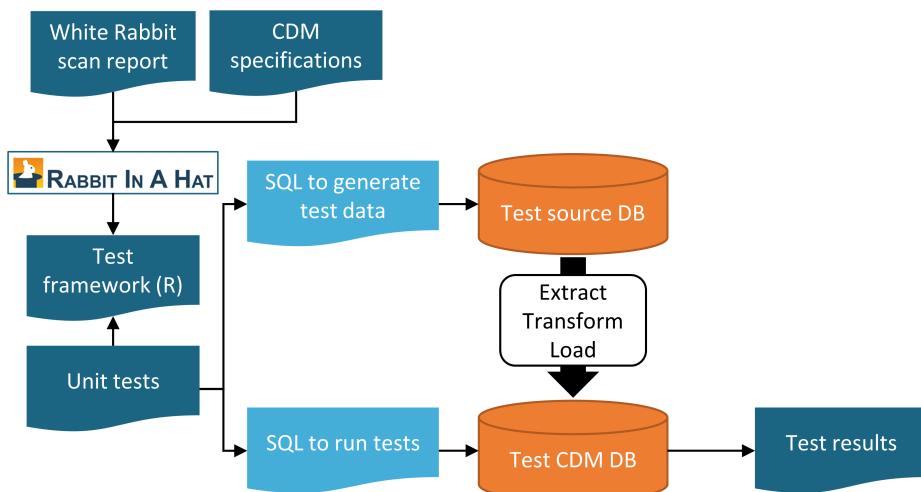


Figure 16.1: Rabbit-in-a-Hat ☐☐☐☐☐☐☐☐☐☐☐☐☐ ETL (Extract-Transform-Load) ☐☐☐☐☐☐☐☐☐☐☐

□: ETL □□□□□□□□□□

ID	□ □	□ □
101	Person gender mappings	PASS
101	Person gender mappings	PASS

ETL

16.2

DQ DQ ACHILLES DQ

## 16.2.1

Vocabulary  
Vocabulary<sup>1</sup>  
MethodEvaluation R  
checkCohortSourceCodes ATLAS  
16.2  
440383 (“ ”) 3  
9 3.11 ICD-10 F32.8 F32.89  
9 ICD-10  
10 ICD-9 ICD-  
10 F32.9

% per month	Max monthly %	Person count	Description
	26.81	92,019,885	<b>Depressive Disorder</b>
	6.64	15,969,198	<b>Depressive disorder</b> 440383
	6.64	15,686,275	311 (ICD9CM) Depressive disorder, not elsewhere classified
	0.46	188,230	F328 (ICD10CM) Other depressive episodes
	0.38	94,693	F3289 (ICD10CM) Other specified depressive episodes
	3.10	12,010,783	<b>Adjustment disorder with mixed emotional features</b> 433454
	3.07	9,839,712	30928 (ICD9CM) Adjustment disorder with mixed anxiety and depressed mood
	3.03	2,049,618	F4323 (ICD10CM) Adjustment disorder with mixed anxiety and depressed mood
	0.04	121,453	3091 (ICD9CM) Prolonged depressive reaction
	3.17	9,237,192	<b>Dysthymia</b> 433440

Figure 16.2: checkCohortSourceCodes

R findOrphanSourceCodes  
16.3 ICD-10 J85.0 4324261

---

<sup>1</sup><https://github.com/OHDSI/Vocabulary-v5.0/issues>

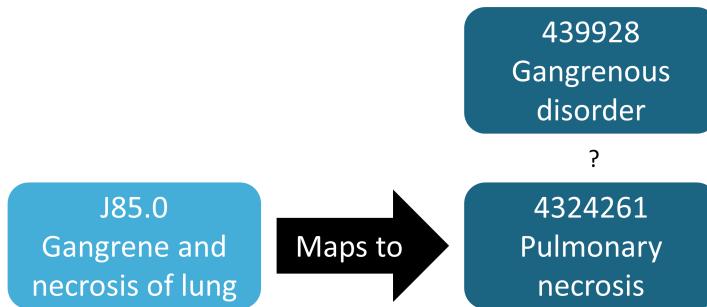


Figure 16.3: □□□□□□□□□□□□□□□□

## 16.3 ACHILLES

□□□□□ CDM □□□□□□□□□□ ACHILLES □□□□□□□□□□

```
library(Achilles)
connDetails <- createConnectionDetails(dbms = "postgresql",
                                         server = "localhost/ohdsi",
                                         user = "joe",
                                         password = "supersecret")

cdmDbSchema <- "my_cdm_data"
cdmVersion <- "5.3.0"
```

ACHILLES

```
result <- achilles(connectionDetails,
                     c
## Data Quality Dashboard  {#dqInPractice}
```

CDM Data Quality Dashboard \eref{achillesInPractice} CDM

```
cdmDbSchema <- "my_cdm_data.dbo"
```

Dashboard

```
viewDqDashboard(jsonPath)
```

Dashboard 16.4 Kahn

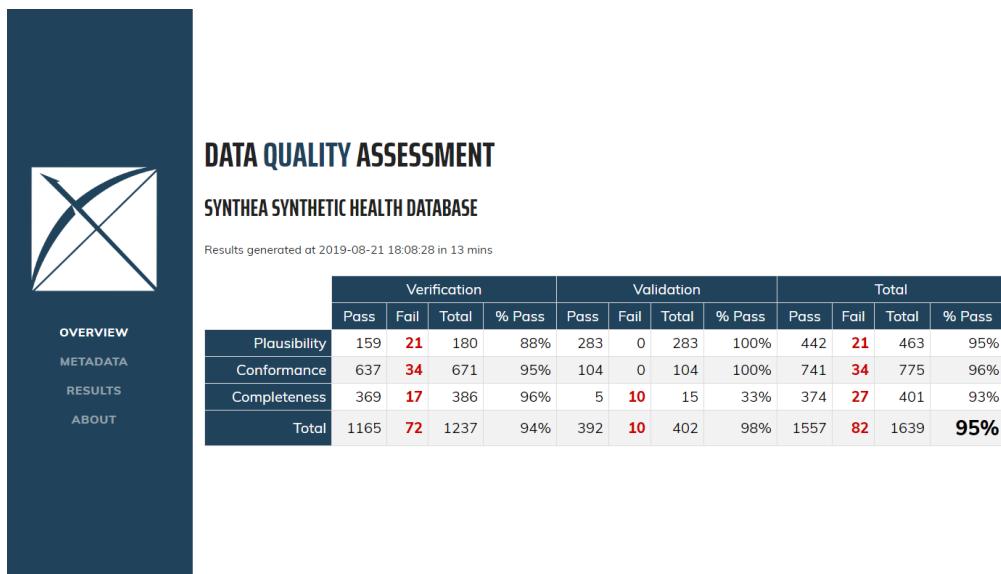


Figure 16.4: Data Quality Dashboard

16.4

#### **B.4**

```
library(MethodEvaluation)
json <- readChar("cohort.json", file.info("cohort.json")$size)
sql <- readChar("cohort.sql", file.info("cohort.sql")$size)
checkCohortSourceCodes(connectionDetails,
```

## RESULTS

### SYNTHIA SYNTHETIC HEALTH DATABASE

Results generated at 2019-08-22 14:15:06 in 29 mins

							Column visibility	CSV
						Search:		
STATUS	CONTEXT	CATEGORY	SUBCATEGORY	LEVEL	DESCRIPTION	% RECORDS		
				FIELD				
+	FAIL	Verification	Plausibility	Atemporal	FIELD The number and percent of records with a value in the gap_days field of the DRUG_ERA table less than 0. (Threshold=0%).	24.07%		
+	FAIL	Verification	Completeness	None	FIELD The number and percent of records with a value of 0 in the standard concept field race_concept_id in the PERSON table. (Threshold=0%).	16.74%		
+	FAIL	Verification	Conformance	Relational	FIELD The number and percent of records that have a value in the ethnicity_concept_id field in the PERSON table that does not exist in the CONCEPT table. (Threshold=0%).	16.15%		
+	PASS	Verification	Completeness	None	FIELD The number and percent of records with a NULL value in the condition_end_date of the CONDITION_OCCURRENCE. (Threshold=100%).	13.24%		
+	PASS	Verification	Completeness	None	FIELD The number and percent of records with a NULL value in the condition_end_datetime of the CONDITION_OCCURRENCE. (Threshold=100%).	13.24%		

Showing 71 to 75 of 1,327 entries (filtered from 1,639 total entries)

Previous 1 ... 14 15 16 ... 266 Next

Figure 16.5: Data Quality Dashboard

```
cdmDatabaseSchema = cdmDbSchema,
cohortJson = json,
cohortSql = sql,
outputFile = "output.html")
```

16.6

9 ICD-10

Angioedema

```
orphans <- findOrphanSourceCodes(connectionDetails,
                                    cdmDatabaseSchema = cdmDbSchema,
                                    conceptName = "Angioedema",
                                    conceptSynonyms = c("Angioneurotic edema",
                                                       "Giant hives",
                                                       "Giant urticaria",
                                                       "Periodic edema"))
```

View(orphans)

□□□

□□

□□ID

□□□□□□

T78.3XXS Angioneurotic edema, sequela  
10002425 Angioedemas

ICD10CM 508  
MedDRA 0

□□□	□□	□□ID	□□□□□□□□
148774	Angioneurotic Edema of Larynx	CIEL	0
402383003	Idiopathic urticaria and/or angioedema	SNOMED	0
232437009	Angioneurotic edema of larynx	SNOMED	0
10002472	Angioneurotic edema, not elsewhere classified	MedDRA	0

"Angioneurotic edema, sequelae"

16.5






16.6

```
install.packages(c("SqlRender", "DatabaseConnector", "remotes"))
remotes::install_github("ohdsi/Achilles")
remotes::install_github("ohdsi/DataQualityDashboard")
remotes::install_github("ohdsi/Eunomia", ref = "v1.0.0")
```

Eunomia  R  CDM 

```
connectionDetails <- Eunomia:::getEunomiaConnectionDetails()
```

# CDM main

□□ 16.1. Eunomia □□□□□□□□□ ACHILLES □□□□□□□□□

□□ 16.2. Eunomia □□□□□□□□□ Data Quality Dashboard □□□□□□□□□

□ □ 16.3. DQD □

% per month	Max monthly %	Person count	Description
	60.60	24,189,656	<b>Inpatient or ER visit</b>
	39.50	15,003,249	<b>Emergency Room Visit 9203</b>
	39.50	15,003,249	ER (None) No matching concept
	23.90	9,186,407	<b>Inpatient Visit 9201</b>
	23.90	9,186,407	IP (None) No matching concept
	0.27	76,711	<b>Angioedema</b>
	0.27	76,711	<b>Angioedema 432791</b>
	0.26	64,726	9951 (ICD9CM) Angioneurotic edema, not elsewhere classified
	0.20	8,822	T783XXA (ICD10CM) Angioneurotic edema, initial encounter
	0.09	3,163	T783XXD (ICD10CM) Angioneurotic edema, subsequent encounter

# Chapter 17

□□□□□□□: Joel Swerdel, Seng Chan You, Ray Chen & Patrick Ryan

OHDSI 

17.1

```
ACE ACE  
## {#CohortValidation}
```

# OHDSI Chapter 13 “ACE Valuator”

17.1.1

		Gold Standard	
		True	False
Cohort Definition	True	True Positive	False Positive
	False	False Negative	True Negative

Figure 17.1: □□□□□□□□□□□□□□

incorrect

1. **PPV** =  $\frac{\text{True Positives}}{\text{True Positives} + \text{False Positives}}$
  2. **NPV** =  $\frac{\text{True Negatives}}{\text{True Negatives} + \text{False Negatives}}$
  3. **PPV** =  $\frac{\text{True Positives}}{\text{True Positives} + \text{False Positives}}$
  4. **NPV** =  $\frac{\text{True Negatives}}{\text{True Negatives} + \text{False Negatives}}$

100% Rubbo et al. (2015) 33 1



PPV 1

6 

17.1.2

1. OHDSI IRB
  2. ATHENA ATLAS  
4163261 ID  
4314337 ID 4048666
  3. ATLAS PERS
  4. EHR
  5. 1
  - 6.
  7. 1 2 3
  8. PPV
  9. CUIMC
  10. PPV

MI Rubbo et al. (2015) # PheEvaluator

OHDSI et al., 2019) PheValuator Patient Level Prediction



Patient-Level Prediction (Chapter 14 et al., 2013)

### 17.1.3 PheEvaluator

---

<sup>1</sup><https://github.com/OHDSI/PheEvaluator>

□ □ □ □ 1: xSpec □ □ □ □ □ □

17.2 ATLAS MI xSpec

Cohort #10934

MI xSpec Cohort

Definition ? Concept Sets Generation Reporting Export

[460] MI xSpec Model

### Cohort Entry Events

Events having any of the following criteria:

a condition occurrence of [460] Myocardial Infarction ▼ + Add attribute...▼ Delete Criteria

with continuous observation of at least 365 ▼ days before and 0 ▼ days after event index date

Limit initial events to: earliest event ▼ per person.

Restrict initial events to:

having all ▼ of the following criteria: + Add criteria to group...▼ Delete Criteria

with at least 1 ▼ using all occurrences of:

a condition occurrence of [460] Myocardial Infarction ▼ + Add attribute...▼ Delete Criteria

X with a Visit occurrence of: Inpatient Visit Add Import

where event starts between 0 ▼ days Before ▼ and 5 ▼ days After ▼ index start date [add additional constraint](#)

restrict to the same visit occurrence  
 allow events from outside observation period

and with at least 4 ▼ using all occurrences of:

a condition occurrence of [460] Myocardial Infarction ▼ + Add attribute...▼ Delete Criteria

where event starts between 1 ▼ days After ▼ and 365 ▼ days After ▼ index start date [add additional constraint](#)

restrict to the same visit occurrence  
 allow events from outside observation period

Limit initial events to: earliest event ▼ per person.

Remove initial event restriction

□□□□ 2: xSens □□□□□□

17.3 ATLAS MI xSens

□ □ □ □ **3:** □ □ □ □ □ □ □

```
createPhenoModel
```

The screenshot shows the 'Cohort Entry Events' section of a software application. At the top, there's a navigation bar with tabs: 'Definition' (selected), 'Concept Sets', 'Generation', 'Reporting', and 'Export'. To the right of the tabs are several icons: a green folder, a blue 'X', a white square with a blue border, a blue gear, and a red trash can. Below the tabs is a search bar containing the text 'MI xSens Cohort'. The main area is titled 'Cohort Entry Events' and contains the following text: 'Events having any of the following criteria:' followed by a list of conditions. One condition is shown: 'a condition occurrence of [460] Myocardial Infarction ▾'. To the right of this list are two buttons: '+ Add Initial Event ▾' and 'Delete Criteria'. Below this, there are two input fields: 'with continuous observation of at least [0 ▾] days before and [0 ▾] days after event index date' and 'Limit initial events to: earliest event ▾ per person.' At the bottom left is a green button labeled 'Restrict initial events'.

Figure 17.3: xSens

```

setwd("c:/temp")
library(PheEvaluator)
connectionDetails <- createConnectionDetails(
  dbms = "postgresql",
  server = "localhost/ohdsi",
  user = "joe",
  password = "supersecret")

phenoTest <- createPhenoModel(
  connectionDetails = connectionDetails,
  xSpecCohort = 10934,
  cdmDatabaseSchema = "my_cdm_data",
  cohortDatabaseSchema = "my_results",
  cohortDatabaseTable = "cohort",
  outDatabaseSchema = "scratch.dbo", #
  trainOutFile = "5XMI_train",
  exclCohort = 1770120, #xSens
  prevCohort = 1770119, #
  modelAnalysisId = "20181206V1",
  excludedConcepts = c(312327, 314666),
  addDescendantsToExclude = TRUE,
  cdmShortName = "myCDM",
  mainPopnCohort = 0, #
  lowerAgeLimit = 18,
  upperAgeLimit = 100
)

```

```
upperAgeLimit = 90,  
gender = c(8507, 8532),  
startDate = "20100101",  
endDate = "20171231")
```

createEvalCohort □□□□ PatientLevelPrediction □□□□□ applyModel □□□□□□□□□□□□□

□□□□

```
setwd("c:/temp")
connectionDetails <- createConnectionDetails(
  dbms = "postgresql",
  server = "localhost/ohdsi",
  user = "joe",
  password = "supersecret")

evalCohort <- createEvalCohort(
  connectionDetails = connectionDetails,
  xSpecCohort = 10934,
  cdmDatabaseSchema = "my_cdm_data",
  cohortDatabaseSchema = "my_results",
  cohortDatabaseTable = "cohort",
  outDatabaseSchema = "scratch.dbo",
  testOutFile = "5XMI_eval",
  trainOutFile = "5XMI_train",
  modelAnalysisId = "20181206V1",
  evalAnalysisId = "20181206V1",
  cdmShortName = "myCDM",
  mainPopnCohort = 0,
  lowerAgeLimit = 18,
  upperAgeLimit = 90,
  gender = c(8507, 8532),
  startDate = "20100101",
  endDate = "20171231")
```

## {#GeneralizabilityOfEvidence}

**OHDSI** vs. **Other**  
vs. **Other** vs. **Other**  
vs. **Other**

17.2



- OHDSI
  - PheValuator
  - OHDSI



# Chapter 18

□□□□□□□: Martijn Schuemie

A horizontal row of 20 empty square boxes, intended for students to draw or write in.

18.1.1 CRM

9.1 OHDSI

Methods Library

## Library Methods

18.1

18.1.1

OHDSI R DatabaseCo

R□□□□□□□□□□□□□□ CDM□□□□□□□□□□□□□□ OHDSI

## Methods Library R

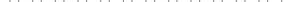
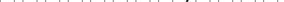
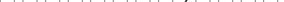
## 16.1.2

2000)  1

2008) 4

- `sampledPatients`: `sampledPatients <- takeSample(patients, sampleSize = 100)`
  - `1`: `1 <- sampledPatients[1]`

### 18.1.3

-  : 1  1 
  -  : 2  2 

## 18.1.4 Methods Library

OHDSI Methods Library

GitHub drat

OHDSI License V2 R C++ SQL Java OHDSI

18.1.5

GitHub git

OHDSI

- **4.3.2**: 4.3.2  
• **4.3.3**: 4.3.3  
• **4.4.0**: 4.4.0  
• **5.0.0**: 4.4.0

---

<sup>1</sup><https://github.com/OHDSI/CohortMethod/issues>

<sup>2</sup><http://forums.ohdsi.org/>

18.1.6

R 1

18.1.7

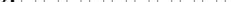
GitHub drat

18.1.8

OHDSI

18.1.9

OHDSI 

OHDSI  OHDSI  OHDSI 

18.1.10

OHDSI GitHub<sup>4</sup> GitHub https://github.com/security OHDSI

18.1.11

OHDSI GitHub GitHub https://github.com/security

18.2

18.2.1

18.2.2

<sup>3</sup><https://ohdsi.github.io/MethodsLibrary/>

<sup>4</sup><https://github.com/>

18.3



# Chapter 19

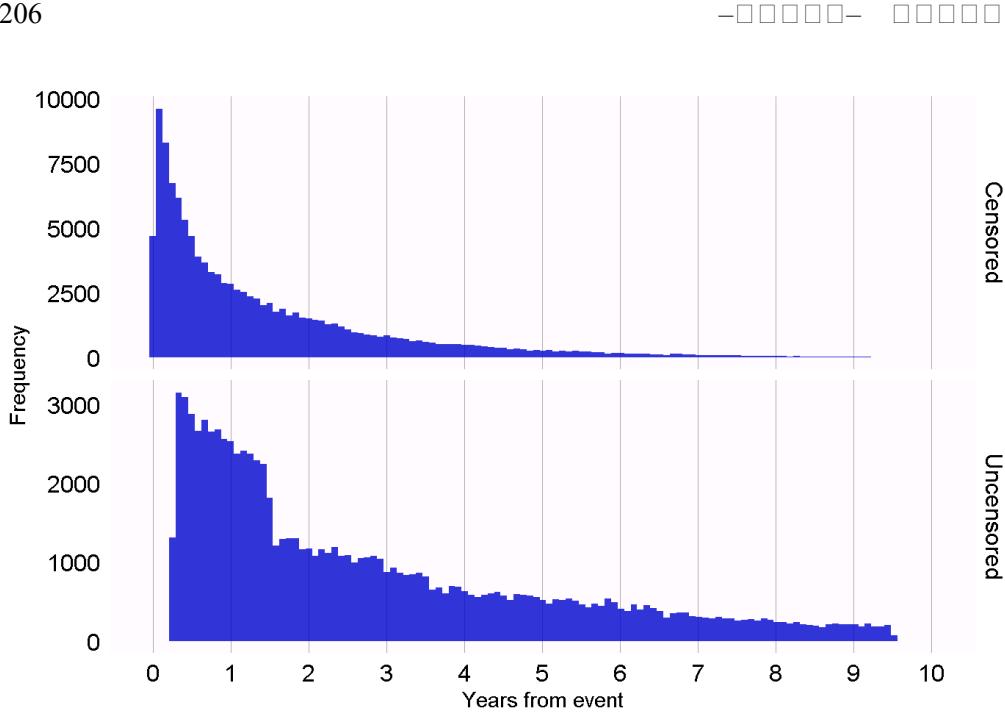
□□□□□: Martijn Schuemie

19.1

# OHDSI CohortMethod



19.2



## 19.2.1

(Lipsitch et al., 2010) (Zaadstra et al., 2008) MS



19.2.2

et al., 2014)

OHDSI

et al., 2013) 10

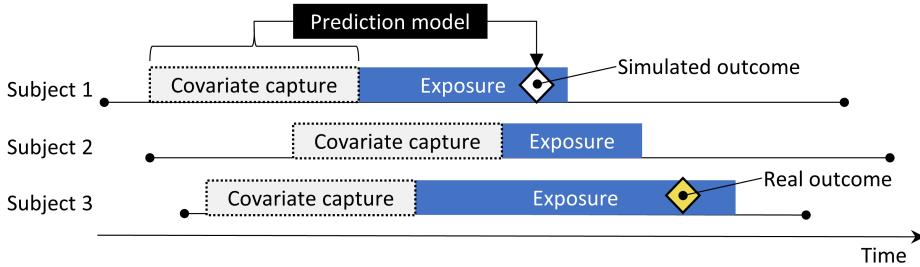


Figure 19.2: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

19.2.3

- **AUC**
  - 95
  - $1/( )^2$
  - **MSE**
  - $\mathbf{1} \alpha = 0.05$
  - $\mathbf{2} \alpha = 0.05$
  -

19.2.4 P

$\alpha = 0.05$  (et al., 2014)

$\hat{\theta}_i$   $\tau_i$   $i = 1, \dots, n$   
 $\theta_i$   $0$   $i$   $\beta_i$   $\theta_i$   
 $\beta_i$   $p$   $\hat{\theta}_i$   $\theta_i + \beta_i$   
 $\hat{\tau}_i^2$   $p$   $\beta_i$   
 $\beta_i$   $\mu$   $\sigma^2$   $\mu$   
 $\sigma^2$

A horizontal row of fifteen empty square boxes, intended for children to write their names in, likely as part of a classroom activity or name recognition exercise.

$$\beta_i \sim N(\mu, \sigma^2) \quad \square \square \square \quad \hat{\theta}_i \sim N(\theta_i + \beta_i, \tau_i^2)$$

$$L(\mu, \sigma | \theta, \tau) \propto \prod_{i=1}^n \int p(\hat{\theta}_i | \beta_i, \theta_i, \hat{\tau}_i) p(\beta_i | \mu, \sigma) d\beta_i$$

$\hat{\mu}$   $\hat{\sigma}$   $\hat{\theta}_{n+1}$   $\hat{\tau}_{n+1}$  {n##}

ACEi THZ

19.2.5

100

ACEi and THZ combined								Optimize	
Concept Set Expression		Included Concepts <span style="border: 1px solid black; padding: 2px;">14</span>		Included Source Codes		Explore Evidence		Export	Compare
Showing 1 to 14 of 14 entries							Search: <input type="text"/>	Previous <span style="border: 1px solid black; padding: 2px;">1</span>	Next
Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	<input type="checkbox"/>	Exclude	<input checked="" type="checkbox"/>	Descendants	<input type="checkbox"/> Mapped
1342439	38454	<a href="#">trandolapril</a>	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
1334456	35296	<a href="#">Ramipril</a>	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
1331235	35208	<a href="#">quinapril</a>	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
1373225	54552	<a href="#">Perindopril</a>	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
1310756	30131	<a href="#">moexipril</a>	Drug	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

C.1 76

## Evidence for all conditions for ACEi and THZ combined

19.2.6

Query Language Chapter  
10 SQL R SQL R

OHDSI 

```
library(MethodEvaluation)
```

#      ACE*i* = 1                          -

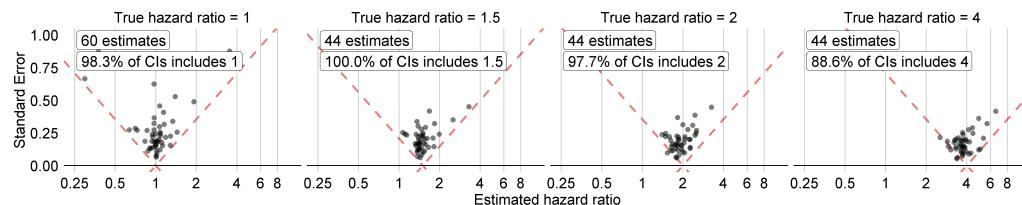
```
    outcomeId = ncs)

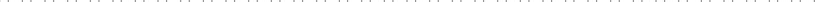
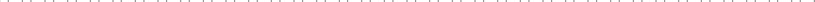
pcs <- synthesizePositiveControls(
  connectionDetails = connectionDetails,
  cdmDatabaseSchema = cdmDbSchema,
  exposureDatabaseSchema = cohortDbSchema,
  exposureTable = cohortTable,
  outcomeDatabaseSchema = cohortDbSchema,
  outcomeTable = cohortTable,
  outputDatabaseSchema = cohortDbSchema,
  outputTable = cohortTable,
  createOutputTable = FALSE,
  modelType = "survival",
  firstExposureOnly = TRUE,
  firstOutcomeOnly = TRUE,
```

```
removePeopleWithPriorOutcomes = TRUE,  
washoutPeriod = 365,  
riskWindowStart = 1,  
riskWindowEnd = 0,  
endAnchor = "cohort end",  
exposureOutcomePairs = eoPairs,  
effectSizes = c(1.5, 2, 4),  
cdmVersion = cdmVersion,  
workFolder = file.path(outputFolder, "pcSynthesis"))
```

Methods Library

19.2.7



**Figure 19.5:** =1>

[MethodEvaluation](https://ohd ## OHDSI)

OHDSI

188






---

<sup>1</sup><http://data.ohdsi.org/MethodEvalViewer/>



Figure 19.6: 95% GI = IBD =



# **Part V**

# **OHDSI**



# Chapter 20

□□□□□□□□: *Sara Dempster & Martijn Schuemie*

OHDSI OHDSI OHDSI R SQL CDM OMOP  
ETL 7 #

20.0.1

Level Estimation 13 / 19

20.0.2

Level Prediction PLE EHR

20.0.3

20.0.4

OHDSI 2, 8, 12, 13, 14

OHDSI ??

20.0.5

20.0.6 CDM

OHDSI OMOP CDM  
5 ETL  
CDM

CDM OMOP 6 OHDSI  
CDM OMOP  
##

20.0.7

(T2DM) Chapter 8 3

## T2DM

OHDSI

20.0.8

PI E

2009

OHDSI ATLAS  
@ref(Ohd ##)





# Chapter 21

— — OHDSI

□□□□: Kristin Kostka, Greg Klebanov & Sara Dempster

OHDSI Research Network

## 21.1 OHDSI

OHDSI CDM



 OHDSI

- OHDSI
  - OMOP
  - OMOP-CDM

21.2 OHDSI

## 21.2.1 OHDSI

### 21.2.2 OHDSI



OHDSI CDM

OHDSI OMOP CDM

OHDSI OHDSI OHDSI CDM OHDSI

OHDSI OHDSI OHDSI

- OHDSI GitHub
- CDM R SQL
- OHDSI OHDSI
- OHDSI GitHub
- R Shiny data.ohdsi.org

OHDSI

### 21.2.3 OHDSI

OHDSI

OHDSI

OHDSI

### 21.2.4 OHDSI

OHDSI OHDSI OHDSI CDM OHDSI OHDSI

OHDSI

- OHDSI
- CDM RStudio
- OHDSI R
- OHDSI R
- OHDSI R



Chapter ?(ExtractTransformLoad) ETL ?(DataQuality)

Level Effect Estimation PLE Patient Level Prediction PLP OHDSI

Level Effect Estimation PLE Patient Level Prediction PLP OHDSI

## OHDSI

OHDSI

- OHDSI
- OHDSI
- OHDSI

### 21.2.5

CDM  
CMS Synthetic Public  
Use Files  
Mitre  
SyntheticMass  
Synthea  
OHDSI  
20  
IRB  
OHDSI

### 21.2.6

OHDSI  
OHDSI  
GitHub  
OHDSI  
OHDSI  
SFTP  
Amazon S3  
/  
OHDSI  
OHDSI  
GitHub  
Shiny

### 21.2.7

OHDSI  
GitHub  
Shiny



OHDSI  
JANE  
Name Estimator  
Journal/Author  
<sup>1</sup>

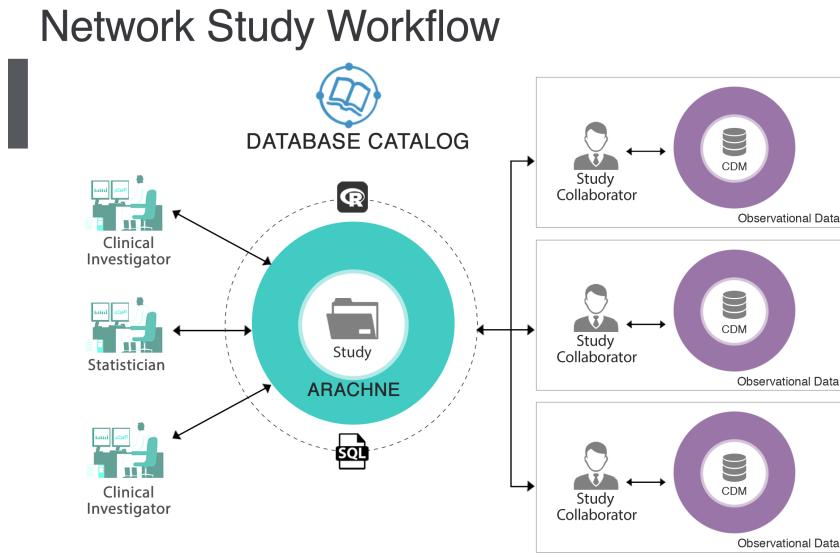


Figure 21.1: ARACHNE

ARACHNE ACHILLES ATLAS

## 21.3 OHDSI

OHDSI OHDSI

GitHub [data.ohdsi.org](https://data.ohdsi.org) R Shiny  
CDM OHDSI  
5 CDM

<sup>2</sup><http://www.icmje.org/recommendations/browse/roles-and-responsibilities/defining-the-role-of-authors-and-contributors.html> ## □□: □□□□□□□□□□□□□□□□□□

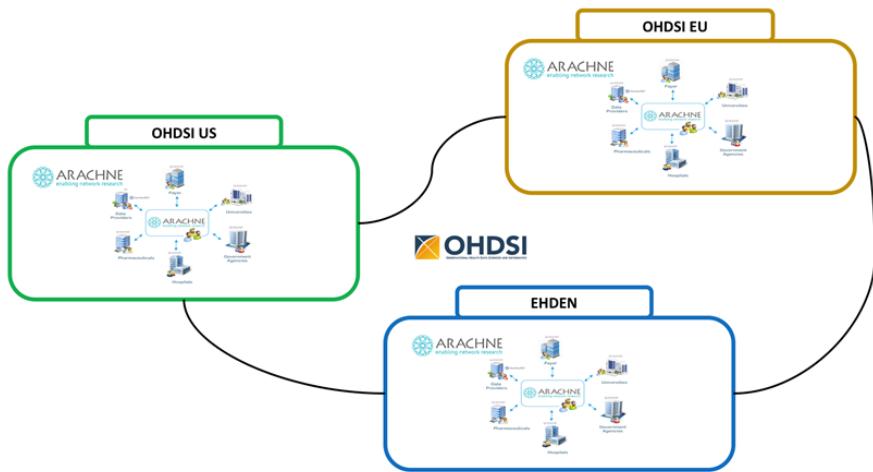


Figure 21.2: ARACHNE

## 21.4



- OHDSI CDM OHDSI
- OHDSI OHDSI Study Nurture Committee
- OHDSI GitHub R Shiny OHDSI



# Chapter A



# Chapter B

— — — — —

## B.1 ACE

- ACE B 1

365 0 1

1

1

- **30**
  - **0**

30 □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □

□ □ □ □ □ □ □ □ □

Table B.1: ACE

ID			
1308216	□□□□□□	□□□	□□
1310756	□□□□□□□□	□□□	□□
1331235	□□□□□	□□□	□□
1334456	□□□□□	□□□	□□
1335471	□□□□□□	□□□	□□
1340128	□□□□□□	□□□	□□
1341927	□□□□□□	□□□	□□
1342439	□□□□□□□□	□□□	□□
1363749	□□□□□□□□	□□□	□□
1373225	□□□□□□□□	□□□	□□

## B.2 ACE

-  ACE  B.2 

365 0 1

10

□□□□#1□□□□□1□□□□□□□□□□□□□□□□□

A horizontal row of twelve empty rectangular boxes, intended for children to write their names in, likely as part of a classroom activity.

- □□□□□□□□□□□□□365□□□□0□□□□□□□□□□□□□1□□□□□□□□□□□□  
B 3□□□□□

□□□□#2□□□□□□□□□□□□□□□□□□



A horizontal row of twelve empty square boxes, likely used for grading student responses.

- 0 7 B 4

per person

- **30**
  - **0**

Table B.2: ACE□□□

ID				
1308216	□□□□□□	□□	□□	□□□□□□
1310756	□□□□□□□□	□□□	□□	□□□
1331235	□□□□□	□□□	□□	□□□
1334456	□□□□□	□□□	□□	□□□
1335471	□□□□□□	□□□	□□	□□□
1340128	□□□□□□	□□□	□□	□□□
1341927	□□□□□□	□□□	□□	□□□
1342439	□□□□□□□□	□□□	□□	□□□
1363749	□□□□□□□□	□□□	□□	□□□
1373225	□□□□□□□□	□□□	□□	□□□

Table B.3: □ □ □ □ □ □

□□□□□ID □□□□□□ □□ □□ □□□□□  
316866 □□□□□□ □□□ □□ □□□

Table B.4: □ □ □ □

ID				
904542	□□□□□□□	□□□	□□	□□□□
907013	□□□□□□	□□□	□□	□□□□
932745	□□□□□□	□□□	□□	□□□□
942350	□□□□□□	□□□	□□	□□□□
956874	□□□□□□	□□□	□□	□□□□
970250	□□□□□□□□□	□□□	□□	□□□□
974166	□□□□□□□□□□□	□□□	□□	□□□□
978555	□□□□□□□	□□□	□□	□□□□

□□□□□ID	□□□□□□	□□	□□	□□□□□
991382	□□□□□	□□□	□□	□□□
1305447	□□□□□	□□□	□□	□□□
1307046	□□□□□□□	□□□	□□	□□□
1307863	□□□□□	□□□	□□	□□□
1308216	□□□□□□	□□□	□□	□□□
1308842	□□□□□□	□□□	□□	□□□
1309068	□□□□□□	□□□	□□	□□□
1309799	□□□□□□	□□□	□□	□□□
1310756	□□□□□□□	□□□	□□	□□□
1313200	□□□□□	□□□	□□	□□□
1314002	□□□□□□	□□□	□□	□□□
1314577	□□□□□□	□□□	□□	□□□
1317640	□□□□□□□	□□□	□□	□□□
1317967	□□□□□□	□□□	□□	□□□
1318137	□□□□□□	□□□	□□	□□□
1318853	□□□□□□	□□□	□□	□□□
1319880	□□□□□□	□□□	□□	□□□
1319998	□□□□□□□	□□□	□□	□□□
1322081	□□□□□□□	□□□	□□	□□□
1326012	□□□□□□	□□□	□□	□□□
1327978	□□□□□□□	□□□	□□	□□□
1328165	□□□□□□	□□□	□□	□□□
1331235	□□□□□	□□□	□□	□□□
1332418	□□□□□□	□□□	□□	□□□
1334456	□□□□□	□□□	□□	□□□
1335471	□□□□□□	□□□	□□	□□□
1338005	□□□□□□□	□□□	□□	□□□
1340128	□□□□□□	□□□	□□	□□□
1341238	□□□□□	□□□	□□	□□□
1341927	□□□□□□	□□□	□□	□□□
1342439	□□□□□□□□	□□□	□□	□□□
1344965	□□□□□□□	□□□	□□	□□□
1345858	□□□□□□	□□□	□□	□□□
1346686	□□□□□□□	□□□	□□	□□□
1346823	□□□□□□□	□□□	□□	□□□
1347384	□□□□□□□	□□□	□□	□□□
1350489	□□□□□	□□□	□□	□□□
1351557	□□□□□□□	□□□	□□	□□□
1353766	□□□□□□□□	□□□	□□	□□□
1353776	□□□□□□	□□□	□□	□□□
1363053	□□□□□□	□□□	□□	□□□
1363749	□□□□□□□	□□□	□□	□□□
1367500	□□□□□	□□□	□□	□□□

ID				
1373225	□□□□□□□	□□□	□□	□□□□□□
1373928	□□□□□□	□□□	□□	□□□
1386957	□□□□□□	□□□	□□	□□□
1395058	□□□□□□□	□□□	□□	□□□
1398937	□□□□□	□□□	□□	□□□
40226742	□□□□□□□	□□□	□□	□□□
40235485	□□□□□□□	□□□	□□	□□□

### B.3 AMI

- **B.5**

A horizontal row of 30 empty square boxes for writing responses.



5.5.5.5

Table B.5: □□□□□□□□□□

□□□□□ID	□□□□□□	□□	□□	□□□□□□
314666	□□□□□□	□□	□□	□□□
4329847	□□□□	□□□	□□	□□□

**Table B.6:** □□□□□□□□□□

B.4

A horizontal row of ten empty square boxes, each with a thin black border, intended for children to draw or write in.






A horizontal row of 20 empty square boxes, intended for children to write their names in, likely as part of a classroom activity or name recognition exercise.

□ □ □ **30** □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □ □

□ □ □ □ □ □ □ □

Table B.7: □□□□□

□□ID □□□ □□ □ □□□□□  
432791 □□□□□ □□□ □□ □□□

**Table B.8:** □□□□□□□□□□

<input type="checkbox"/>	ID	<input type="checkbox"/>		<input type="checkbox"/>		<input type="checkbox"/>
262		<input type="checkbox"/>				
9201		<input type="checkbox"/>				
9203		<input type="checkbox"/>				

□□ID □□□ □□ □ □□□□□

B.5

-  B.9 

10

□ □ □ □ □ 1 □ □ □ □ □ 1 □ □ □ □ □ □ □ □ □ □ □ □






□□□□3□□□□□□ACE□□□□□□□□□□□□



□□□□□□□□□□□□□□□□□□□□□□□ B.9 □□□□□□□□□□

- **30**
  - **0**

□□□□□□□□□0□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

□ □ □ □ □ □ □ □

Table B.9: □□□□□□□□□□□□□□□□□□□□□□□

□□ID	□□□	□□	□	□□□□□
907013	□□□□□	□□□	□□	□□□
974166	□□□□□□□□□□□	□□□	□□	□□□
978555	□□□□□□	□□□	□□	□□□
1395058	□□□□□□□	□□□	□□	□□□

Table B.10: □□□□□□

□□ID	□□□	□□	□	□□□□□
316866	□□□□□□□	□□□	□□	□□□

Table B.11: □□□□□□

□□ID	□□□	□□	□	□□□□□
904542	□□□□□□□□	□□□	□□	□□□
907013	□□□□□	□□□	□□	□□□
932745	□□□□□	□□□	□□	□□□
942350	□□□□□	□□□	□□	□□□
956874	□□□□□	□□□	□□	□□□
970250	□□□□□□□□□	□□□	□□	□□□
974166	□□□□□□□□□□□	□□□	□□	□□□
978555	□□□□□□	□□□	□□	□□□
991382	□□□□□□	□□□	□□	□□□
1305447	□□□□□	□□□	□□	□□□
1307046	□□□□□□□	□□□	□□	□□□
1307863	□□□□□	□□□	□□	□□□
1308216	□□□□□□	□□□	□□	□□□
1308842	□□□□□□	□□□	□□	□□□
1309068	□□□□□□	□□□	□□	□□□
1309799	□□□□□□	□□□	□□	□□□
1310756	□□□□□□□	□□□	□□	□□□
1313200	□□□□□	□□□	□□	□□□
1314002	□□□□□□	□□□	□□	□□□
1314577	□□□□□□	□□□	□□	□□□
1317640	□□□□□□□	□□□	□□	□□□
1317967	□□□□□□	□□□	□□	□□□
1318137	□□□□□□□	□□□	□□	□□□
1318853	□□□□□□	□□□	□□	□□□
1319880	□□□□□□	□□□	□□	□□□
1319998	□□□□□□□	□□□	□□	□□□

ID				
1322081	□□□□□□□	□□□	□□	□□□□□□
1326012	□□□□□□□	□□□	□□	□□□
1327978	□□□□□□□□	□□□	□□	□□□
1328165	□□□□□□□	□□□	□□	□□□
1331235	□□□□□□	□□□	□□	□□□
1332418	□□□□□□□	□□□	□□	□□□
1334456	□□□□□□	□□□	□□	□□□
1335471	□□□□□□□	□□□	□□	□□□
1338005	□□□□□□□□	□□□	□□	□□□
1340128	□□□□□□□	□□□	□□	□□□
1341238	□□□□□□	□□□	□□	□□□
1341927	□□□□□□□	□□□	□□	□□□
1342439	□□□□□□□□□	□□□	□□	□□□
1344965	□□□□□□□□	□□□	□□	□□□
1345858	□□□□□□□	□□□	□□	□□□
1346686	□□□□□□□□	□□□	□□	□□□
1346823	□□□□□□□□	□□□	□□	□□□
1347384	□□□□□□□□	□□□	□□	□□□
1350489	□□□□□□	□□□	□□	□□□
1351557	□□□□□□□□	□□□	□□	□□□
1353766	□□□□□□□□□	□□□	□□	□□□
1353776	□□□□□□□	□□□	□□	□□□
1363053	□□□□□□□	□□□	□□	□□□
1363749	□□□□□□□□	□□□	□□	□□□
1367500	□□□□□□	□□□	□□	□□□
1373225	□□□□□□□□	□□□	□□	□□□
1373928	□□□□□□□	□□□	□□	□□□
1386957	□□□□□□□	□□□	□□	□□□
1395058	□□□□□□□□	□□□	□□	□□□
1398937	□□□□□□	□□□	□□	□□□
40226742	□□□□□□□□	□□□	□□	□□□
40235485	□□□□□□□□	□□□	□□	□□□

B.6

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-  B.12 

10 of 10

- 1 B.13 0
  - 365 0 B.1

5 / 5

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Table B.12: □□□□□□□□

ID				
907013	███████	NO	YES	NO
974166	██████████████████	NO	YES	NO
978555	██████████	NO	YES	NO
1307863	███████	NO	YES	NO
1308216	██████████	NO	YES	NO
1308842	██████████	NO	YES	NO
1310756	██████████	NO	YES	NO
1317640	██████████	NO	YES	NO
1318137	██████████	NO	YES	NO
1318853	██████████	NO	YES	NO
1319880	██████████	NO	YES	NO
1326012	██████████	NO	YES	NO
1328165	██████████	NO	YES	NO
1331235	███████	NO	YES	NO
1332418	██████████	NO	YES	NO
1334456	███████	NO	YES	NO
1335471	██████████	NO	YES	NO
1340128	██████████	NO	YES	NO
1341927	██████████	NO	YES	NO
1342439	██████████████	NO	YES	NO
1346686	██████████	NO	YES	NO
1347384	██████████	NO	YES	NO
1351557	██████████	NO	YES	NO

□□□□□ID	□□□□□□	□□	□□	□□□□□
1353776	□□□□□□	NO	YES	NO
1363749	□□□□□□	NO	YES	NO
1367500	□□□□□	NO	YES	NO
1373225	□□□□□□□	NO	YES	NO
1395058	□□□□□□□	NO	YES	NO
40226742	□□□□□□□	NO	YES	NO
40235485	□□□□□□□	NO	YES	NO

Table B.13: □□□□

□□□□□ID	□□□□□□	□□	□□	□□□□□
904542	□□□□□□□□	NO	YES	NO
907013	□□□□□	NO	YES	NO
932745	□□□□□	NO	YES	NO
942350	□□□□□	NO	YES	NO
956874	□□□□□	NO	YES	NO
970250	□□□□□□□□□	NO	YES	NO
974166	□□□□□□□□□□	NO	YES	NO
978555	□□□□□□	NO	YES	NO
991382	□□□□□	NO	YES	NO
1305447	□□□□□	NO	YES	NO
1307046	□□□□□□□	NO	YES	NO
1307863	□□□□□	NO	YES	NO
1308216	□□□□□□	NO	YES	NO
1308842	□□□□□□	NO	YES	NO
1309068	□□□□□□	NO	YES	NO
1309799	□□□□□□	NO	YES	NO
1310756	□□□□□□□	NO	YES	NO
1313200	□□□□□	NO	YES	NO
1314002	□□□□□□	NO	YES	NO
1314577	□□□□□□	NO	YES	NO
1317640	□□□□□□□	NO	YES	NO
1317967	□□□□□□	NO	YES	NO
1318137	□□□□□□	NO	YES	NO
1318853	□□□□□□	NO	YES	NO
1319880	□□□□□□	NO	YES	NO
1319998	□□□□□□□	NO	YES	NO
1322081	□□□□□□□	NO	YES	NO
1326012	□□□□□□	NO	YES	NO
1327978	□□□□□□□	NO	YES	NO
1328165	□□□□□□	NO	YES	NO
1331235	□□□□□	NO	YES	NO

ID				
1332418	□□□□□□□	NO	YES	NO
1334456	□□□□□□	NO	YES	NO
1335471	□□□□□□□	NO	YES	NO
1338005	□□□□□□□□	NO	YES	NO
1340128	□□□□□□□	NO	YES	NO
1341238	□□□□□□	NO	YES	NO
1341927	□□□□□□	NO	YES	NO
1342439	□□□□□□□□□	NO	YES	NO
1344965	□□□□□□□□	NO	YES	NO
1345858	□□□□□□□	NO	YES	NO
1346686	□□□□□□□□	NO	YES	NO
1346823	□□□□□□□□	NO	YES	NO
1347384	□□□□□□□□	NO	YES	NO
1350489	□□□□□□	NO	YES	NO
1351557	□□□□□□□□	NO	YES	NO
1353766	□□□□□□□□□	NO	YES	NO
1353776	□□□□□□□	NO	YES	NO
1363053	□□□□□□□	NO	YES	NO
1363749	□□□□□□□	NO	YES	NO
1367500	□□□□□□	NO	YES	NO
1373225	□□□□□□□□	NO	YES	NO
1373928	□□□□□□□	NO	YES	NO
1386957	□□□□□□□	NO	YES	NO
1395058	□□□□□□□□	NO	YES	NO
1398937	□□□□□□	NO	YES	NO
40226742	□□□□□□□□	NO	YES	NO
40235485	□□□□□□□□	NO	YES	NO

Table B-14: □□□□□□

B.7 3

## B.8 ACE

- ACE□□□□□ B.15□□□□□

ACE□□□□ B.15□□□□□□□□□□□□□

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Table B.15: ACE□□□

ID				
1308216	□□□□□□	NO	YES	NO
1310756	□□□□□□□	NO	YES	NO
1331235	□□□□□	NO	YES	NO
1334456	□□□□□	NO	YES	NO
1335471	□□□□□□	NO	YES	NO
1340128	□□□□□□	NO	YES	NO
1341927	□□□□□□	NO	YES	NO
1342439	□□□□□□□□	NO	YES	NO
1363749	□□□□□□	NO	YES	NO
1373225	□□□□□□□	NO	YES	NO

B.9 ARB

□□□□□□□

ID				
1308842	██████████	NO	YES	NO
1317640	███████████	NO	YES	NO
1346686	███████████	NO	YES	NO
1347384	███████████	NO	YES	NO
1351557	███████████	NO	YES	NO
1367500	███████	NO	YES	NO
40226742	███████████	NO	YES	NO
40235485	███████████	NO	YES	NO

B.10

**B.8** **B.17** **ACE** **B.15**

**Table B.17:** □□□□□□□□□□□□□□□□□□□□□□□□□□□□□

ID				
907013	000000	NO	YES	NO
974166	0000000000	NO	YES	NO
978555	000000	NO	YES	NO
1395058	000000	NO	YES	NO

B.11 dCCB

Table B.18:  dCCB

ID				
1318137	□□□□□□	NO	YES	NO
1318853	□□□□□□	NO	YES	NO
1319880	□□□□□□	NO	YES	NO
1326012	□□□□□□	NO	YES	NO

ID				
1332418		NO	YES	NO
1353776		NO	YES	NO

## B.12                      ndCCB

Table B.19: □□□□□□□□□□□□□□□□□ndCCB□

ID				
1307863		NO	YES	NO
1328165		NO	YES	NO

B.13

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**Table B.20:** □□□□□□□□

ID					
1307046	██████████	NO	YES	NO	
1313200	███████	NO	YES	NO	
1314002	██████████	NO	YES	NO	
1314577	██████████	NO	YES	NO	
1319998	██████████	NO	YES	NO	
1322081	██████████	NO	YES	NO	
1327978	██████████	NO	YES	NO	
1338005	██████████	NO	YES	NO	
1345858	███████	NO	YES	NO	
1346823	██████████	NO	YES	NO	
1353766	████████████	NO	YES	NO	
1386957	███████	NO	YES	NO	

**B.14**

*ACE*□□□□□ *B.8* □□□□□□□□□□□□□□□ (□ B.21) □ *ACE*□□□ (□ B.15) □□□□□□□□□□□  
 □□□□□□□□□

Table B.21: □□□□□□□

□□ID	□□□	□□	□□	□□□
932745	□□□□□	NO	YES	NO
942350	□□□□□	NO	YES	NO
956874	□□□□□	NO	YES	NO

**B.15**

*ACE*□□□□□ *B.8* □□□□□□□□□□□□□□□ (□ B.22) □ *ACE*□□□ (□ B.15)  
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Table B.22: □□□□□□□□□

□□ID	□□□	□□	□□	□□□
904542	□□□□□□□	NO	YES	NO
991382	□□□□□□	NO	YES	NO

**B.16 1**

*ACE*□□□□□ *B.8* □□□□□□□□□□□□□□□ *I*□□□□□ (□ B.23) □ *ACE*□□□ (□ B.15) □□□□□□□□□  
 □□□□□□□□□

Table B.23: □□□□1□□□□□

□□ID	□□□	□□	□□	□□□
1341238	□□□□□	NO	YES	NO
1350489	□□□□□	NO	YES	NO
1363053	□□□□□□	NO	YES	NO



# Chapter C

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100 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118 119 120 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147 148 149 150 151 152 153 154 155 156 157 158 159 160 161 162 163 164 165 166 167 168 169 170 171 172 173 174 175 176 177 178 179 180 181 182 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197 198 199 200

## C.1 ACEi THZ

ID	
434165	███████████████████
436409	███████
199192	██/████████████████
4088290	███████
4092879	███████
44783954	███████
75911	███████████
137951	███████████
77965	██████████
376707	██████████
4103640	███████
73241	███████████████████
133655	███████
73560	██████████
434327	███████
4213540	███████████████████
140842	███████████████
81378	███████████████
432303	███████████
4201390	███████████
46269889	███████████████████

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□□□□□ID	□□□□□□□□□
134438	□□□□□□
78619	□□□□□
201606	□□□□□
76786	□□□□
4115402	□□□□
45757370	□□□□□□□□□
433111	□□□□□
433527	□□□□□
4170770	□□□□
4092896	□□□□□□□□□
259995	□□□□□□□□
40481632	□□□□□□□□□
4166231	□□□□□
433577	□□□□□□
4231770	□□□□□□□
440329	□□□□□□□□□□□□
4012570	□□□□□□□□□
4012934	□□□□□□□□□
441788	□□□□□□□□□□□□□□
4201717	□□□□□□□
374375	□□□□
4344500	□□□□□□□□□□□□□□□□□
139099	□□□
444132	□□□□
196168	□□□□□□□
432593	□□□□□□□□
434203	□□□□□□□
438329	□□□□□□□□□□
195873	□□□□
4083487	□□□
4103703	□□□□□□□□□
4209423	□□□□□□□□
377572	□□□□□□□□□□□
40480893	□□□□□□□□□□□□□□□
136368	□□□□□□□□□□□□
140648	□□□□□□□□□□□□
438130	□□□□□□□
4091513	□□
4202045	□□□□□□□□□□□□
373478	□□
46286594	□□□□□□□□□□□□□
439790	□□□□
81634	□□□□

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□□□□ID	□□□□□□□
380706	□□□□□□
141932	□□□□□□
36713918	□□□□□□□□□
443172	□□□□□□□□□□□
81151	□□□□□
72748	□□□□□□□□□□□□□□□□
378427	□□□□
437264	□□□□□□
194083	□□□□□□□□□
140641	□□□□□
440193	□□□□
4115367	□□□□



# **Chapter D**

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16. □□□□

# Chapter E

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A horizontal row of 20 empty rectangular boxes, likely for students to write their answers in a worksheet.

E.1

5.1

E.1

Table E.1: PERSON□□□□□

□□□□	□	□□
PERSON_ID	2	□□□□□□
GENDER_CONCEPT_ID	8507	□□□□□ ID □ 8507□
YEAR_OF_BIRTH	1974	
MONTH_OF_BIRTH	8	
DAY_OF_BIRTH	4	
BIRTH_DATETIME	1974-08-04 00:00:00	□□□□□□□□□□□□□00:00:00□□□□□
DEATH_DATETIME	NULL	
RACE_CONCEPT_ID	8516	□□□□□□□□□□□□□ ID □ 8516□
ETHNICITY_CONCEPT_ID	38003564	38003564
LOCATION_ID		□□□□□□□□□□□□□
PROVIDER_ID		□□□□□□□
CARE_SITE		□□□□□□□□□□
PERSON_SOURCE_VALUE	NULL	□□□□□□□□□□
GENDER_SOURCE_VALUE	Man	□□□□□□□□□□□□□

□	□	□
GENDER_SOURCE_	0	
CONCEPT_ID		
RACE_SOURCE_	African	□□□□□□□□□□□□
VALUE	American	
RACE_SOURCE_	0	
CONCEPT_ID		
ETHNICITY_SOURCE_	NULL	
VALUE		
ETHNICITY_SOURCE_	0	
CONCEPT_ID		

□ □ □ □ 5.2

Table E.2: OBSERVATION PERIOD

□ □ □ □ 5.3

Table E.3: DRUG EXPOSURE

□□□□	□	□□
DRUG_EXPOSURE_	2019-05-01	□□□□□□□□□□□□□□□
START_DATETIME	00:00:00	
DRUG_EXPOSURE_	2019-05-31	□□□ + □□□□□□□□□
END_DATE		
DRUG_EXPOSURE_	2019-05-31	□□□□□□□□□□□□□□
END_DATETIME	00:00:00	
VERBATIM_END_DATE	NULL	□□□□□□□□□
DRUG_TYPE_	38000177	38000177 □□□□□□□□□□□□□□
CONCEPT_ID		
STOP_REASON	NULL	
REFILLS	NULL	
QUANTITY	NULL	□□□□□□□□□
DAYS_SUPPLY	30	□□□□□□□□□□□□□□
SIG	NULL	□□□□□□□□□
ROUTE_CONCEPT_ID	4132161	4132161 □□□□□□□□
LOT_NUMBER	NULL	□□□□□□□□□
PROVIDER_ID	NULL	□□□□□□□□□
VISIT_OCCURRENCE_	NULL	□□□□□□□□□□□□□□□
ID		
VISIT_DETAIL_ID	NULL	
DRUG_SOURCE_	76168009520	□□□□□NDC□□□□
VALUE		
DRUG_SOURCE_	583945	583945 □□□□□□□□
CONCEPT_ID		(NDC□□□□76168009520□)□
ROUTE_SOURCE_	NULL	
VALUE		

## □□□□ 5.4

□□□□□□□□□□□□□□□ CONDITION\_OCCURRENCE□□□□□□□□□□□□□□

```
library(DatabaseConnector)
connection <- connect(connectionDetails)
sql <- "SELECT *
FROM @cdm.condition_occurrence
WHERE condition_concept_id = 192671;"

result <- renderTranslateQuerySql(connection, sql, cdm = "main")
head(result)
```

```
##   CONDITION_OCCURRENCE_ID PERSON_ID CONDITION_CONCEPT_ID ...
## 1                      4657        273             192671 ...
## 2                      1021        61              192671 ...
```

```
## 3           5978     351      192671 ...
## 4           9798     579      192671 ...
## 5          9301     549      192671 ...
## 6          1997     116      192671 ...
```

5.5

□□□□□□□□□□ CONDITION OCCURRENCE □□□□ CONDITION SOURCE VALUE

```
sql <- "SELECT *
FROM @cdm.condition_occurrence
WHERE condition_source_value = 'K92.2';"

result <- renderTranslateQuerySql(connection, sql, cdm = "main")
head(result)
```

```
##   CONDITION_OCCURRENCE_ID PERSON_ID CONDITION_CONCEPT_ID ...
## 1                 4657      273          192671 ...
## 2                 1021       61          192671 ...
## 3                 5978      351          192671 ...
## 4                 9798      579          192671 ...
## 5                 9301      549          192671 ...
## 6                1997      116          192671 ...
```

5.6

## **OBSERVATION PERIOD**

```
library(DatabaseConnector)
connection <- connect(connectionDetails)
sql <- "SELECT *
FROM @cdm.observation_period
WHERE person_id = 61;"

renderTranslateQuerySql(connection, sql, cdm = "main")
```

E.2

□ □ □ □ 6.1

□□ ID 192671 (“□□□□”)

□ □ □ □ 6.2

ICD-10CM□□□□

- K29.91 “□□□□□□□□□□□□□□□□□□”
  - K92.2 “□□□□□□□□□□□□”

ICD-9CM□□□□

- 578 “□□□□”
  - 578.9 “□□□□□□□□□□□□□□”

□ □ □ □ 6.3

MedDRA□□□□□

- “□□□□” (□□ID 35707864)
  - “□□□” (□□ID 35707858)

E.3

□ □ □ □ 7.1

- A)  CDM
  - B)  ETL
  - C)  ETL
  - D)

□ □ □ □ 7.2

PERSON_ID	A123B456	
GENDER_CONCEPT_ID	8532	
YEAR_OF_BIRTH	NULL	
MONTH_OF_BIRTH	NULL	
DAY_OF_BIRTH	NULL	
RACE_CONCEPT_ID	0	WHITE
ETHNICITY_CONCEPT_ID	8527	8527
ID		0
PERSON_SOURCE_VALUE	A123B456	
GENDER_SOURCE_VALUE	F	
RACE_SOURCE_VALUE	WHITE	
ETHNICITY_SOURCE_VALUE	NONE	
PROVIDED		

□ □ □ □ 7.3

VISIT_OCCURRENCE_ID	1
PERSON_ID	11
VISIT_START_DATE	2004-09-26
VISIT_END_DATE	2004-09-30
VISIT_CONCEPT_ID	9201
VISIT_SOURCE_VALUE	inpatient

E.4

□ □ □ □ 8.1

1. □□□□
  2. □□□□□□□□
  3. □□□□□□□□

□ □ □ □ 8.2

## SQL|R {#SqlAndRanswers}

□ □ 10.1

```
library(DatabaseConnector)
connection <- connect(connectionDetails)
sql <- "SELECT COUNT(*) AS person_count
FROM @cdm.person;"

renderTranslateQuerySql(connection, sql, cdm = "main")
```

```
##    PERSON_COUNT  
## 1          2694
```

10.2

1 DRUG EXPOSURE

```
library(DatabaseConnector)
connection <- connect(connectionDetails)
sql <- "SELECT COUNT(DISTINCT(person_id)) AS person_count
FROM @cdm.drug_exposure
INNER JOIN @cdm.concept_ancestor
    ON drug_concept_id = descendant_concept_id
INNER JOIN @cdm.concept ingredient
    ON ancestor_concept_id = ingredient.concept_id
WHERE LOWER(ingredient.concept_name) = 'celecoxib'
    AND ingredient.concept_class_id = 'Ingredient'
    AND ingredient.standard_concept = 'S';"

renderTranslateQuerySql(connection, sql, cdm = "main")
```

```
##      PERSON_COUNT  
## 1          1844
```

```
library(DatabaseConnector)
connection <- connect(connectionDetails)

sql <- "SELECT COUNT(DISTINCT(person_id)) AS person_count
FROM @cdm.drug_era
INNER JOIN @cdm.concept ingredient
  ON drug_concept_id = ingredient.concept_id
WHERE LOWER(ingredient.concept_name) = 'celecoxib'
  AND ingredient.concept_class_id = 'Ingredient'
  AND ingredient.standard_concept = 'S';"

renderTranslateQuerySql(connection, sql, cdm = "main")
```

```
##      PERSON_COUNT  
## 1          1844
```

□ □ 10.3

#### CONDITION OCCURRENCE

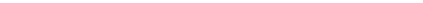
```
library(DatabaseConnector)
connection <- connect(connectionDetails)
sql <- "SELECT COUNT(*) AS diagnose_count
FROM @cdm.drug_era
INNER JOIN @cdm.concept ingredient
    ON drug concept id = ingredient.concept id"
```

```
    INNER JOIN @cdm.condition_occurrence
        ON condition_start_date >= drug_era_start_date
           AND condition_start_date <= drug_era_end_date
    INNER JOIN @cdm.concept_ancestor
        ON condition_concept_id = descendant_concept_id
    WHERE LOWER(ingredient.concept_name) = 'celecoxib'
        AND ingredient.concept_class_id = 'Ingredient'
        AND ingredient.standard_concept = 'S'
        AND ancestor_concept_id = 192671;"
```

```
##    DIAGNOSE_COUNT  
## 1        41
```

E.5

□ □ 11.1

- 
  - 16 
  - 

E.1

Cohort Entry Events

Events having any of the following criteria:

+ Add Initial Event ▾
+ Add attribute... ▾
Delete Criteria

a drug era of **diclofenac** ▾

**X** for the first time in the person's history

**X** with age in years at era start **Greater or Equal To** ▾ **16**

with continuous observation of at least **365** ▾ days before and **0** ▾ days after event index date

Limit initial events to: **earliest event** ▾ per person.

**Restrict initial events**

E.2 Diclofenac

NSAIDs□□□□□□□□□E.4□□□□□□□NSAIDs□□□□□□□□□NSAID□□□□□□

Concept Set Expression   Included Concepts (11473)   Included Source Codes   Export   Import

Name:  
diclofenac

Show 25 entries   Search:

Showing 1 to 1 of 1 entries

	Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	Exclude	Descendants	Mapped
	1124300	3355	Diclofenac	Drug	Standard	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>

Classification   Non-Standard   Standard

Figure E.2: □□□□□□□□□□□□□□□□□□

Inclusion Criteria

New inclusion criteria

Without prior exposure to any NSAID   Copy   Delete

1. Without prior exposure to any NSAID  
*Excluding subjects with prior exposure to any NSAID*

Excluding subjects with prior exposure to any NSAID  
having all of the following criteria:

+ Add criteria to group...   Delete Criteria

with exactly 0 using all occurrences of:  
a drug exposure of **NSAIDs**   + Add attribute...

where **event starts** between All days Before and 1 days Before  
**index start date** [add additional constraint](#)

restrict to the same visit occurrence  
 allow events from outside observation period

Limit qualifying events to: earliest event per person.

Figure E.3: NSAID□□□□□□□□□□□□

Concept Set Expression   Included Concepts (23112)   Included Source Codes   Export   Import

Name:  
NSAIDs

Show 25 entries   Search:

Showing 1 to 1 of 1 entries

	Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	Exclude	Descendants	Mapped
	21603933	M01A	ANTIIINFLAMMATORY AND ANTRHEUMATIC PRODUCTS, NON-STEROIDS	Drug	Classification	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>

Classification   Non-Standard   Standard

Figure E.4: NSAIDs□□□□□□□□

E.5

**Inclusion Criteria**

New inclusion criteria

Without prior exposure to any NSAID  
Excluding subjects with prior exposure to any NSAID

Without prior diagnose of cancer  
Excluding subjects with prior cancer diagnosis

having all of the following criteria:

+ Add criteria to group... □

**2. Without prior diagnose of cancer**  
Excluding subjects with prior cancer diagnosis

with exactly 0 occurrences of:  
a condition occurrence of Broad malignancies + Add attribute... □

where event starts between All days Before and 0 days Before  
index start date add additional constraint  
 restrict to the same visit occurrence  
 allow events from outside observation period

Delete Criteria

Limit qualifying events to: earliest event per person.

**Figure E.5:** □□□□□□□□□□□□□□□□□□

E.6

Concept Set Expression	Included Concepts 4401	Included Source Codes	Export	Import				
Name:								
Broad malignancies								
Show	25 ▾	entries	Search:					
Showing 1 to 1 of 1 entries								
Previous	1			Next				
Cart	Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	Exclude	Descendants	Mapped
Cart	443392	363346000	Malignant neoplastic disease	Condition	Standard	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>

Figure E.6: □□□□□□□□□□□□□□□□

11.2

SOL 2

```
library(DatabaseConnector)
connection <- connect(connectionDetails)
sql <- "SELECT person_id AS subject_id,
  condition_start_date AS cohort_start_date
INTO #diagnoses
FROM @cdm.condition_occurrence
WHERE condition_concept_id IN (
  SELECT descendant_concept_id
  FROM @cdm.concept_ancestor
```

**Cohort Exit**

**Event Persistence:**  
Event will persist until:  ▾

**Continuous Exposure Persistence:**  
Specify a concept set that contains one or more drugs. A drug era will be derived from all drug exposure events for any of the drugs within the concept set, using the specified persistence window as a maximum allowable gap in days between successive exposure events and adding a specified surveillance window to the final exposure event. If no exposure event end date is provided, then an exposure event end date is inferred to be event start date + days supply in cases when days supply is available or event start date + 1 day otherwise. This event persistence assures that the cohort end date will be no greater than the drug era end date.

Concept set containing the drug(s) of interest:  ▾

- Persistence window: allow for a maximum of  days between exposure records when inferring the era of persistence exposure
- Surveillance window: add  days to the end of the era of persistence exposure as an additional period of surveillance prior to cohort exit.

**Censoring Events:**

Exit Cohort based on the following criteria:

No censoring events selected.

**Figure E.7:** □□□□□□□□□□

```
        WHERE ancestor_concept_id = 4329847 --  
    )  
    AND condition_concept_id NOT IN (  
        SELECT descendant_concept_id  
        FROM @cdm.concept_ancestor  
        WHERE ancestor_concept_id = 314666 --  
    );"  
  
renderTranslateExecuteSql(connection, sql, cdm = "main")
```

ER COHORT DEFINITION ID

```
sql <- "INSERT INTO @cdm.cohort (
    subject_id,
    cohort_start_date,
    cohort_definition_id
)
SELECT subject_id,
    cohort_start_date,
    CAST (1 AS INT) AS cohort_definition_id
FROM #diagnoses
INNER JOIN @cdm.visit_occurrence
    ON subject_id = person_id
        AND cohort_start_date >= visit_start_date
        AND cohort_start_date <= visit_end_date
WHERE visit_concept_id IN (9201, 9203, 262); --      ER;"
```

```
renderTranslateExecuteSql(connection, sql, cdm = "main")
```

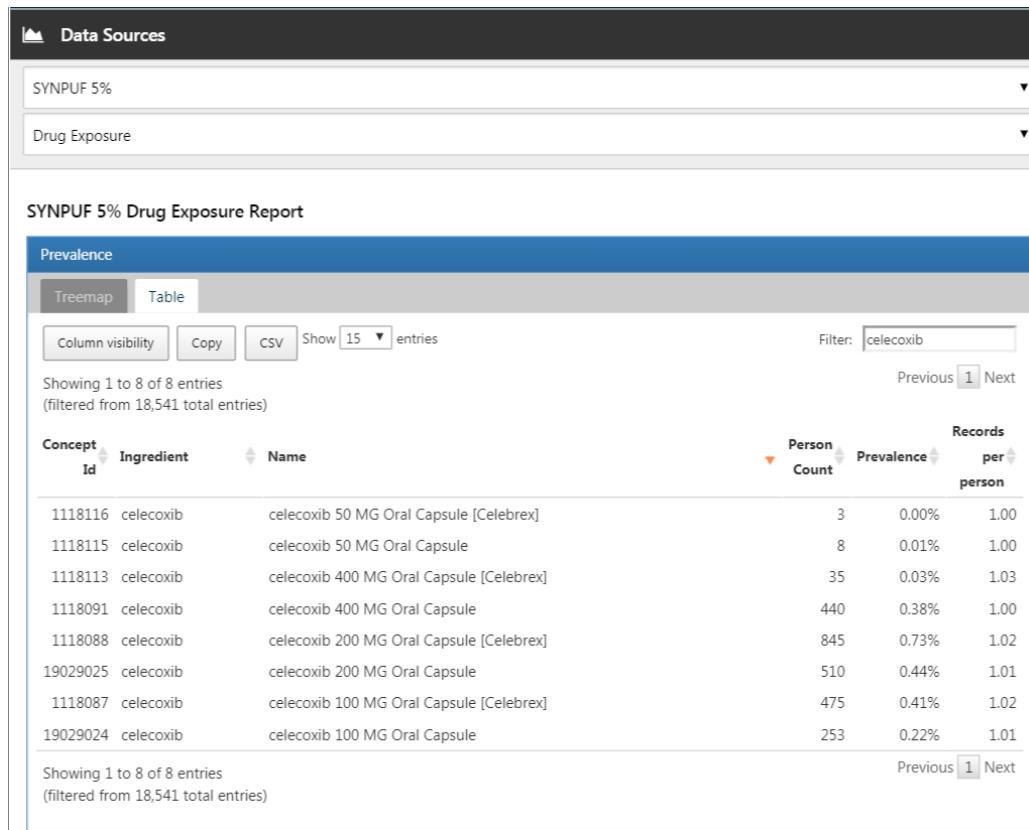
VISIT\_OCCURRENCE\_ID:

```
sql <- "TRUNCATE TABLE #diagnoses;  
DROP TABLE #diagnoses;"  
  
renderTranslateExecuteSql(connection, sql)
```

E.6

□ □ 12.1

ATLAS Data Sources  
E.8



**Figure E.8:** □□□□□□□□□□□□

□ □ 12.2

Cohort Definitions

Search

E.9

← Celecoxib new users ➔ Celecoxib

**Q Search**

**Search Import**

celexcoxib

[Advanced Options](#)

Column visibility CSV Show 15 entries Filter:

Showing 1 to 1 of 1 entries

Id	Code	Name	Class	RC	DRC	Domain	Vocabulary
1118084	140587	celexcoxib	Ingredient	2,587	5,184	Drug	RxNorm

Showing 1 to 1 of 1 entries

**Vocabulary**

- RxNorm Extension (1376)
- NDC (1337)
- SPL (449)
- DPD (167)
- [SNOMED \(75\)](#)

**Class**

- Ingredient (7)**
- Clinical Drug Form (5)
- Clinical Drug Comp (5)
- Lab Test (5)
- [More \(6\)](#)

**Domain**

- Drug (3570)
- Measurement (18)
- Observation (1)
- Meas Value (1)

**Standard Concept**

- Non-Standard (1831)
- Standard (1292)**
- Classification (467)

Figure E.9: □□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

E.9 +



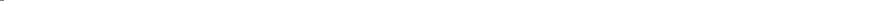
 Characterizations

E.11  

□□□□□□□□□□□□□□□□ 1 □□□□□□□□□□□□□□□□□□□□□□

E.12 □ □ □ □

□ □ 12.3

 Cohort Definitions  G  
E.13 

## New Cohort Definition

Celecoxib new users Save  Cancel 

- [Definition !\[\]\(fb19c0689fa6de7a5107037a8c7e9217\_img.jpg\)](#)
- [Concept Sets](#)
- [Generation](#)
- [Reporting](#)
- [Export](#)

enter a cohort definition description here

### Cohort Entry Events

Events having any of the following criteria:

a drug era of Celecoxib   Add Initial Event 

 Delete Criteria

 for the first time in the person's history

with continuous observation of at least 365  days before and 0  days after event index date  
 Limit initial events to: earliest event  per person.

Restrict initial events

**Figure E.10:** □□□□□□□□□□□□□□□□□□□□□□□□□□□□□

E.15 + 

E.16  Incidence Rates

E.17 □□□

E.7

□ □ 13.1

```
library(CohortMethod)
nsaids <- c(1118084, 1124300) #
covSettings <- createDefaultCovariateSettings(
  excludedCovariateConceptIds = nsaids,
  addDescendantsToExclude = TRUE)

#
cmData <- getDbCohortMethodData(
  connectionDetails = connectionDetails,
```

**New Characterization**

Celecoxib new users characterization

Design Executions Utilities

**Cohort characterization** is defined as the process of generating cohort level descriptive summary statistics from person level covariate data. Summary statistics of these person level covariates may be count, mean, sd, var, min, max, median, range, and quantiles. In addition, covariates during a period may be stratified into temporal units of time for time-series analysis such as fixed intervals of time relative to cohort\_start\_date (e.g. every 7 days, every 30 days etc.), or in absolute calendar intervals such as calendar-week, calendar-month, calendar-quarter, calendar-year.

**Cohort definitions**

Import

Show 10 entries Search:

ID	Name	Actions
1771701	Celecoxib new users	Edit cohort Remove

Showing 1 to 1 of 1 entries Previous 1 Next

**Feature analyses**

Import

Show 10 entries Search:

ID	Name	Description	Actions
15	Drug Group Era Any Time Prior	One covariate per drug rolled up to ATC groups in the drug_era table overlapping with any time prior to index.	Remove
27	Condition Group Era Any Time Prior	One covariate per condition era rolled up to groups in the condition_era table overlapping with any time prior to index.	Remove

Showing 1 to 2 of 2 entries Previous 1 Next

Figure E.11: □□□□□□

**Characterization #69**

Celecoxib new users characterization

Design Executions Utilities

Executions > Reports for SYNPUF 5%

Date: 08/23/2019 12:53 PM Design: -1840810470 Results: 2 reports

**Filter panel**

Cohorts	Analyses	Domains
Celecoxib new users	Condition Group Era Any Time P	Condition, Drug

**CONDITION / Condition Group Era Any Time Prior**

Export Show 10 entries Search:

Covariate	Explore	Concept ID	Count	Pct
Pain	Explore	4329041	1,140	78.62%
Pain finding at anatomical site	Explore	4132926	1,135	78.28%
Inflammation of specific body systems	Explore	4178818	1,135	78.28%
Arthropathy	Explore	73553	1,122	77.38%

**Figure E.12:** □□□□□□□

◀ GI bleed ▶ GI bleed

**Search**

Search Import

Gastrointestinal hemorrhage

Advanced Options

Column visibility Copy CSV Show 15 entries Filter: Previous 1 2 Next

Showing 1 to 15 of 25 entries

Vocabulary	Id	Code	Name	Class	RC	DRC	Domain	Vocabulary
SNOMED (17)	192671	74474003	Gastrointestinal hemorrhage	Clinical Finding	919	37,144	Condition	SNOMED
ICD10CM (2)	4338544	87763006	Lower gastrointestinal hemorrhage	Clinical Finding	0	15,617	Condition	SNOMED
ICD9CM (2)								
DRG (2)								
NIDERT (1)								
Class	4100660	27719009	Acute gastrointestinal hemorrhage	Clinical Finding	0	9,852	Condition	SNOMED
Clinical Finding (17)								

**Figure E.13:** □□□□□□□□□□□□□□□□□□□

Concept Set Expression   Included Concepts (191)   Included Source Codes   Export   Import

Name: GI bleed

Show 25 entries   Search:  Previous | 1 | Next

Showing 1 to 1 of 1 entries

Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	Exclude	Descendants	Mapped
192671	74474003	Gastrointestinal hemorrhage	Condition	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>

Classification   Non-Standard   Standard

Figure E.14: □□□□□□□□□□□□□□□□□□□□□□□□

New Cohort Definition

GI bleed

Definition   Concept Sets   Generation   Reporting   Export

enter a cohort definition description here

Cohort Entry Events

Events having any of the following criteria:

a condition occurrence of **GI bleed**

with continuous observation of at least  days before and  days after event index date

Limit initial events to:  earliest event per person.

Figure E.15: □□□□□□□□□□□□□□□□□□

New Incidence Rate Analysis

Incidence of GI bleed after celecoxib initiation

Definition   Concept Sets   Generation   Utilities

Study Cohorts

Target Cohorts: #1771701:Celecoxib new users

Outcome Cohorts: #1771702:GI bleed

Time At Risk

Time at risk defines the time window relative to the cohort start or end date with an offset to consider the person 'at risk' of the outcome.

- Time at risk starts with  start date  plus  days.
- Time at risk ends with  start date  plus  1095 days.

No study window defined.

**Stratify Criteria:** You can provide optional stratification criteria to the analysis that will divide the population into unique groups based on their satisfied criteria.

Please select a qualifying inclusion criteria to edit

Figure E.16: □□□□□□

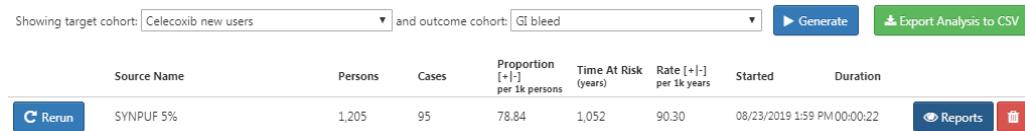


Figure E.17: □□□□□

```
cdmDatabaseSchema = "main",
targetId = 1,
comparatorId = 2,
outcomeIds = 3,
exposureDatabaseSchema = "main",
exposureTable = "cohort",
outcomeDatabaseSchema = "main",
outcomeTable = "cohort",
covariateSettings = covSettings)
summary(cmData)
```

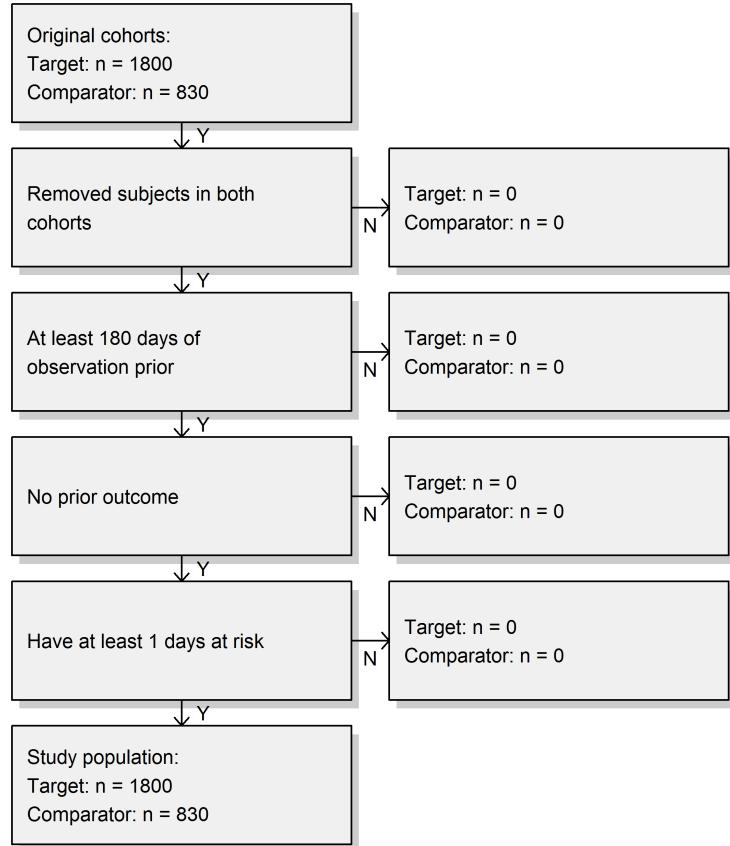
```
## CohortMethodData
##
##      ID 1
##      ID 2
##      ID 3
##
##          1800
##          830
##
##          479          479
## 3
##
##          389
##          26923
```

13.2

A horizontal row of 20 empty rectangular boxes, likely for students to write their answers in a worksheet.

```
studyPop <- createStudyPopulation(  
  cohortMethodData = cmData,  
  outcomeId = 3,  
  washoutPeriod = 180,  
  removeDuplicateSubjects = "remove all",  
  removeSubjectsWithPriorOutcome = TRUE,
```

```
riskWindowStart = 0,  
startAnchor = "cohort start",  
riskWindowEnd = 99999)  
drawAttritionDiagram(studyPop)
```



□ □ 13.3

Cox 

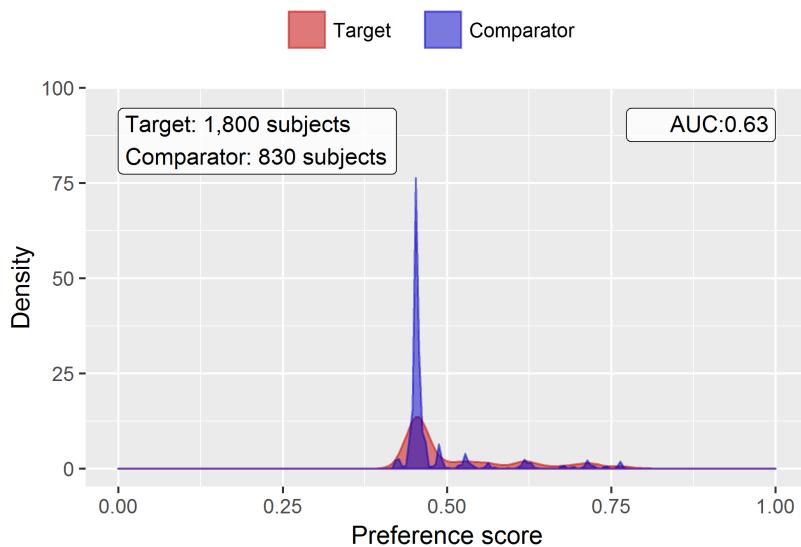
```
model <- fitOutcomeModel(population = studyPop,
                           modelType = "cox")
model
```

```
##      COX  
##      FALSE
```

```
##      FALSE
##      FALSE
##      OK
##
##          .95     .95
##      1.34612    1.10065    1.65741  0.29723    0.1044
```

□ □ 13.4

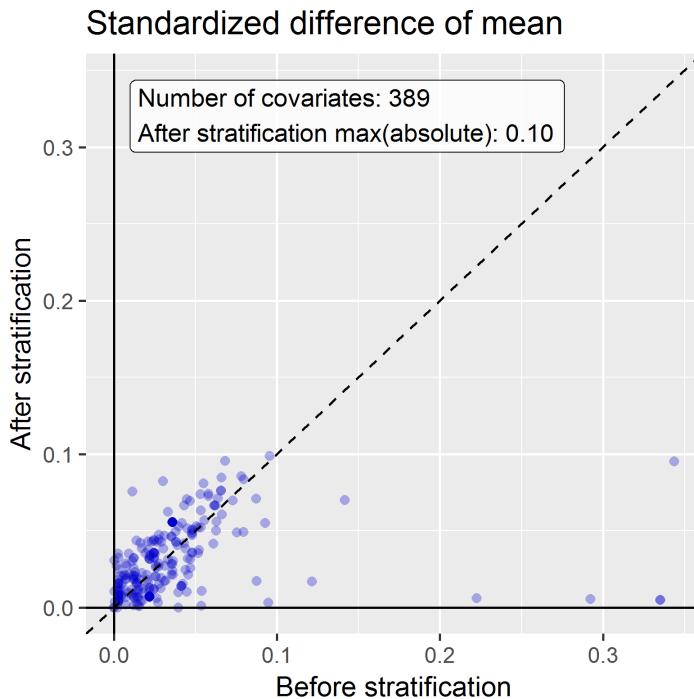
```
ps <- createPs(cohortMethodData = cmData,  
                population = studyPop)  
plotPs(ps, showCountsLabel = TRUE, showAucLabel = TRUE)
```



0.63 AUC

□ □ 13.5

```
showMaxLabel = TRUE,  
beforeLabel = "  ",  
afterLabel = "  ")
```



0.1 x > 0.3

□ □ 13.6

PS□□□□ Cox □□□□□□□□□□□□□□□□□□

```
adjModel <- fitOutcomeModel(population = strataPop,
                           modelType = "cox",
                           stratified = TRUE)
adjModel
```

```
##      cox
##      TRUE
##      FALSE
##      FALSE
##      OK
##
##          .95    .95
##      1.13211  0.92132  1.40008  0.12409  0.1068
```

E.8

□ □ 14.1

getPlpData

```
## plpData
##
##           ID -1
##     ID 3
##
##           2630
##
##           479      479
## 3
##
##           245
##
##           54079
```

□ □ 14.2

```
population <- createStudyPopulation(plpData = plpData,
                                     outcomeId = 3,
                                     washoutPeriod = 364,
                                     firstExposureOnly = FALSE,
                                     removeSubjectsWithPriorOutcome = TRUE,
                                     priorOutcomeLookback = 9999,
                                     riskWindowStart = 1,
                                     riskWindowEnd = 365,
                                     addExposureDaysToStart = FALSE,
                                     addExposureDaysToEnd = FALSE,
                                     minTimeAtRisk = 364,
                                     requireTimeAtRisk = TRUE,
                                     includeAllOutcomes = TRUE)

nrow(population)
```

```
## [1] 2578
```

□ □ 14.3

LASSO  runPlp 

```
lassoModel <- setLassoLogisticRegression(seed = 0)

lassoResults <- runPlp(population = population,
                        plpData = plpData,
                        modelSettings = lassoModel,
                        testSplit = 'person',
                        testFraction = 0.25,
                        nfold = 2,
                        splitSeed = 0)
```

## LASSO

Shiny□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□□

```
viewPlp(lassoResults)
```

E 18 AUC 0.645

The screenshot shows a Shiny application interface titled "PatientLevelPrediction Explorer". The top navigation bar includes tabs for "Internal Validation" and "External Validation". Below the navigation bar, a horizontal menu bar contains links for "Evaluation Summary", "Characterization", "ROC", "Calibration", "Demographics", "Preference", "Box Plot", and "Settings". The main content area is titled "Evaluation Summary" and displays a table of performance metrics. The table has columns for "Metric", "test", and "train". The data rows are as follows:

Metric	test	train
1 AUC	0.645	0.7112
2 AUC_lb95ci	0.589	0.6815
3 AUC_ub95ci	0.700	0.7409
4 AUPRC	0.286	0.3615
5 BrierScaled	0.062	0.0860
6 BrierScore	0.144	0.1382

At the top left of the table, there is a dropdown menu labeled "Show 25 entries". At the top right, there is a search bar labeled "Search: [ ]".

Figure E.18: □□□□□□□Shiny□□□□

## E.9

### □□ 16.1

ACHILLES□□□□□□□:

```
library(ACHILLES)
result <- achilles(connectionDetails,
                      cdmDatabaseSchema = "main",
                      resultsDatabaseSchema = "main",
                      sourceName = "Eunomia",
                      cdmVersion = "5.3.0")
```

### □□ 16.2

□□□□□□□□□□□□□□□□□□□□:

```
DataQualityDashboard::executeDqChecks(
  connectionDetails,
  cdmDatabaseSchema = "main",
  resultsDatabaseSchema = "main",
  cdmSourceName = "Eunomia",
  outputFolder = "C:/dataQualityExample")
```

### □□ 16.3

□□□□□□□□□□□□□□□□□□□□:

```
DataQualityDashboard::viewDqDashboard(  
  "C:/dataQualityExample/Eunomia/results_Eunomia.json")
```

## E.10

## E.11 NA



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