Cohort Definition Examples

Contents

```
1 Type 2 diabetes mellitus
2 Type 1 diabetes mellitus
3
3 Atrial Fibrillation
4
library(Capr)
# Function to test that cohort compiles
assertCohortCompiles <- function(cohortDefinition) {
    sql <- CirceR::cohortExpressionFromJson(as.json(ch))
    sql <- CirceR::buildCohortQuery(sql, options = CirceR::createGenerateOptions(generateStats = TRUE))
    stopifnot(nchar(sql) > 1)
    invisible(NULL)
}
```

This vignette provides a number of example cohorts built with Capr from the OHDSI *Phenotype Phebruary* event.

1 Type 2 diabetes mellitus

Persons with new type 2 diabetes mellitus at first diagnosis

https://atlas-phenotype.ohdsi.org/#/cohortdefinition/88/definition

Persons with new type 2 diabetes and no prior T1DM or secondary diabetes

https://atlas-phenotype.ohdsi.org/#/cohortdefinition/89/export

```
library(Capr)
cs0 \leftarrow cs(descendants(443238, 201820, 442793),
          descendants(exclude(195771, 201254, 435216, 761051, 4058243, 40484648)),
          name = "Type 2 diabetes mellitus (diabetes mellitus excluding T1DM and secondary)")
cs1 <- cs(descendants(201254, 435216, 40484648),
          name = "Type 1 diabetes mellitus")
cs2 <- cs(descendants(195771),
          name = "Secondary diabetes mellitus")
ch <- cohort(</pre>
  entry = entry(
    conditionOccurrence(cs0),
    observationWindow = continuousObservation(priorDays = 365)
  ),
  attrition = attrition(
    't1d' = withAll(
    exactly(0, conditionOccurrence(cs1), duringInterval(eventStarts(-Inf, 0)))
   ),
   'secondaryDiabetes' = withAll(
    exactly(0, conditionOccurrence(cs2), duringInterval(eventStarts(-Inf, 0)))
  ),
  exit = exit(
    endStrategy = observationExit()
  )
```

Persons with new type 2 diabetes mellitus at first dx rx or lab

https://atlas-phenotype.ohdsi.org/#/cohortdefinition/90

```
conditionOccurrence(cs0),
  drugExposure(cs4),
  measurement(cs3, valueAsNumber(bt(6.5, 30)), unit("%")),
  measurement(cs3, valueAsNumber(bt(48, 99)), unit("mmol/mol")),
  observationWindow = continuousObservation(priorDays = 365)
),
  attrition = attrition(
  'no T1D' = withAll(
    exactly(0, conditionOccurrence(cs1), duringInterval(eventStarts(-Inf, 0)))
    ),
  'no secondary diabettes' = withAll(
    exactly(0, conditionOccurrence(cs2), duringInterval(eventStarts(-Inf, 0)))
  ))
  exit = exit(
  endStrategy = observationExit()
)
```

2 Type 1 diabetes mellitus

Persons with new type 1 diabetes

https://atlas-phenotype.ohdsi.org/#/cohortdefinition/92

Persons with new type 1 diabetes and no prior T2DM or secondary diabetes

https://atlas-phenotype.ohdsi.org/#/cohortdefinition/91

```
),
attrition = attrition(
   "no prior T2DM" = withAll(exactly(0, conditionOccurrence(cs0), duringInterval(eventStarts(-Inf, 0))
   "no prior secondary T1DM" = withAll(exactly(0, conditionOccurrence(cs2), duringInterval(eventStarts))
)
```

3 Atrial Fibrillation

Persons with a trial fibrillation per Wharton et al 2021

https://atlas-phenotype.ohdsi.org/#/cohortdefinition/93

Persons with atrial fibrillation per Subramanya et al 2021

https://atlas-phenotype.ohdsi.org/#/cohortdefinition/94

```
afib <- cs(descendants(313217),
           name = "Atrial fibrillation")
ip \leftarrow cs(descendants(262, 9201),
         name = "Inpatient or inpatient ER visit")
op <- cs(descendants(9202, 9203),
         name = "Outpatient or ER visit")
ch <- cohort(</pre>
  entry = entry(
    conditionOccurrence(
      conceptSet = afib,
      nestedWithAll(
        # an inpatient afib
        atLeast(
          aperture = duringInterval(eventStarts(-Inf, 0), eventEnds(0, Inf)),
          query = visit(conceptSet = ip)
      )
    ),
    conditionOccurrence(
      conceptSet = afib,
      nestedWithAll(
        # 2 afib in outpatient
        atLeast(
          x = 1,
          aperture = duringInterval(eventStarts(-Inf, 0), eventEnds(0, Inf)),
          query = visit(
            conceptSet = op,
            nestedWithAll(
```