## A first example of how to extract Frequent Patterns

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## Connect to the database

```
### Define database parameters
cdmdatabaseschema = "main"
resultsdatabaseschema = "main"
connectionDetails <- Eunomia::getEunomiaConnectionDetails()
connection <- connect(connectionDetails)

## Connecting using SQLite driver

#on.exit(DatabaseConnector::disconnect(connection)) #Close db connection on error or exit

Define cohort

# Define cohort
cohort <- readSql(".../data/cohorts/Eunomia_MI_cohort.sql")</pre>
```

```
# Define cohort
cohort <- readSql("../data/cohorts/Eunomia_MI_cohort.sql")

renderTranslateExecuteSql(connection, cohort, cdm = "main")

## |
## Executing SQL took 0.0446 secs

sql <- "ALTER TABLE #diagnoses ADD cohort_definition_id INT NOT NULL DEFAULT(1)"

# Execute the script to receive the data
renderTranslateExecuteSql(connection, sql)

## |
## Executing SQL took 0.000895 secs

querySql(connection, "SELECT count(*) FROM diagnoses;")

## COUNT(*)
## 1 67</pre>
```

## Get the data and close the connection

```
# Define covariate settings
TemporalcovariateSettings_eunomia <- createTemporalCovariateSettings(useConditionOccurrence = TRUE,
                                                       temporalStartDays = seq(-(60*365), -1, by = 1),
                                                       temporalEndDays = seq(-(60*365)+1, 0, by = 1)
# Extract covariates
TemporalcovariateData_eunomia <- getDbCovariateData(connection = connection,</pre>
                         cdmDatabaseSchema = cdmdatabaseschema,
                         cohortDatabaseSchema = resultsdatabaseschema,
                         cohortTable = "diagnoses",
                         rowIdField = "subject_id",
                         covariateSettings = TemporalcovariateSettings_eunomia,
                         cohortTableIsTemp = TRUE)
## Sending temp tables to server
## Constructing features on server
## Executing SQL took 40.8 secs
## Fetching data from server
## Fetching data took 0.176 secs
disconnect(connection)
Prepare the data
getInputFileForFrequentPatterns(covariateDataObject = TemporalcovariateData_eunomia, fileToSave = "exam
## Extracting temporal data...
## Extracting covariate names...
## Generating input file for frequent pattern mining...
## Input data has been created succesfully and saved in example1.txt
Running Analysis
frequentPatterns <- runFrequentPatterns(algorithm = "SPADE", inputFile = "example1.txt", outputFile = "</pre>
## Analysing 67 sequence IDs...
## Running frequent pattern algorithm...
## Running algorithm
```

```
## [1] "java -jar ../inst/java/spmf.jar run SPADE example1.txt results_example1.txt 0.5 true"

## The command line that has been running is: java -jar ../inst/java/spmf.jar run SPADE example1.txt re

str(frequentPatterns)

## 'data.frame': 2469 obs. of 3 variables:

## $ Sequence: chr "Streptococcal sore throat " "Osteoarthritis " "Acute bronchitis " "Coronary arter

## $ Count : chr "34" "61" "60" "66" ...

## $ Support : num 0.507 0.91 0.896 0.985 0.955 ...

dfWithIDs <- getIdDataFrame("results_example1.txt")

dim(dfWithIDs)</pre>
```

## [1]

67 2470