## Project 1. Solution

Santiago Castro Dau, June Monge, Rachita Kumar, Sarah Lötscher

## Problem 1: Conditional independence and BNs

**a**)

The condition  $A \perp B|C$  holds for this Bayesian network a). Proof:

$$P(A,B,C) = P(A|C)P(B|C)P(C)$$
 Using Bayes' rule, and substituting 
$$P(A,B|C) = \frac{P(A,B,C)}{P(C)}$$
 
$$= \frac{P(A|C)P(B|C)P(C)}{P(C)}$$

= P(A|C)P(B|C) $\implies A \perp B|C$ 

Hence Proved.

In general,  $A \perp B$  does not hold for this Bayesian network. Proof:

Consider

$$\begin{split} P(A,B) &= \sum_{C} P(A,B,C) \\ &= \sum_{C} P(A|C)P(B|C)P(C) \\ &= P(A)P(B) \sum_{C} \frac{P(C|B)P(C|A)}{P(C)} \\ &\implies A \perp B \text{ does not generally hold.} \end{split}$$

b)

The condition  $A \perp B$  holds for this Bayesian network b). Proof:

$$\begin{split} P(A,B,C) &= P(A)P(B)P(C|A,B) \\ P(A,B) &= \sum_{C} P(A,B,C) \\ &= \sum_{C} P(A)P(B)P(C|A,B) \\ &= P(A)P(B) \sum_{C} P(C|A,B) \\ &= P(A)P(B) \\ &\Longrightarrow A \perp B \end{split}$$

Hence Proved.

In general,  $A \perp B | C$  does not hold for this Bayesian network. Proof:

$$P(A, B|C) = \frac{P(A, B, C)}{P(C)}$$
$$= \frac{P(C|A, B)P(A)(B)}{P(C)}$$

Multiplying and diving by P(C|A) and using Bayes' rule,

$$= \frac{P(C|A,B)P(A)P(B)}{P(C)} \frac{P(C|A)}{P(C|A)}$$
$$= \frac{P(C|A,B)P(A|C)P(B)}{P(C|A)}$$

Multiplying and diving by P(C|B)P(C) and using Bayes' rule,

$$= \frac{P(C|A,B)P(A|C)P(B)}{P(C|A)} \frac{P(C|B)}{P(C|B)}$$
$$= P(A|C)P(B|C) \frac{P(C|A,B)P(C)}{P(C|A)P(C|B)}$$

Since  $\frac{P(C|A,B)P(C)}{P(C|A)P(C|B)}$  will be equal to 1 in only rare cases, in general,  $A \perp B|C$  does not hold for this Bayesian network.

## Problem 2: Markov blanket

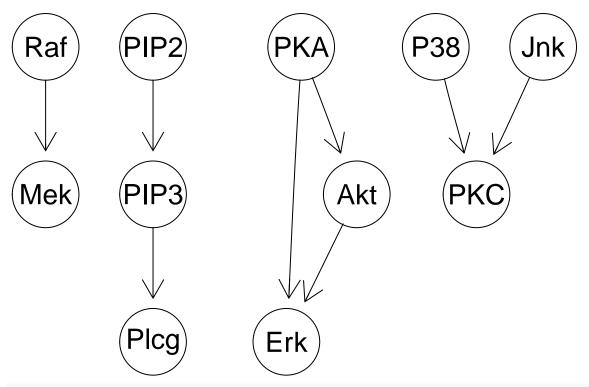
 $MB(D) = \{B, F, E, G, C\}$  Consider,

$$P(D|A, B, C, E, F, G) = \frac{P(A, B, C, D, E, F, G)}{P(A, B, C, E, F, G)}$$
  
Since  $A \perp B, C, D, E, F, G$   
$$P(D|A, B, C, E, F, G) = \frac{P(A)P(B, C, D, E, F, G)}{P(A)P(B, C, E, F, G)}$$
$$= P(D|MB(D))$$

## Problem 3: Learning Bayesian networks from protein data

a)

```
## Set seed
set.seed(2022)
## Read and index data
data <- vroom(file = "sachs.data.txt", show_col_types = FALSE)</pre>
data = data %>% rowid_to_column(var = 'row_id')
## Visualize data dimensions.
dim(data)
## [1] 853 12
  • N = 853
  • n = 11
## Split data
train = data %>% sample_frac(0.8, replace = FALSE)
test = anti_join(data, train, by = "row_id")
## Remove index column used for splitting
train = train %>% select(-row_id)
test = test %>% select(-row_id)
## Initializing score objects
score_object_train = scoreparameters(scoretype = 'bge', train)
score_object_test = scoreparameters(scoretype = 'bge', test)
b)
## Infer network on train data
infered_network = orderMCMC(score_object_train)
## Construct graph object with the inferred network
graph_object = graphAM(adjMat = infered_network$DAG,
                       edgemode='directed',
                       values=NA)
## Plot graph
plot(graph_object)
```



## Evaluate the log BGe score of the test data against the estimated DAG.
DAGscore(score\_object\_test, infered\_network\$DAG)

## [1] -9456.304

```
c)
## Define function for average BGe score and average number of edges
avg_BGe_score_and_n_edges <- function(am) {</pre>
  ## Have to load libraries inside the function otherwise it can not find other
  ## functions even if they are alredy loaded.
 library(BiDAG)
  library(dplyr)
  ## Init objects to keep track of BGe score and number of edges
  n_{edges} = c()
  BGe scores = c()
  for (i in 1:10) {
   ## Split data
   train = data %>% sample_frac(0.8, replace = FALSE)
   test = anti_join(data, train, by = "row_id")
   ## Remove id column used for splitting
   train = train %>% select(-row_id)
   test = test %>% select(-row_id)
    ## Initializing score objects
   score_object_train = scoreparameters(scoretype = 'bge',
```

train,

```
bgepar = list(am = am, aw = NULL))
    score_object_test = scoreparameters(scoretype = 'bge',
                                        test,
                                        bgepar = list(am = am, aw = NULL))
    ## Infer network on train data
    infered_network = orderMCMC(score_object_train)
   ## Append number of edges of the inferred network
   n_edges = c(n_edges, sum(infered_network$DAG))
   ## Append BGe score
   BGe_scores = c(BGe_scores, DAGscore(score_object_test,
                                        infered_network$DAG))
  }
  ## Return mean number of edges and mean BGe score
  return(list("am_value" = am,
              "mean_n_edges" = mean(n_edges),
              "mean_BGe_score" = mean(BGe_scores)))
}
## Create list of alpha mu values (am values)
am_values = list(10e-5, 10e-3, 10e-1, 10, 10e2)
## Run the each average pair (for each am value) on parallel
## You might have to change this if you computer doesn't have 5 cores!!
results = mclapply(X = am_values,
                   FUN = avg_BGe_score_and_n_edges,
                   mc.cores = 5,
                   mc.preschedule = FALSE)
## Run this in case your OS is Windows ()
# results = lapply(X = am_values,
                   FUN = avg_BGe_score_and_n_edges)
## Print results
do.call(rbind, results)
        am_value mean_n_edges mean_BGe_score
## [1,] 1e-04
                5.2
                              -9577.575
                6.9
                              -9482.639
## [2,] 0.01
## [3,] 1
                 7
                              -9383.069
## [4,] 10
                7.4
                              -9618.116
## [5,] 1000
                17.8
                              -35574.84
## The closer x is to 0 the more negative the log(x) is. Therefore the least neg
## average log(BGe_score) corresponds to the largest BGe score (hence am = 1).
## Plot relearned DAG with whole data set
## Initializing score objects
score_object_data = scoreparameters(scoretype = 'bge',
                                    data %>% select(-row_id),
                                    bgepar = list(am = 1, aw = NULL))
```

