## 732A96/TDDE15 ADVANCED MACHINE LEARNING

### LAB 1: GRAPHICAL MODELS

JOSE M. PEÑA IDA, LINKÖPING UNIVERSITY, SWEDEN

# 1. Instructions

# Deadline for individual and group reports See LISAM.

#### What and how to hand in

Each student must send a report to LISAM with his/her solutions to the lab. The file should be named FirstName\_LastName.pdf. The report must be concise but complete. It should include (i) the code implemented or the calls made to existing functions, (ii) the results of such code or calls, and (iii) explanations for (i) and (ii).

In addition, students must discuss their lab solutions in a group. Each group must compile a collaborative report that will be used for presentation at the seminar. The report should clearly state the names of the students that participated in its compilation and a short description of how each student contributed to the report. This report should be submitted via LISAM. The file should be named Group\_X.pdf where X is the group number. Please, upload also a copy of the group report to the collaborative workspace folder in LISAM. The collaborative reports are corrected and graded. The individual reports are also checked, but feedback on them will not be given. A student passes the lab if the group report passes the seminar and the individual report has reasonable quality, otherwise the student must complete his/her individual report by correcting the mistakes in it.

Attendance to the seminar is obligatory. In the seminar, some groups will be responsible for presenting their group reports. Each student in these groups must be prepared to individually present an arbitrary part of the report. The selection of the speakers is done randomly during the seminar. In the seminar, some groups will act as opponents to the reports provided by the presenters. The opponent group should examine the group report of the presenter group before the seminar (available in the collaborative workspace folder in LISAM), and prepare a minimum of three questions, comments and/or improvements. The opponent group will ask these questions during the seminar. Check LISAM for the list of presenter and opponent groups.

## Resources

The lab is designed to be solved with the R packages bnlearn and gRain. You may also want to use the RStudio development environment.

- Literature:
  - \* Package documentation.
  - \* Højsgaard, S. Graphical Independence Networks with the gRain Package for R. *Journal of Statistical Software* 46, 2012.
  - \* Scutari, S. Learning Bayesian Networks with the bnlearn R Package. *Journal of Statistical Software* 35, 2010.
- Hint: Spend the first hour exploring the site www.bnlearn.com. Try the code in www.bnlearn.com/examples.

### 2. QUESTIONS

The purpose of the lab is to put in practice some of the concepts covered in the lectures.

(1) Show that multiple runs of the hill-climbing algorithm can return non-equivalent Bayesian network (BN) structures. Explain why this happens. Use the Asia dataset which is included in the bnlearn package. To load the data, run data("asia").

**Hint**: Check the function hc in the bnlearn package. Note that you can specify the initial structure, the number of random restarts, the score, and the equivalent sample size (a.k.a imaginary sample size) in the BDeu score. You may want to use these options to answer the question. You may also want to use the functions plot, arcs, vstructs, cpdag and all.equal.

(2) Learn a BN from 80 % of the Asia dataset. The dataset is included in the bnlearn package. To load the data, run data("asia"). Learn both the structure and the parameters. Use any learning algorithm and settings that you consider appropriate. Use the BN learned to classify the remaining 20 % of the Asia dataset in two classes: S = yes and S = no. In other words, compute the posterior probability distribution of S for each case and classify it in the most likely class. To do so, you **have** to use exact or approximate inference with the help of the bnlearn and gRain packages, i.e. you are not allowed to use functions such as predict. Report the confusion matrix, i.e. true/false positives/negatives. Compare your results with those of the true Asia BN, which can be obtained by running

```
dag = model2network("[A][S][T|A][L|S][B|S][D|B:E][E|T:L][X|E]").
```

Hint: You already know the Lauritzen-Spiegelhalter algorithm for inference in BNs, which is an exact algorithm. There are also approximate algorithms for when the exact ones are too demanding computationally. For exact inference, you may need the functions bn.fit and as.grain from the bnlearn package, and the functions compile, setFinding and querygrain from the package gRain. For approximate inference, you may need the functions prop.table, table and cpdist from the bnlearn package. When you try to load the package gRain, you will get an error as the package RBGL cannot be found. You have to install this package by running the following two commands (answer no to any offer to update packages):

```
source("https://bioconductor.org/biocLite.R")
biocLite("RBGL")
```

(3) In the previous exercise, you classified the variable S given observations for all the rest of the variables. Now, you are asked to classify S given observations only for the so-called Markov blanket of S, i.e. its parents plus its children plus the parents of its children minus S itself. Report again the confusion matrix.

**Hint**: You may want to use the function mb from the bnlearn package.

(4) Repeat the exercise (2) using a naive Bayes classifier, i.e. the predictive variables are independent given the class variable. See p. 380 in Bishop's book or Wikipedia for more information on the naive Bayes classifier. Model the naive Bayes classifier as a BN. You have to create the BN by hand, i.e. you are not allowed to use the function naive.bayes from the bnlearn package.

**Hint**: Check http://www.bnlearn.com/examples/dag/ to see how to create a BN by hand.

(5) Explain why you obtain the same or different results in the exercises (2-4).