### Bayesian Learning

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# Learning Bayesian Models

How can we learn a Bayesian model from a database of past cases?

We consider two basic problems:

- Parameter learning: We have an structure for the Bayesian Network, but we miss the conditional probabilities associated with the factorization
- ② Structure learning: We have neither the structure nor the conditional probabilities



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Maximum Likelihood Estimation (MLE): Given a model BN and database of cases D, what are the most probable parameters of the corresponding probability tables for BN?

For a BN with n variables, we need to estimate n probability tables

The probability table for a variable  $x_i$  where the total number of cases to be considered with its set of parent variables is  $q_i$  has

$$(r_i - 1)q_i$$

independent probability parameters



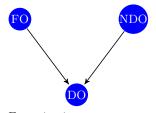


Factorization:

P(DO|FO)P(FO)

Independent parameters:

- For P(FO) :  $\{P(FO=0)\}$
- For P(DO|FO) :  $\{P(DO=0|FO=0),$ P(DO = 0|FO = 1)



Factorization:

Independent parameters:

- For  $P(FO) : \{P(FO=0)\}$
- For  $P(NDO) : \{P(NDO=0)\}$

P(DO|FO,NDO)P(FO)P(NDO)

• For P(DO|FO,NDO):  $\{P(DO = 0|FO = 0, NDO =$ 0), P(DO = 0 | FO = 0, NDO =1), P(DO = 0|FO = 1, NDO =0), P(DO = 0|FO = 1, NDO = 1)



Given a BN, a variable  $x_i$ , our database D, and a vector of values  $w_{i,j}$  assigned to its vector of parent variables  $\pi_i$ , the MLE estimation for the conditional probability:

$$\theta_{i,j,k} = \Pr(x_i = k \mid \pi_i = w_{i,j})$$

is

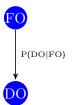
$$\hat{\theta_{i,j,k}} = \frac{N_{i,j,k}}{N_{i,j}}$$

assuming multinomial random variables, where:

- $N_{i,j,k} = |\{r \mid r \in D, r[x_i = k, \pi_i = w_{i,j}]\}|$ . That is, the number of rows in D where the parent variables  $\pi_i$  have assigned the values  $w_{i,j}$  and  $x_i$  the value k
- $N_{i,j} = \sum_{k=1}^{r_i} N_{i,j,k}$



Consider the following BN for the FO-DO problem and a database of cases D



Estimations:

• 
$$P(FO = 0) = \frac{4}{8}$$

• 
$$P(DO = 0|FO = 0) = \frac{2}{4},$$
  
 $P(DO = 0|FO = 1) = \frac{1}{4}$ 

DO	FO
0	0
0	0
1	0
1	0
0	1
1	1
1	1
1	1



Consider the alternative BN for the FO-DO problem and the same database of cases D



Estimations:

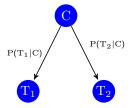
- $P(DO = 0) = \frac{3}{8}$
- $P(FO=0|DO=0) = \frac{2}{3},$   $P(FO=0|DO=1) = \frac{2}{5}$

DO	FO
0	0
0	0
1	0
1	0
0	1
1	1
1	1
_1	1

What model predicts with less uncertainty the values for the variables?



Consider the following BN for the two-tests cancer example



Such that we do not know its corresponding probability tables, but we have instead a database of samples, obtained from past clinical records



Suppose the following database D for the cancer problem:

С	T1	T2
0	0	1
0	0	0
0	0	0
0	1	0
0	0	0
0	0	0
0	0	0
1	1	1

We get the estimations:

• 
$$P(C = 0) = \frac{7}{8}$$

• 
$$P(T1=0|C=0) = \frac{6}{7},$$
  
 $P(T1=0|C=1) = \frac{0}{1}$ 

• 
$$P(T2=0|C=0) = \frac{6}{7},$$
  
 $P(T2=0|C=1) = \frac{0}{1}$ 



Observe that depending on the particular database D, some probabilities may be estimated to be equal to 0, even if the real probability could be very small, but not 0!

As this can produce problems when performing inference, some learning algorithms add an small bias factor so that probabilities equal to 0 are replaced by very small probabilities



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For example, with the previous estimations for the two-tests cancer example, if we now want to compute:

$$P(C=1|T_1=1,T_2=0) = \frac{P(C=1)P(T_1=1|C=1)P(T_2=0|C=1)}{P(T_1=1,T_2=0)}$$

we get the following answer:

$$\frac{1}{8} \frac{1}{1} \frac{0}{1} \frac{1}{P(T_1 = 1, T_2 = 0)} = 0$$

So, because one of the factors was 0, independently of the fact that other factors were greater than 0, the final result is 0.

This crisp result is not very helpful!



### Learning the Structure of a Bayesian Network

What is the best Bayesian Network that explains a given data set D?

Ingredients for a learning algorithm for Bayesian Networks:

- Scoring: When a Bayesian Network is better than other for explaining D?
- 2 Search space: What is the search space of possible Bayesian Networks we want to consider?
- Search strategy: If the search space has exponential size, how do we search the best Bayesian Network?



Given our data set D, and a possible Bayesian Network BN for our problem, a possible scoring model for how good is BN is computing the probability of obtaining all the samples in D from BN:

### P(D|BN)

The higher that value is, the more confident we can be with the model BN because the data set D will be closer to what BN tipically will generate when used as a model for simulating the outcomes obtained from the real problem



Observe that from probability theory we have that:

$$P(D|BN)P(BN) = P(D,BN)$$

Typical scoring metrics developed for bayesian learning algorithms use rather P(D,BN) than P(D|BN), so that it is possible to include a prior probability P(BN) on the space of possible bayesian network models

That is, to induce some bias towards some preferred bayesian models, if we have additional information that may indicate us that some models are more natural than others



We can compute P(D,BN) integrating over all the possible probability distributions  $B_p$  consistent with BN:

$$P(D,BN) = P(BN) \int_{B_p} (P(D|BN,B_p)f(B_p|BN)dB_p)$$

### Where:

- P(D|BN, B<sub>p</sub>) is the probability of observing the samples in D, assuming the data is generated with network BN and probability distribution B<sub>p</sub>.
- $f(B_p|BN)$  is the probability of having probability distribution  $B_p$  assuming the correct network is BN, and assuming that all the  $B_p$  consistent with BN are uniformly distributed



In UPSM, the set of prior probabilities over the set of Bayes networks (BN) follows an uniform distribution (all the BN are equally likely in the absence of any data set D)

So, we can substitute P(BN) by an arbitrary constant C:

$$P(D,BN) = C \int_{B_p} P(D|BN,B_p) f(B_p|BN) dB_p$$



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Assuming all the samples  $r_h$  from D are independently generated:

$$P(D|BN, B_p) = \prod_{h=1}^{|D|} P(r_h|BN, B_p)$$

Considering the factorization given by BN and B<sub>p</sub>:

$$\prod_{h=1}^{|D|} P(r_h|BN, B_p) = \prod_{i=1}^n \prod_{i=1}^{q_i} \prod_{k=1}^{r_i} P(x_i = k | \pi_i = w_{i,j})^{N_{i,j,k}}$$

Finally, working further with the term  $f(B_p|BN)$  we obtain:

$$P(D,BN) = C \prod_{i=1}^n \prod_{j=1}^{q_i} (r_i-1)! \int_{\theta_{i,j,1}} \dots \int_{\theta_{i,j,r_i}} \prod_{k=1}^{r_i} \theta_{i,j,k}^{N_{i,j,k}} d\theta_{i,j,1} \dots d\theta_{i,j,r_i}$$



And working with Dirichlet's integrals:

$$P(D,BN) = C \prod_{i=1}^{n} \prod_{j=1}^{q_i} \frac{(r_i-1)!}{(N_{i,j}+r_i-1)!} \prod_{k=1}^{r_i} N_{i,j,k}!$$

This scoring scheme gives a higher value to a BN that predicts with higher probability the samples in D

Thats is, to a BN where the samples in D are more likely to be obtained.

Check the original paper where this scoring model (and the learning algorithm K2 that uses it) is presented!



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Example: Consider the two possible models for the FO-DO problem and our previous database D

#### Estimations:

• 
$$P(FO = 0) = \frac{4}{9}$$

• 
$$P(DO = 0|FO = 0) = \frac{2}{4},$$
  
 $P(DO = 0|FO = 1) = \frac{1}{4}$ 

• UPSM:

• For FO: 
$$\frac{4!4!}{(8+1)!}$$
  
• For DO:  $\frac{2!2!}{(4+1)!} \frac{3!1!}{(4+1)!}$ 

• Final product: 0,00002646

#### Estimations:

• 
$$P(DO = 0) = \frac{3}{8}$$

• 
$$P(FO=0|DO=0) = \frac{2}{3}$$
,  
  $P(FO=0|DO=1) = \frac{2}{5}$ 

• UPSM:

• Final product: 0,00002756

So, the model P(DO)P(FO|DO) seems to be slightly better (from the point of view of UPSM)



Example: Consider the two following models for the two-tests cancer problem and our previous database D

Factorization for BN<sub>1</sub>:

Factorization for 
$$BN_2$$
:

$$P(C)P(T_1|C)P(T_2|C)$$

$$P(T_1)P(C|T_1)P(T_2|C)$$

Factors for  $P(BN_1, D)$ :

Factors for  $P(BN_2, D)$ :

$$\begin{array}{c|cccc} & j=0 & j=1 \\ \hline T_1 & \frac{6!1!}{(7+1)!} & \frac{0!1!}{(1+1)!} \\ T_2 & \frac{6!1!}{(7+1)!} & \frac{0!1!}{(1+1)!} \\ C & \frac{7!1!}{(8+1)!} \end{array}$$

$$P(BN_1, D) = \frac{1}{9.8^3 \cdot 7^2 \cdot 4} = 11.07 \cdot 10^{-6} \ P(BN_2, D) = \frac{1}{9.8^2 \cdot 7^3 \cdot 6} = 8.44 \cdot 10^{-7}$$
  
So, BN<sub>1</sub> seems to be a better model for explaining D with more accuracy



When has the factor associated with a variable  $x_i$  and parent row j in the score P(BN, D):

$$\frac{(r_i-1)!}{(N_{i,j}+r_i-1)!}\prod_{k=1}^{r_i}N_{i,j,k}!$$

a higher score?

This factor is associated with the conditional probability:

$$P(x_i = k | \pi_i = w_{i,j})$$



### Best case for UPSM: No uncertainty on x<sub>i</sub>

There is a case v with  $N_{i,j,v} = N_{i,j}$  (the other v' terms are 0). Then:

$$\frac{(r_i-1)!}{(N_{i,j}+r_i-1)!}\prod_{k=1}^{r_i}N_{i,j,k}! = \frac{(r_i-1)!N_{i,j}!}{(N_{i,j}+r_i-1)!}$$

That is, with parent case j the model predicts a unique possible value v for  $x_i$  (no uncertainty, perfect prediction!) Even better case: when

$$N_{i,j,v} = N_{i,j}$$
 is much bigger than  $r_i$   $(r_i = o(N_{i,j}))$ 



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### Worst case for UPSM: Maximal uncertainty on x<sub>i</sub>

All the  $N_{i,j,k}$  are equal:  $N_{i,j,k} = N_{i,j}/r_i$ 

$$\frac{(r_i-1)!}{(N_{i,j}+r_i-1)!} \prod_{k=1}^{r_i} N_{i,j,k}! = \left(\frac{N_{i,j}}{r_i}\right)!^{r_i} \frac{(r_i-1)!}{(N_{i,j}+r_i-1)!}$$

That is, maximal uncertainty on the value predicted for  $x_i$  (all the possible values equally likely)

Even worse case: when  $N_{i,j,k} = N_{i,j}/r_i = 1$  (so  $N_{i,j} = r_i$ ). Then:

$$\left(\frac{N_{i,j}}{r_i}\right)!^{r_i}\frac{(r_i-1)!}{(N_{i,i}+r_i-1)!}=\frac{(r_i-1)!}{(r_i+r_i-1)!}$$



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### Some relevant properties of the UPSM scoring model:

- Local scoring scheme: the factor contributed to the score by a node  $x_i$  depends only on its set of parents  $\pi_i$ , so it does not change when we change the parent set of other nodes
- Incremental computation: Due to the local scoring property, it can be computed incrementally for a BN that is incrementally build: this is helpful for the efficient implementation of greedy search algorithms



# Scoring: Penalising Too Complex Models

Remember that in machine learning, a general principle is that simpler hypothesis are preferred when many of them seem to be equally good

Too complex hypothesis can overfit to the data: they will predict well our data set, but give poor answers with new data

With UPSM, adding more parent nodes to a node  $x_i$  will almost always increase its scoring, but it may lead to overfit as the model could capture particular dependencies of the data set D



# Scoring: Penalising Too Complex Models

Remember that in machine learning, a general principle is that simpler hypotheses are preferred when many of them seem to be equally good

Too complex hypotheses can overfit to the data: they will predict well our data set, but give poor answers with new data



# Scoring: Penalising Too Complex Models

A possible way to penalise too complex models is to substract from the UPSM scoring for a BN a measure of the size of the model:

$$\frac{\log |D|}{2} \sum_{i=1}^{n} q_i (r_i - 1)$$

where remember that  $q_i$  is the number of different cases associated with the parents  $\pi_i$  of  $x_i$  on the BN being scored

The scoring that incorporates the substraction of such model size is called Bayesian Information Criterion (BIC) or Minimum Description Length (MDL)



### Looking For a Good Model

Once we have a way of scoring how good is a BN model with respect to our database D, the next step is to pick a search algorithm for learning the best possible BN for our problem, given our database D

In principle, the best option would be to perform systematic search trough the space of possible Bayesian networks that explain D

But even with the restriction that a BN is an acyclicic graph, the search space is still of exponential size!

So, the machine learning methods developed for Bayesian networks have focused on incomplete methods, based mainly on local search



# A Greedy Local Search Algorithm: K2

#### How it works:

- Generate some ordering between the nodes
- Initialize the set of parents for every node to the empty set
- For each node  $x_i$ , incrementally and in a greedy fashion, selects its set of parent nodes  $\pi_i$  from the set of nodes that precede it in the ordering using UPSM to score each possible parent

Even if we work with a fixed ordering, still the possible number of Bayesian networks with n nodes is exponential:

$$2^{n(n-1)/2}$$

Exercise: Use Combinatorial Analysis to check why this is the number of networks for a fixed ordering between the nodes



# A Greedy Local Search Algorithm: K2

### Function LearnBNetK2(D ,Ordering, u)

```
for each v<sub>i</sub> in Ordering do
     \pi_{i} := \{\};
     P_{\text{old}}(x_i) := UPSM(x_i, \pi_i, D);
     improving := true :
     while improving \wedge |\pi_i| < u do
           Let z be a node \in Prev(x<sub>i</sub>, Ordering) \ \pi_i that maximizes
           UPSM(x_i, \pi_i \cup \{z\}, D);
           P_{new}(x_i) := UPSM(x_i, \pi_i \cup \{z\}, D);
          if P_{new}(x_i) > P_{old}(x_i) then
                P_{old}(x_i) := P_{new}(x_i);
               \pi_i := \pi_i \cup \{z\}:
           else
                improving := false;
```



### A Greedy Local Search Algorithm: K2

Using an upper bound on the number of parents for each node (u) on the algorithm is a way to overcome a possible overfit of the model obtained

But of course, it is not clear how to select at the beginning a good value for u

The local search scheme used by K2 can actually be used with any other scoring model that has the property of being local: the scoring given by a node  $x_i$  depends only on its set of parents  $\pi_i$ , and not on the parents of other nodes



# Greedy Local Search is not Perfect

Database D:

$x_i$	$p_1$	$p_2$	$p_3$
1	0	0	1
0	0	0	0
1	0	1	0
0	0	1	1
0	1	1	1
1	1	1	0
0	1	0	0
1	1	0	1

Consider finding the best parent set for  $x_i$  with at most two nodes, taken from  $\{p_1, p_2, p_3\}$ First, observe that:

UPSM(x<sub>i</sub>, {}, D) = 
$$\frac{4!4!}{9!} = \frac{1}{630}$$

But for any pi:

$$UPSM(x_i, \{p_j\}, D) = \frac{2!2!}{5!} \frac{2!2!}{5!} = \frac{1}{900}$$



### Greedy Local Search is not Perfect

Database D:

$x_i$	$p_1$	$p_2$	$p_3$
1	0	0	1
0	0	0	0
1	0	1	0
0	0	1	1
0	1	1	1
1	1	1	0
0	1	0	0
1	1	0	1

Then, we have that starting with any p<sub>i</sub>:

$$UPSM(x_i, \{p_j\}, D) = \frac{2!2!}{5!} \frac{2!2!}{5!} = \frac{1}{900}$$

but the effect of adding a second parent depends on the first one:

• If we start with  $p_1$ , with  $p_i = p_2$  or  $p_i = p_3$ :

$$UPSM(x_i,\{p_1,p_j\},D) = \left(\frac{1!1!}{3!}\right)^4 = \frac{1}{1296}$$



### Greedy Local Search is not Perfect

On the other hand, if we extend the maximum number of parents to three, starting with parent set  $\{p_1\}$  the previous computations show that it makes no profit to add a second parent, but observe that:

Database D:

$x_i$	$p_1$	$p_2$	$p_3$
1	0	0	1
0	0	0	0
1	0	1	0
0	0	1	1
0	1	1	1
1	1	1	0
0	1	0	0
1	1	0	1

UPSM(x<sub>i</sub>, {p<sub>1</sub>, p<sub>2</sub>, p<sub>3</sub>}, D) = 
$$\left(\frac{1!}{2!}\right)^8 = \frac{1}{256}$$

So, even if adding a second parent node to  $\{p_1\}$  does not improve the score, adding it two more parents leads to a better score than only with  $\{p_1\}$ 

But still observe that the best global solution is the parent set:  $\{p_2, p_3\}$  that is never reached starting from  $\{p_1\}$ 

Concluding: the greedy local search approach does not guarantee to find the best global set of parents!



The Speed Dating Problem

### Speed Dating

From wikipedia: Speed dating is a formalised matchmaking process or dating system whose purpose is to encourage people to meet a large number of new people

How can we know when two persons are gonna match each other?



Consider the following database of previous dates with a client, whith characteristics of the mates, and whether they were liked by our client

ideo	sex	talks	vege	blade	youlike
izq	mas	nada	$_{ m si}$	si	si
izq	fem	nada	si	no	no
izq	fem	nada	$_{ m si}$	si	si
cen	mas	poco	si	no	no
cen	mas	poco	no	si	si
cen	mas	poco	no	no	no
$\operatorname{der}$	mas	mucho	no	si	no
izq	fem	mucho	no	no	si
$\operatorname{der}$	fem	mucho	no	si	no
izq	$_{ m fem}$	mucho	no	no	si

Is there any sound pattern with sex, blade and youlike?



Can we learn a model to explain what makes people to be liked by our client?

If we can learn this, we can try to reduce the number of unsuccessful meetings between our client and other people

Many dating companies on internet can take profit of such learning process

A possibility is to learn a bayesian model for the joint probability distribution of the variables of the problem



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We run 10 times the K2 algorithm with a random ordering of its variables in each run and maximum number of parents 3. Results ordered by model score

Run #	Log(BayesScore)	Total # of edges
3	-51.52	7
10	-51.78	8
1	-52.22	8
5	-52.94	6
2	-54.31	9
9	-54.92	5
7	-55.11	6
8	-55.16	6
4	-56.32	6
6	-57.32	3



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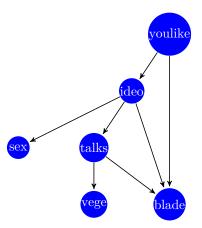
Suppose the following new person arrives ideo sex talks vege blade youlike izquierdas mas nada yes no yes

But if the person in new, we still do not know the value for the variable youlike. But we would like to predict it!

So, we would like to know how much likely are the results youlike=yes and youlike=no, given the known attributes of the new person (conditioned to them)



Bayesian network of best model found (# 3):





How well our ten learned models predict the value of the variable youlike for our test instance with known attribute values E ?

Run #	Log(BayesScore)	P(youlike = yes E)
3	-51.52	0.156
10	-51.78	0.083
1	-52.22	0.083
5	-52.94	0.156
2	-54.31	0.083
9	-54.92	0.885
7	-55.11	0.67
8	-55.16	0.656
4	-56.32	0.67
6	-57.32	0.5



Observe that the models with the lowest bayes score (last 5 models of the previous table) are the ones that predict with high probability the right answer

In the data set used for learning, with these attributes the answer is almost always youlike=no

So, it is reasonable that a learning method focused on adjusting well to the training data (to explain it with high confidence) should give models where for these attributes the answer with highest probability is youlike=no

So, we have here a problem derived by the overfitting to the training data, that is a common problem with many other learning techniques

