

Two distinct problems

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1. You know all the variables, but some values are missing in some

instances, e.g.,

Х	Y	Z
0 1 0	1 ? ?	1 0 ?
	• • •	

This makes the search problem a lot harder, but still doable

2. There are hidden (latent) variables which you *never* observe,

e.g.

0			
Х	Y	Z	Н
0 1 0	1 1 0	1 0 0	

Scoring structure using MDL

• Recall that for a graph G, the MDL score has the form:

$$score_{MDL} = m \sum_{i} MI_{\hat{p}}(X_{i}, Parents(X_{i})) - Penalty(G)$$

- To get p̂, we need to compute the parameters of the graph G, from our incomplete data
- Simple idea: use gradient descent or EM to compute (as best we can) max. likelihood parameters given the data
- The penalty term depends on the size of the graph, <u>not</u> the parameters, so it will not be affected.

A simple algorithm

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- 1. Start with a graph structure G
- 2. Repeat as long as desired:
 - (a) Consider all graphs G' that can be obtained by adding or deleting an arc from G (these are G's successors)
 - (b) For each structure G', run EM (or gradient ascent) to fit its parameters.
 - (c) Compute score MDL(G') for each G'
 - (d) Pick a G' out of the candidates using your favorite method (e.g., greedily or using simulated annealing)

(e) $G \leftarrow G'$

The simple algorithm is too slow!

- If we have n random variables, <u>in each search step</u> there are n² possible successors for G (we can pick any pair of variables and add an arc, if none is there, or remove an arc, if they are connected
- Of course, this is a worst-case estimate, because some of the resulting structures may be illegal
- Finding the parameters of the network requires some number of EM iterations
- Then to compute the score, we need to compute the likelihood of the data, which is basically a step of inference
- We need a better idea!

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Structural EM (Friedman, 1997)

- Recall the interpretation of the EM algorithm in parameter estimation
 - Start with a guess for the parameters
 - <u>Complete the data</u> by assigning the most likely values to the missing variables.
 - Improve the parameter guess based on the completed data, and iterate
- So let's *use our current network G* to complete the data!
- In our previous algorithm, we completed the data separately using each successor *G*[']
- But G and G' differ only by one arc!
- So using G to complete the data cannot be too bad...



Example: Two versions of the algorithm

- Hard EM: pick the <u>most likely values</u> for Y^2 and Z^3 , then install them and use the resulting data set to score the successors G'
- Soft EM
 - Consider <u>all possible</u> assignments of values for Y^2 and Z^3 , which gives us several completed data sets
 - The score for the successors G' is obtained as an <u>expected value</u>, by averaging the scores obtained from each data set

Example: Soft EM

- Consider all possible combinations of values for Y^2 and Z^3 : $\langle Y^2 = 0, Z^3 = 0 \rangle, \langle Y^2 = 1, Z^3 = 0 \rangle, ...$
- This gives us 4 data sets, call them D_{00} , D_{01} , D_{10} , D_{11}
- Because the data is i.i.d., the likelihood of each data set is: $p(D_{ij}) = p(Y = i | X = 1, Z = 0, \langle G, \theta \rangle) p(Z = j | X = 0, Y = 1, \langle G, \theta \rangle)$
- For every G', evaluate 4 scores, score_{*ij*}(G'), one corresponding to each completion of the data, D_{ij}

$$\operatorname{score}_{MDL}(G') \approx \sum_{i \in \{0,1\}} \sum_{j \in \{0,1\}} p(D_{ij}) \operatorname{score}_{ij}(G')$$

• Note that the number of data sets created in the "soft" version is *exponential* in the number of missing values

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Making the algorithm more efficient

- Recall from lecture 16 that the likelihood of the (complete) data can be *decomposed* based on the network structure
- Likewise, the MDL score can be computed by looking at the mutual information of a node and it parents, which can be computed locally at each node, using counts
- So we keep sufficient statistics (counts) at each node
- The fact that there are missing values only means we need to keep alternate counts *at the nodes* for which values are missing.
- When going from G to a successor G', we recompute the score only for the families that are affected
- Every *k*th search step, we have to do EM again to compute a new completion of the data set

Theoretical properties

• For any two graphs G_1 and G_2 , we have:

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score_{MDL}(G_2) - score_{MDL}(G_1)

\geq E[score_{MDL}(G_2)|completed data] - E[score_{MDL}(G_1)|completed data]
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- So if SEM moves from graph G_1 to a graph G_2 that seems to have a better expected MDL score (according to the possible data completions), then the true MDL score of G_2 is also better than the true MDL score of G_1
- The difference between the two MDL scores is at least as big as predicted by SEM
- Hence, the score is guaranteed to converge to a local maximum
- Of course, like in regular EM, multiple restarts will help get a better network in the end.

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What about Bayesian scoring?

• Recall the Bayesian score:

 $score_{\mathsf{Bayes}} = p(G|D) = p(G)p(D|G) = p(G)\int p(D|G,\theta)p(\theta|G)d\theta$

- We have to evaluate the integral for all graphs G!
- Evaluating the integral can also be quite expensive!
 - We can pick a few graphs that are most likely, and evaluate it only for those
 - Alternatively, use stochastic integration, but it turns nasty...

Computational hardship

- The computation of parameters for every candidate is very expensive
- We cannot tell beforehand whether it's really worth doing it (how good will a candidate be?)
- Works only if we limit the search space to a small number of networks

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Dealing with hidden variables

This is much harder!

- How can we tell there is something hidden?
- How many hidden variables should be introduce?
- How should they link to the rest of the network?



How do we get the structure with hidden variables?

- How many hidden variables should be introduce?
 - As few as possible! Most applications introduce at most one....
- How should they link to the rest of the network?
 - Make a guess for an the structure (e.g. by looking at large cliques or strongly connected subsets of nodes)
 - Then use EM to estimate parameters!

Hidden variables: Case study (Heckerman)

- Complete data from over 10000 Wisconsin high school graduates: sex (2 values), socio-economic status (4 values), IQ (4 values), parental encouragement (2 values), college plans (2 values)
- Goal is to find causal relationships between the variables
- Best structure found:



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Hidden variables: Case study (2)

- They considered adding 1-4 hidden variables, each with between 2-6 possible values.
- Best structure has one hidden variable, *H*,with two possible values



- This is $2 \cdot 10^{10}$ more likely than the previous best!
- In general, bushy networks are an indication of potential hidden variables