We wish to learn both the parameters and structure of a Bayesian Network.

# Parameters

Consider the disease model from the previous lecture:

![](https	:	//paper	_
attachments. drop box. com/s	$E_E 29353D8DE6A7F32069419A77A11$	E9F7A8BD49E718AC081FD5B	$F77701FB86FF68_{1}5909054763$
drawing.jpg)			

Cases 1 and 3 have incomplete data, whereas 2 had **complete**. If every example is complete, the data set is called complete. If complete, the maximum likelihood parameters are unique. We find the likelihood of a parameter set like so:

 $![](https : //paper - attachments.dropbox.com/s_E29353D8DE6A7F32069419A77A11E9F7A8BD49E718AC081FD5F77701FB86FF68_15909057905drawing + 1.jpg)$ 

Parameter Estimation With Complete Data

![](https//paperdrawing + 2.jpg

The large table is our empirical distribution .

This is formed by collapsing our distribution.  $\theta \{ \neg S | H \} = \frac{Pr(\neg s, h)}{Pr(h)} = \frac{5}{6}.$ 

$$\theta_{\lbrace} \neg S | H \rbrace = \frac{Pr(\neg s, h)}{Pr(h)} = \frac{5}{6}.$$

This is our maximum likelihood parameter estimate!

### Parameter Estimation With Incomplete Data

This uses an iterative algorithm called **expected maximum**.

Say we wish to fill the row <h, s, ?>.

We start with an arbitrary guess at the CPT.

$$CPT_1 \rightarrow BN_1 \rightarrow Pr_1(.).$$

We must choose from {TFT, TFF}, so we assign probabilities according to the probability function

$$x = Pr1(e|h, \neg s)$$

$$y = Pr(\neg e|h, \neg s)$$

We then take the assignment with the greater probability, yielding

$$CPT_2 \to BN_2 \to Pr_2(.)$$
.

We only needed to surmise one term here, but we may need to iterate.

The probabilities Pr1(.) & Pr2(.) are guaranteed to be non-increasing, so this converges.

This is thus a form of local search algorithm, so we know we may have to run repeatedly.

Note: this is the thinking behind the original algorithm, but it isn't how it is performed in practice.

#### Structure

Consider the following three structures:

 $![](https : //paper - attachments.dropbox.com/s_E29353D8DE6A7F32069419A77A11E9F7A8BD49E718AC081FD5F77701FB86FF68_15909086033drawing + 4.jpg)$ 

What do I need to optimize to choose between these three structures? We have learned many of these algorithms, so we won't discuss in detail, but:

- 1. local search methods ( approximate methods)
  - $\rightarrow$  we transform the structure looking for a better score

We must thus specify movement within the neighborhood structure:  $\rightarrow$  legal operations: add/remove/reverse edge

- 2. systemic search methods (exact methods)
  - $\rightarrow$  A\* is a good example

Why can't we just use maximum likelihood? We face overfitting.

Say we are using likelihood to compare; then  $C > B > \overline{A}$ .

We will thus end up with a complete DAG, no matter what.

Thus we need a model that balances structure complexity with likelihood.

Why? Consider:

 $![](https : //paper - attachments.dropbox.com/s_E29353D8DE6A7F32069419A77A11E9F7A8BD49E718AC081FD5F77701FB86FF68_15909090216drawing + 5.jpg)$ 

The data is clearly a linear fit, but if we estimate to the fourth degree, we get bad data! There is no fixed answer to the problem; we clearly need some function of the form f(likelihood) - g(complexity).

A common one is called MDK, but we need not discuss details.

## Model-Oriented Vs. Query-Oriented Learning

This can also be referred to as (unsupervised vs supervised) or (unlabeled vs labeled). We have done model learning now; we move on to query based.

A model-based approach might give us the following:

 $![](https : //paper - attachments.dropbox.com/s_E29353D8DE6A7F32069419A77A11E9F7A8BD49E718AC081FD5F77701FB86FF68_15909096142drawing + 6.jpg)$ 

Say we promise only to infer upward:

Then we don't 'model' the system and instead prepare for a specific 'query'. Labeling of correct responses is done by humans, and this is done to 'train'. If we want to talk about supervised learning, we need to introduce

## **Arithmetic Circuits**

 $![](https : //paper - attachments.dropbox.com/s_E29353D8DE6A7F32069419A77A11E9F7A8BD49E718AC081FD5F77701FB86FF68_15909102939Shot + 2020 - 05 - 31 + at + 12.30.28 + AM.png)$ 

The (+) symbols represent OR gates The (\*) symbols represent AND gates The  $\theta$  are the parameters The  $\lambda$  are the evidence

When given a query, we can perform weighted model counting on the equivalent arithmetic circuit. If A=True,  $\lambda_A=1$ ; if A=False,  $\lambda_{\neg A}=1$ ; if unknown, both are 1 Thus we can evaluate this in linear time; unfortunately converting to an arithmetic circuit is  $O(nd^w)$ 

BUT what do we do if we don't have the parameters? We use the labeled data:

 $![](https : //paper - attachments.dropbox.com/s_E29353D8DE6A7F32069419A77A11E9F7A8BD49E718AC081FD5F77701FB86FF68_15909113848drawing + 7.jpg)$ 

**Cross Entropy** gives us a measure of the disagreement between the two.

Therefore, we often seek to minimize it across a structure.

We generally use it to perform gradient descent (nth dimensional hill climbing).

Consider the example of recognizing shapes:

 $![](https : //paper - attachments.dropbox.com/s_E29353D8DE6A7F32069419A77A11E9F7A8BD49E718AC081FD5F77701FB86FF68_15909119570drawing + 8.jpg)$ 

Though pixel is functional in general, we allow it to be probabilistic to permit noise.

It will thus be inferred to be very close to 0/1.

A lot of heights will be zeroed as well depending on the column.

We can thus see that we have a lot of **background knowledge** for which we could simply substitute and reduce cases.

The simplest case of this is  $A \iff B$ ,