

![CausalGraphandConditionalProbabilityTables](https://paperkit.net/assets/images/illustrations/illustration-10.png) : <https://paperkit.net/assets/images/illustrations/illustration-10.png> –
attachments.dropbox.com/s_5F1FEB98332EEC99C328935F7AB0E929534BB33EF2AB134DF14A85072ADC65A01590219806
Shot + 2020 - 05 - 23 + at + 12.43.14 + AM.png)

Variables:

C — condition

T's — tests to detect the condition

S — sex of the patient

We first compute the marginal distributions to discuss probabilities.

These come in two major forms:

attachments.dropbox.com/s_5F1FEB98332EEC99C328935F7AB0E929534BB33EF2AB134DF14A85072ADC65A01590220321drawing + 5.jpg)

We then use this information to find the **Most Probable Explanation**:

We assume the final consequence (A) and search for the most probable query

Setting A moves us from $32 \rightarrow 8$ states;

result: $\{C=no, S=fe, T1=-ve, T2=-ve\} = 47\%$

This is not applicable in every situation; we generalize into the **Maximum a Posteriori Hypothesis**:

this is more complex and less efficient, but more often applicable

we find a subset of variables & find the MPE

(ex. $S=C \mid A \rightarrow \{C=no, S=fe\} \approx 49.3\%$)

When we seek to make inferences, algorithms fall into two major categories:

1. variable elimination
2. conditioning

In both situations, complexity is tied to the topology of the Bayesian Network

the actual property is **tree width** — it is roughly analogous to connectivity

$MPH = O(nd^w)$ given $n=\#var$, $d=\#val$, $w=width$

We will not define tree width, as it is complex, but we observe a few special cases:

Trees have width 1 (one path from any node to node)

Poly-trees (>1 parent ok) have width = the maximum parent count of any node

These are both singly connected networks; multiply connected leads to a DAG

5

Our first method of performing inference is

0.1 Weighted Model Counting

Consider the statement $\Delta = (A \vee B) \wedge \neg C$

This has 3 variables and suggests 8 worlds

Given an nd-DNNF circuit, we can solve in $O(N)$!

Consider the example on the right; it is trivial! $WMC = 0.04 + 0.10 + 0.00 = 0.14$

We therefore only need a compiler which would transform $\Delta \rightarrow \text{sd-DNNF}$

Unfortunately, in the general case this is intractable

There are tractable subsets of this, though:

7

We can use probability as our weights! $\rightarrow WMC(\Delta \wedge \alpha) = Pr(\alpha)$

So we need to convert $\Delta \rightarrow$ boolean circuit.

We can see that Δ holds in $w_1, w_4, \& w_7$

We thus say $W(A) = W(\neg A) = \dots = W(\neg C) = 1$

We then let $W(P_i) = \theta_i \& W(\neg P_i) = 0$, so $W(A) = W(P_1)W(P_4)W(P_7)$

Thus we can solve probabilistic reasoning by symbol manipulations!

Properties of the algorithm:

1. there are many possible representations
2. it is not sensitive to tree width
3. it is not only applicable to Bayesian Networks

We can see that the size(Bayes) = size(CPT); though size(Bayes) = $O(nd^{k+1})$, size(joint-table) = $O(D^n)$!

This shows that Bayesian Networks are much more space efficient!

The process of modeling logic as a Bayesian Network has 3 steps:

1. define variables & values
2. define edges
3. specify CPT

Variables will then be labeled either query or evidence variables depending on the query.

This is a common approach used in early spam filters and Google ad Rephil.

attachments.dropbox.com/s_5F1FEB98332EEC99C328935F7AB0E929534BB33EF2AB134DF14A85072ADC65A01590256640
drawing + 8.jpg)

Consider the following statements

- The cold causes a sore throat/chill
- The flu causes a sore throat/chill/fever/body ache
- Tonsillitis can show itself in fever/body ache

This network forms a bipartite graph.

We could have used one multivariable disease node.

We use this to give probabilities of a given condition.

If we have complete information, we can model a system as a Bayesian Network.

For incomplete information, however, we must use Expectation Maximization.

For example, suppose the following environment:

attachments.dropbox.com/s_5F1FEB98332EEC99C328935F7AB0E929534BB33EF2AB134DF14A85072ADC65A01590257642drawing + 9.jpg)

What is the marginal $(P|\neg S, \neg B, \neg U)$? — 10.21%!

WHAT?? why is that so high?

It turns out this is because of the false negative rate of the scanning test.

We want a false negative rate of below 5%.

We can address this in one of three ways:

1. get a better scanning test — a false (-) for S of 4.63% meets our requirement
2. lower the success of the procedure — 75.59% would meet our requirement
3. increase $P(L|P)$ — 99.67% meets our requirement

(b) and (c) turn out to be either impractical or in-economical, so our standard approach is to pay!