

# COMP90051 Statistical Machine Learning

Semester 2, 2015

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## 8. PGM Probabilistic Inference



THE UNIVERSITY OF  
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# Probabilistic inference on PGMs

*Computing marginal and conditional distributions from the joint of a PGM using Bayes rule and marginalisation.*

*This deck: how to do it efficiently.*

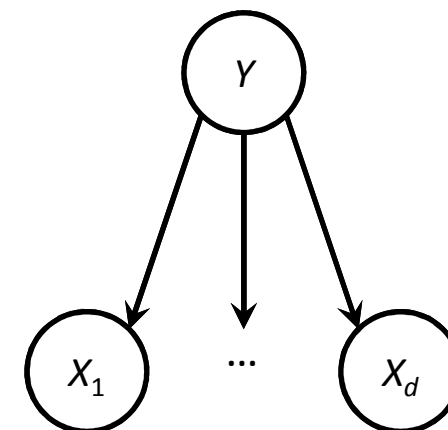
*Based on Andrew Moore's tutorial slides*

# Two familiar examples

- Naïve Bayes (frequentist/Bayesian)

- \* Chooses most likely class given data

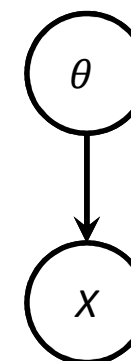
- \*  $\Pr(Y|X_1, \dots, X_d) = \frac{\Pr(Y, X_1, \dots, X_d)}{\Pr(X_1, \dots, X_d)} = \frac{\Pr(Y, X_1, \dots, X_d)}{\sum_y \Pr(Y=y, X_1, \dots, X_d)}$



- Data  $X|\theta \sim N(\theta, 1)$  with prior  $\theta \sim N(0,1)$  (Bayesian)

- \* Given observation  $X = x$  update posterior

- \*  $\Pr(\theta|X) = \frac{\Pr(\theta, X)}{\Pr(X)} = \frac{\Pr(\theta, X)}{\sum_{\theta} \Pr(\theta, X)}$

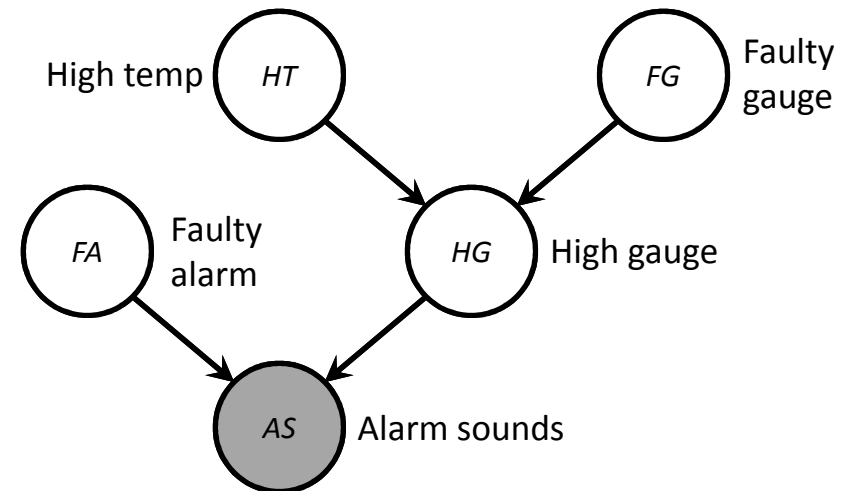


- Joint + Bayes rule + marginalisation  $\rightarrow$  anything

# Nuclear power plant

- **Alarm sounds; meltdown?!**

$$\begin{aligned} \Pr(HT|AS = t) &= \frac{\Pr(HT, AS=t)}{\Pr(AS=t)} \\ &= \frac{\sum_{FG, HG, FA} \Pr(AS=t, FA, HG, FG, HT)}{\sum_{FG, HG, FA, HC} \Pr(AS=t, FA, HR, FG, HT)} \end{aligned}$$



- Numerator (denominator similar)

expanding out sums, joint *summing once over  $2^5$  table*

$$= \sum_{FG} \sum_{HG} \sum_{FA} \Pr(HT) \Pr(HG|HT, FG) \Pr(FG) \Pr(AS = t|FA, HG) \Pr(FA)$$

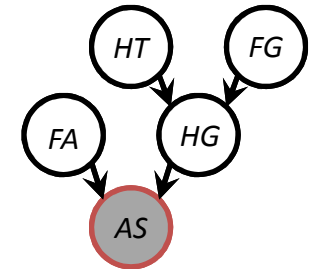
distributing the sums as far down as possible *summing over several smaller tables*

$$= \Pr(HT) \sum_{FG} \Pr(FG) \sum_{HG} \Pr(HG|HT, FG) \sum_{FA} \Pr(FA) \sum_{AS} \Pr(AS = t|FA, HG)$$

# Nuclear power plant (cont.)

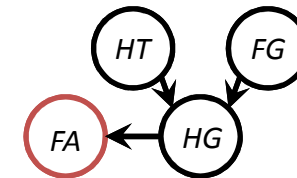
$$= \Pr(HT) \sum_{FG} \Pr(FG) \sum_{HG} \Pr(HG|HT, FG) \sum_{FA} \Pr(FA) \sum_{AS} \Pr(AS = t|FA, HG)$$

**eliminate AS:** since AS observed, really a no-op



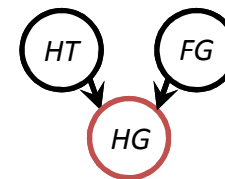
$$= \Pr(HT) \sum_{FG} \Pr(FG) \sum_{HG} \Pr(HG|HT, FG) \sum_{FA} \Pr(FA) m_{AS}(FA, HG)$$

**eliminate FA:** multiplying 1x2 by 2x2



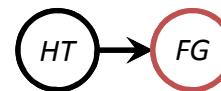
$$= \Pr(HT) \sum_{FG} \Pr(FG) \sum_{HG} \Pr(HG|HT, FG) m_{FA}(HG)$$

**eliminate HG:** multiplying 2x2x2 by 2x1



$$= \Pr(HT) \sum_{FG} \Pr(FG) m_{HG}(HT, FG)$$

**eliminate FG:** multiplying 1x2 by 2x2



$$= \Pr(HT) m_{FG}(HT)$$



Multiplication of tables, followed by summing, is actually matrix multiplication

$$m_{FA}(HG) = \begin{array}{c|c} & \begin{array}{c} FA \end{array} \\ \hline \begin{array}{c} f \\ t \end{array} & \begin{array}{cc} & \begin{array}{c} FA \end{array} \\ \hline f & \begin{array}{cc} 0.6 & 0.4 \end{array} \\ t & \begin{array}{cc} 0.6 & 0.4 \end{array} \end{array} \times \begin{array}{c|c} & \begin{array}{c} HG \end{array} \\ \hline \begin{array}{c} f \\ t \end{array} & \begin{array}{cc} & \begin{array}{c} HG \end{array} \\ \hline f & \begin{array}{cc} 1.0 & 0 \end{array} \\ t & \begin{array}{cc} 0.8 & 0.2 \end{array} \end{array}$$

# Elimination algorithm

Orange background  
= Slide just for fun!

**Eliminate** (Graph  $G$ , Evidence nodes  $E$ , Query nodes  $Q$ )

1. Choose node ordering  $I$  such that  $Q$  appears last
2. Initialise empty list **active**
3. For each node  $X_i$  in  $G$ 
  - a) Append  $\Pr(X_i | \text{parents}(X_i))$  to **active**
4. For each node  $X_i$  in  $E$ 
  - a) Append  $\delta(X_i, x_i)$  to **active**
5. For each  $i$  in  $I$ 
  - a) potentials = Remove tables referencing  $X_i$  from **active**
  - b)  $N_i$  = nodes other than  $X_i$  referenced by tables
  - c) Table  $\varphi_i(X_i, X_{N_i})$  = product of tables
  - d) Table  $m_i(X_{N_i}) = \sum_{X_i} \varphi_i(X_i, X_{N_i})$
  - e) Append  $m_i(X_{N_i})$  to **active**
6. Return  $\Pr(X_Q | X_E = x_E) = \varphi_Q(X_Q) / \sum_{x_Q} \varphi_Q(X_Q)$

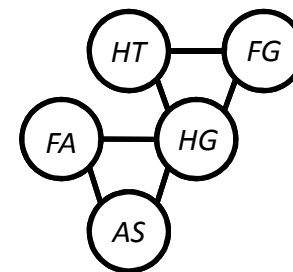
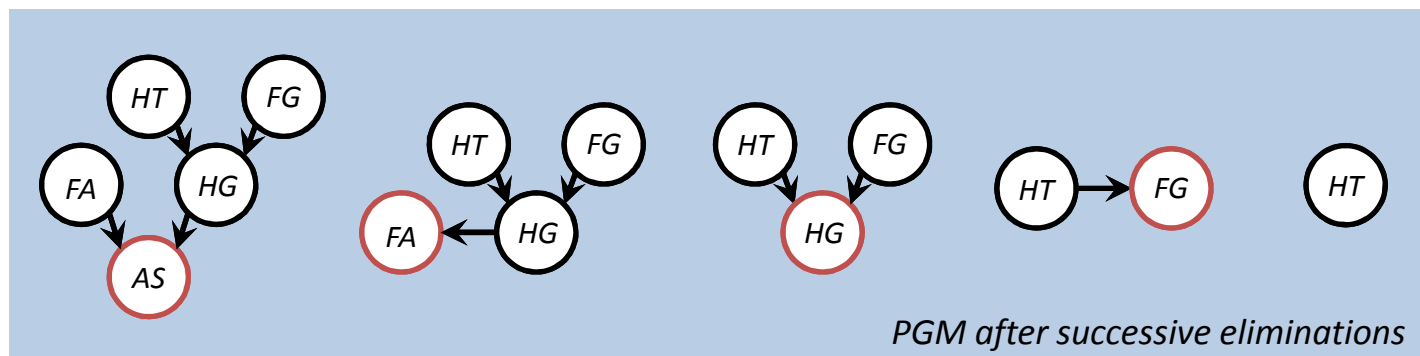
initialise

evidence

marginalise

normalise

# Runtime of elimination algorithm

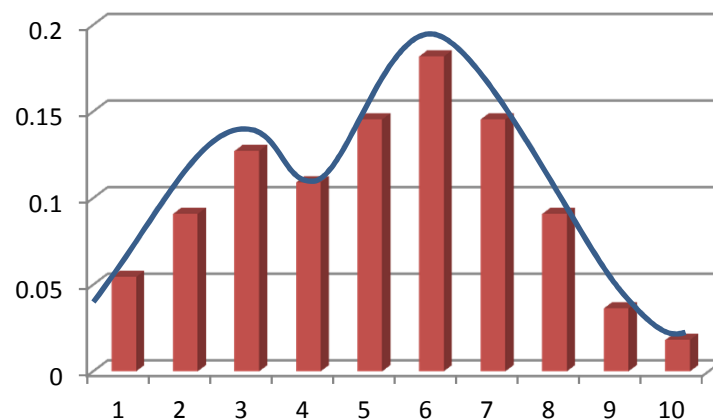


"reconstructed" graph  
From process called  
**moralisation**

- Each step of elimination
  - \* Removes a node
  - \* Connects node's remaining neighbours  
→ **forms a clique** in the "reconstructed" graph  
(cliques are exactly r.v.'s involved in each sum)
- Time complexity **exponential in largest clique**
- Different elimination orderings produce different cliques
  - \* **Treewidth**: minimum over orderings of the largest clique
  - \* Best possible time complexity is exponential in the treewidth

# Probabilistic inference by simulation

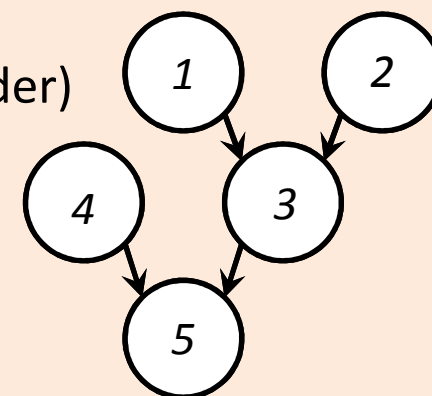
- Exact probabilistic inference can be expensive/impossible
- Can we approximate numerically?
- Idea: **sampling methods**
  - \* Cheaply sample from desired distribution
  - \* Approximate **distribution** by **histogram of samples**





# Monte Carlo approx probabilistic inference

- Algorithm: sample once from joint
  1. Order nodes' parents before children (topological order)
  2. Repeat
    - a) For each node  $X_i$ 
      - i. Index into  $\Pr(X_i | \text{parents}(X_i))$  with parents' values
      - ii. Sample  $X_i$  from this distribution
    - b) Together  $\mathbf{X} = (X_1, \dots, X_d)$  is a sample from the joint
- Algorithm: sampling from  $\Pr(X_Q | X_E = x_E)$ 
  1. Order nodes' parents before children
  2. Initialise set  $S$  empty; Repeat
    1. Sample  $\mathbf{X}$  from joint
    2. If  $X_E = x_E$  then add  $X_Q$  to  $S$
  3. Return: Histogram of  $S$ , normalising counts via divide by  $|S|$
- Sampling++: Importance weighting, Gibbs, Metropolis-Hastings



# Alternate forms of probabilistic inference

- Elimination algorithm produces single marginal
- **Sum-product** algorithm on trees
  - \* 2x cost, supplies all marginals
  - \* Name: Marginalisation is just **sum** of **product** of tables
  - \* “Identical” variants: **Max-product**, for MAP estimation
- In general these are **message-passing algorithms**
  - \* Can generalise beyond trees (beyond scope):  
junction tree algorithm, loopy belief propagation
- **Variational Bayes**: approximation via optimisation

# Summary

- Probabilistic inference on PGMs
  - \* What is it and why do we care?
  - \* Elimination algorithm; complexity via cliques
  - \* Monte Carlo approaches as alternate to exact integration