#### Stat 521A Lecture 4

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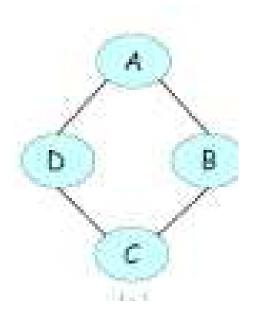
# Admin

• CS auditors: please turn in your form to Joyce Poon, who will pass it to Laks for signing

# Outline

- Aside on canonical parameterization (ex 4.4.14)
- Structured factors (4.4.1.2)
- Structured CPDs (5.2-5.6)
- Temporal models (6.2)

## Degrees of freedom of a UGM

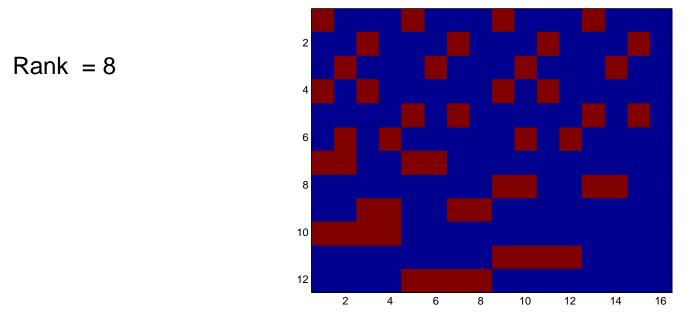


Why do we just need 8 numbers to uniquely parameterize the distribution?

Eg a^1, b^1, c^1, d^1, (a^1,b^1), (b^1,c^1), (c^1,d^1), (a^1,d^1)

#### Num params = rank of feature matrix

- Let F(n, i)=1 iff i'th bit vector turns on n'th feature
- Each feature specifies a value for every pair of nodes connected by an edge, and hence is a vector in R^{16}. 4 edges, 3 unique settings = 12 rows.



Eg a^1, b^1, c^1, d^1, (a^1,b^1), (b^1,c^1), (c^1,d^1), (a^1,d^1)

## Rank of feature matrix

- edges = {[1 2], [1 3], [2 4], [3 4]};
- ndx = 1;
- F = zeros(0, 2^4);
- for e=1:length(edges)
- s = edges{e}(1); t = edges{e}(2);
- for j=1:2
- for k=1:2
- if j==2 && k==2, continue; end
- for x=1:16
- xv= ind2subv([2 2 2 2], x);
- if xv(s)==j && xv(t)==k
- F(ndx,x)=1;
- end
- end
- ndx = ndx + 1;
- end
- end
- end
- rank(F)

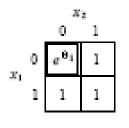


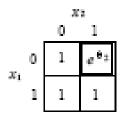
## Log-linear factors

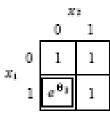
- A factor defined on m discrete rv's with K states needs K<sup>m</sup> parameters.
- Imagine a factor on triples of letters. Instead of having 26<sup>3</sup> numbers, we can define binary features that only turn on for certain values, eg f<sub>ing</sub>(x) = 1 iff x<sub>1</sub>='l',x<sub>2</sub>='n',x<sub>3</sub>='g'. This has weight ω<sub>ing</sub>. We define

$$\phi_c(\mathbf{x}_c) = \exp(\sum_{i=1}^k w_{c,i} f_{c,i}(\mathbf{x}_c))$$

#### Tables are a special case







0

1 - 1

 $x_1$ 

 $\frac{32}{0}$ 

1

1

1

 $e^{\theta_4}$ 

$$f_i(x_1, x_2) = \delta(x_1 = 1, x_2 = 0)$$

 $f_1(x_1, x_2) = \delta(x_1 = 0, x_2 = 0)$ 

 $f_2(x_1, x_2) = \delta(x_1 = 0, x_2 = 1)$ 

$$f_4(x_1, x_2) = \delta(x_1 = 1, x_2 = 1)$$

 $\varphi_{12}(x_1, x_2) = e^{\Theta_1 f_1 + \Theta_2 f_2 + \Theta_3 f_1 + \Theta_4 f_4}$ 

$$\begin{array}{c}
x_{2} \\
0 \\
x_{1} \\
1 \\
\end{array}$$

Jordan, fig 19.1

# **CRF** features

- Typical features used in a CRF model for language processing (X=words, Y=labels)
- $F_1(Y_t, X_t, X_{t-1}, X_t+1) = I(X_{t-1}="New", X_t="York", X_{t+1}="Times", Y_t="Object")$
- $F_2(Y_t, X_t, X_{t-1}, X_t+1) = I(X_{t-1}="New", X_t="York", X_{t+1} \neq "Times", Y_t="Place")$
- Models often have ~100k manually specified features.
- Common to use L1 regularization to sparsify.
- Can also perform feature induction, by eg greedily creating conjunctions or disjunctions

## Exponential family (maxent) models

• Combining all the local potentials

$$p(\mathbf{x}) = \frac{1}{Z} \prod_{c} \phi_{c}(\mathbf{x}_{c})$$

$$\phi_{c}(\mathbf{x}_{c}) = \exp(\sum_{i=1}^{k} w_{c,i} f_{c,i}(\mathbf{x}_{c}))$$

$$p(\mathbf{x}) = \frac{1}{Z} \exp(\sum_{i} w_{i} f_{i}(\mathbf{x}_{c_{i}}))$$

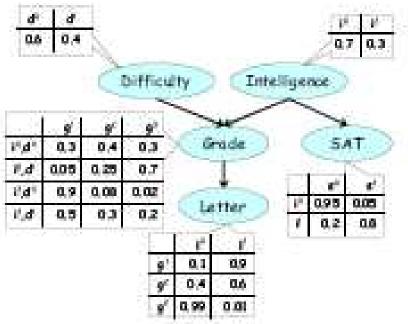
DAGs are a special case where each  $\phi_c(x_c) = p(X_i|Pa(X_i))$  sums to 1, so Z=1

See ch 8



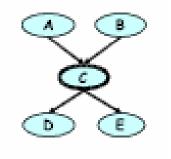
# Tabular CPDs

- If all nodes are discrete and have K values, we can represent p(X\_i|Pa(X\_i)) as a table, with one row per conditioning case (K^#pa), and K columns which sum to 1
- If K and/or #pa is large, this is too many parameters, so we seek more parsimonious representations.



# Deterministic CPDs

- In some cases, the child is a deterministic function of the parents, eg bloodtype is determined by the 2 alleles
- Deterministic nodes often denoted by doubleringed oval.
- Determinism can imply additional (non-graphical) independencies
- Eg  $\dot{D} \perp E \mid A,B$  since C = fn(A,B)



Det-sep

# Context specific independence (CSI)

- Sometimes, the set of edges which are "active" depends on the value of the nodes
- Eg Y is a noisy observation of object X1, or X2. Z specifies the identity of the measurement. Let X =multiplexer(X1,X2, Z). Then X2  $\perp$  Y | Z=1. So our

posterior on X2 is not affected by the measurement. (Data association ambiguity)

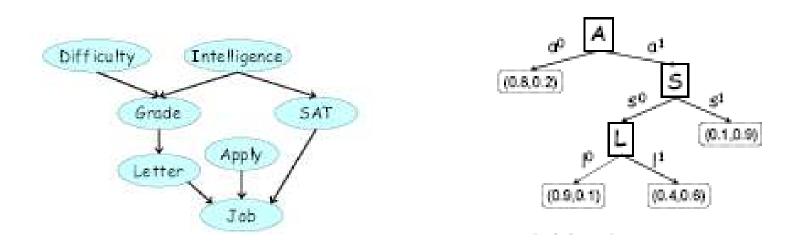
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# Contingently acyclic BNs

- Sometimes we can define a directed graph with cycles, but where some of the edges are not active for a given setting of certain variables C.
- If we can guarantee that the graph is a DAG for each context C=c, the result is a mixture of differently structured BNs.
- This is called a Bayesian multinet.

## **Tree-structured CPDs**

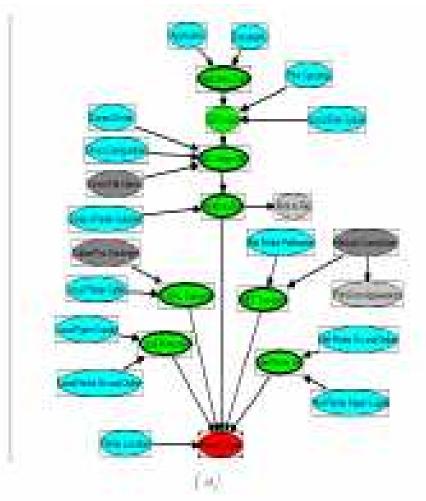
 Different parents can be rendered irrelevant, depending on the values
 P(J|A,S,L)

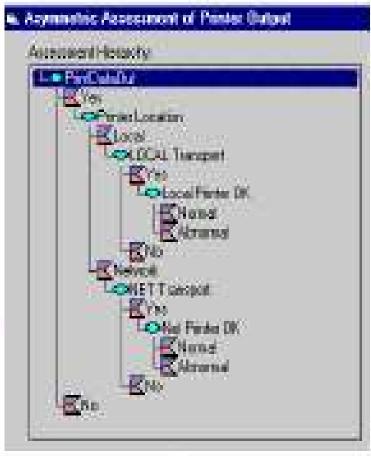


#### Eg. J | S,L if A=0 since we go down left branch of tree

# Printer fault diagnosis in MS windows

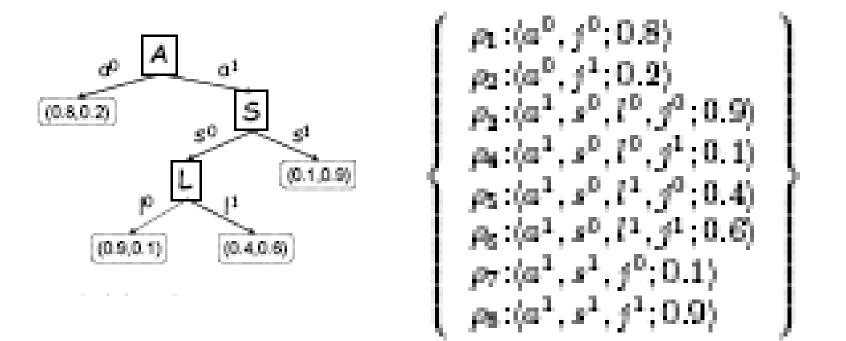
• Uses tree structured CPDs, since different sets of variables are relevant in different contexts





## **Rule-structured CPDs**

Specify a pattern and a value



# Logistic regression (sigmoid BNs)

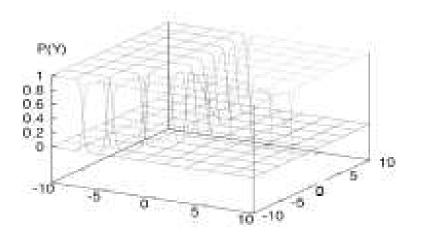
 Suppose all nodes are binary. We can use logreg CPDs

$$p(y = 1 | \mathbf{x}) = \sigma(w_0 + \sum_{i=1}^k w_i x_i) \quad \sigma(u) = \frac{1}{1 + e^{-u}}$$

# **Multinomial logreg**

 If Y is K-ary, and the parents are binary or cts, we can use a softmax function

$$p(y = j | \mathbf{x}) = \frac{\exp(\mathbf{w}_j^T \mathbf{x})}{\sum_{j'=1}^{K} \exp(\mathbf{w}_{j'}^T \mathbf{x})}$$



For K-ary parents, use 1-of-K encoding

## Independence of causal influence

 We can model the effects of many parents by assuming that each parent is corrupted by independent noise, and the results are deterministically combined via a simple function such as OR or MAX

 $\begin{array}{cccc} \chi_1 & & \chi_{l_{\mathcal{E}}} \\ \downarrow & & \downarrow \\ \chi_1 & & \downarrow \\ \chi_1 & & \chi_k \end{array}$ 

## Noisy-or model

- Each Xi in {0,1} gets passed through a noisy wire to produce Zi in {0,1}. 0 maps to 0, 1 maps to 0 wp w<sub>i</sub> (failure probability). λ<sub>i</sub>=1-w<sub>i</sub> is the prob. that Xi alone turns on Y.
- The Zi's are combined in an OR to produce Z. Then Y=Z.
- The only way Y can be off is if all Zi's are off, which means all the wires for Xi st Xi=1 independently failed:

$$p(y = 0 | \mathbf{x}) = \prod_{i:x_i=1} w_i = \prod_{i=1} w_i^{x_i}$$
$$p(y = 1 | \mathbf{x}) = 1 - p(y = 0 | \mathbf{x})$$

Popular in cogsci models of causality

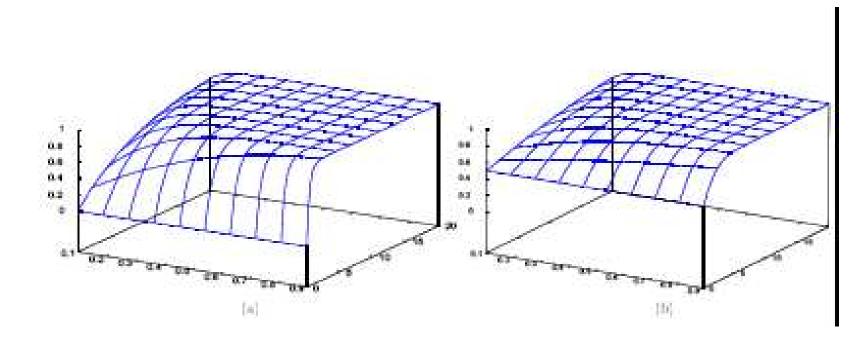
# Example

- P(fever=0|cold=1, flu=0, malaria=0)=0.6
- P(fever=0|cold=0, flu=1, malaria=0)=0.2
- P(fever=0|cold=0, flu=0, malaria=1)=0.1

| Cold | Flu | Malaria | p(Fever=1) | p(Fever=0)                  |
|------|-----|---------|------------|-----------------------------|
| 0    | 0   | 0       | 0.0        | 1.0                         |
| 0    | 0   | 1       | 0.0        | 0.1                         |
| 0    | 1   | 0       | 0.8        | 0.2                         |
| 0    | 1   | 1       | 0.98       | 0.02=0.2	imes 0.1           |
| 1    | 0   | 0       | 0.4        | 0.6                         |
| 1    | 0   | 1       | 0.94       | 0.0.6=0.6	imes 0.1          |
| 1    | 1   | 0       | 0.88       | 0.12=0.6	imes 0.2           |
| 1    | 1   | 1       | 0.988      | 0.012=0.6	imes 0.2	imes 0.1 |

## Leak nodes

 If Y=0 and all Xi=0, the CPD assigns 0 probability to this event. To prevent this, we add a leak node, X0=1, which is always on, to model "any other cause". The leak can fail wp w0.



W0=1

W0=0.5

## **BN20** networks

- In medical diagnosis, it is common to construct 2 layered bipartite networks of binary nodes, mapping diseases to symptoms (findings).
- Because of the large number of parents, the child nodes use noisy-or.
- Conditional on F, the diseases D are correlated.
- The QMR-DT network is a standard testbed for evaluating approximate inference algorithms.

(10

$$F_1 \quad F_2 \quad F_3 \quad \dots \quad F_n$$

# Negative findings

- If Fi=1, the disease parents fight to explain the finding. Hence they become fully correlated.
- But if Fi=0, the parents are independent! Hence the p(Fi=0|Pa(Fi)) likelihood fully factorizes, and does not make inference harder (homework).

 $D_1 \perp D_2 \mid F_i = 0$ 

## Conditional linear Gaussian CPDs

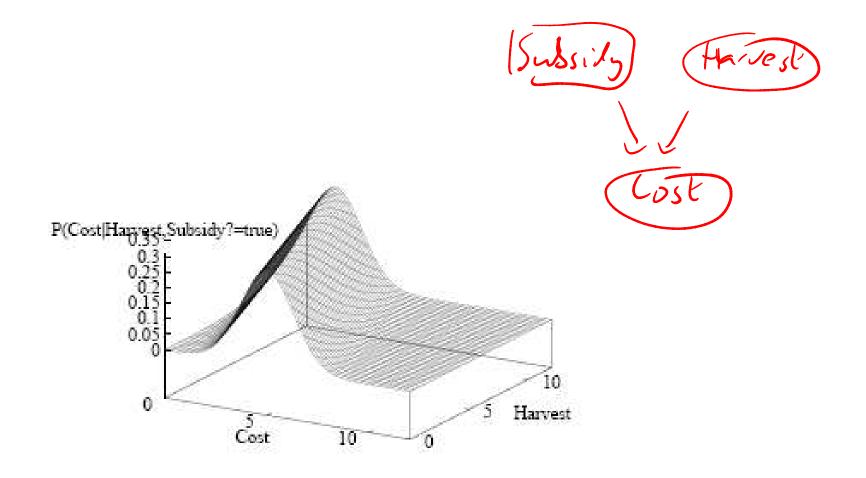
• If Y is continuous and all the parents are cts we can define

$$p(y|\mathbf{x}) = \mathcal{N}(y|\mathbf{x}^T\mathbf{w}, \sigma^2)$$

- Networks of linear Gaussian CPDs define a joint multivariate Gaussian (see ch 7)
- For discrete parents u, we can use 1-of-K and LG, or we can use a different set of parameters for each discrete setting (CLG). The resulting distribution is a mixture of Gaussians, where each discrete setting defines a mixture component.

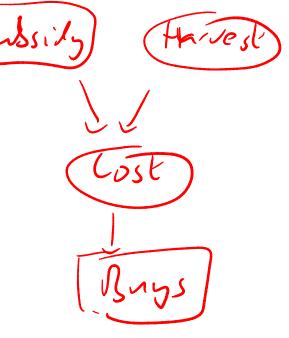
$$p(y|\mathbf{x}, \mathbf{u} = k) = \mathcal{N}(y|\mathbf{x}^T \mathbf{w}_k, \sigma_k^2)$$

# Example of CLG network



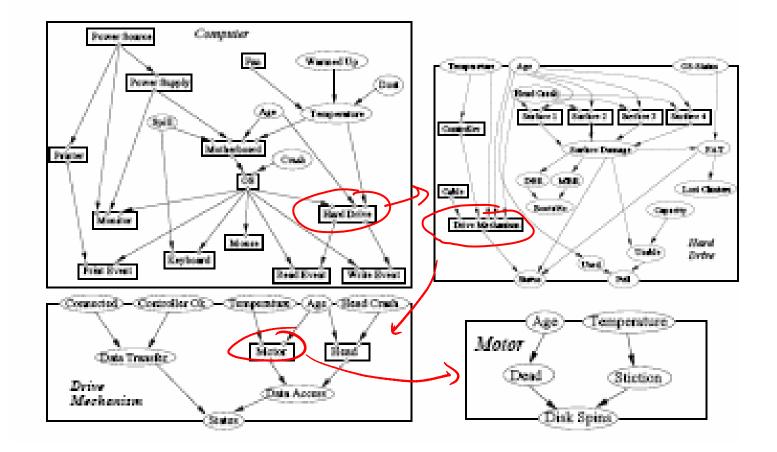
# Hybrid network

P(buys=1|cost) = logreg or probit. Joint distribution is no longer mixture of Gaussians. Closed-form inference no longer possible (see ch14).



## **Encapsulated BNs**

- We can embed a BN inside a CPD, and "hide" the internal nodes using an interface layer.
- This, combined with parameter tying, yields OOBN.





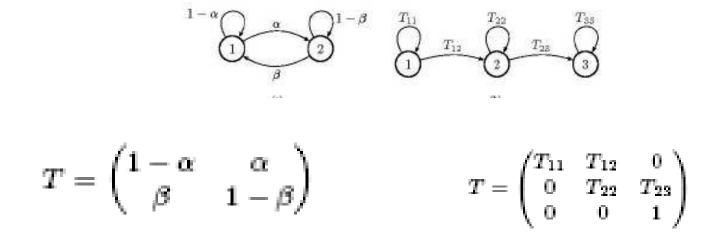
## Markov chains

 We can define a distribution over a semi-infinite sequence X\_1, X\_2, ... by using a discrete-time Markov chain with tied parameters (stationary)

T

#### State transition diagram

Picture of the stochastic finite state automaton

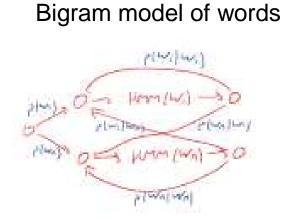


## Hidden Markov Models

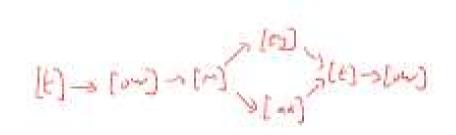
- An HMM is a function of a Markov chain.
- We observe  $V_t$ , hidden state is  $H_t$  in  $\{1, ..., K\}$
- $P(H_t=j|H_{t-1}=i)$  is the transition model
- P(V<sub>t</sub>|H<sub>t</sub>=j) is the observation model (eg mixture of Gaussians)

$$\begin{array}{cccc}
\mu_{1} \rightarrow \mu_{2} \rightarrow \cdots \\
\downarrow & \downarrow \\
\nu_{1} & \nu_{2}
\end{array}$$

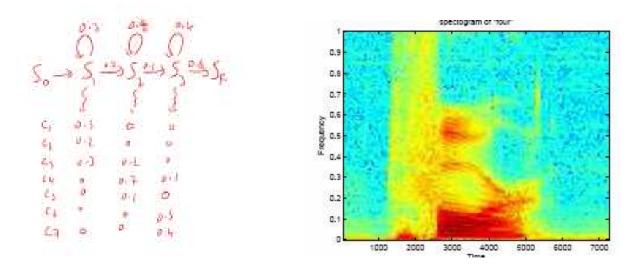
#### HMMs for speech recognition



Pronunciation model : word -> phonemes



Acoustic model: phonemes -> observations



## State space models

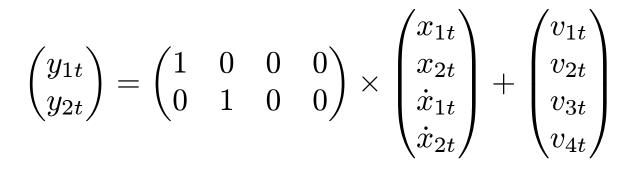
- Same graph (CI assumptions) as HMM, but now X and Y are real-valued vectors
- Special case: linear dynamical system (LDS)

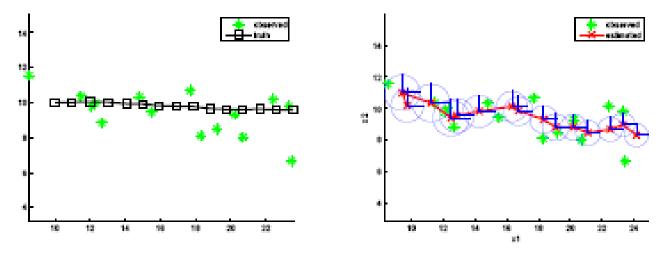
$$p(\mathbf{x}_t | \mathbf{x}_{t-1}) = \mathcal{N}(\mathbf{x}_t | \mathbf{A}\mathbf{x}_{t-1}, \mathbf{Q})$$
$$p(\mathbf{y}_t | \mathbf{x}_t) = \mathcal{N}(\mathbf{y}_t | \mathbf{H}\mathbf{x}_t, \mathbf{R})$$

$$egin{array}{rcl} \mathbf{x}_t &=& \mathbf{A}\mathbf{x}_{t-1} + \mathcal{N}(\mathbf{0},\mathbf{Q}) \ \mathbf{y}_t &=& \mathbf{H}\mathbf{x}_t + \mathcal{N}(\mathbf{0},\mathbf{R}) \end{array}$$

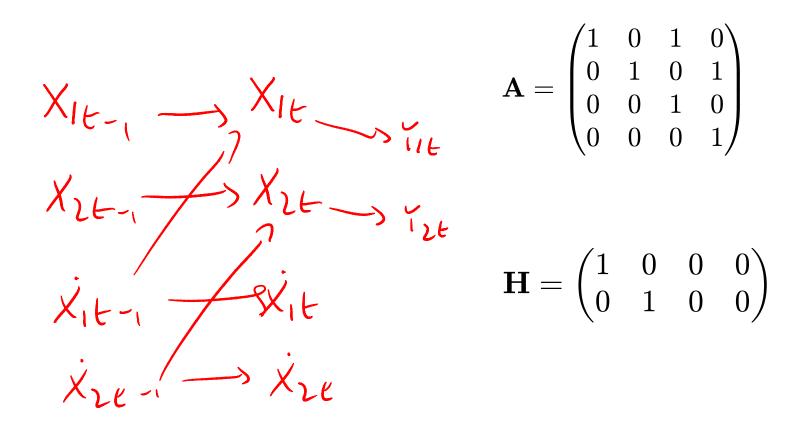
#### Example: tracking in 2D

$$\begin{pmatrix} x_{1t} \\ x_{2t} \\ \dot{x}_{1t} \\ \dot{x}_{2t} \end{pmatrix} = \begin{pmatrix} 1 & 0 & 1 & 0 \\ 0 & 1 & 0 & 1 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix} \times \begin{pmatrix} x_{1t-1} \\ x_{2t-1} \\ \dot{x}_{1t-1} \\ \dot{x}_{2t-1} \end{pmatrix} + \begin{pmatrix} w_{1t} \\ w_{2t} \\ w_{3t} \\ w_{4t} \end{pmatrix}$$



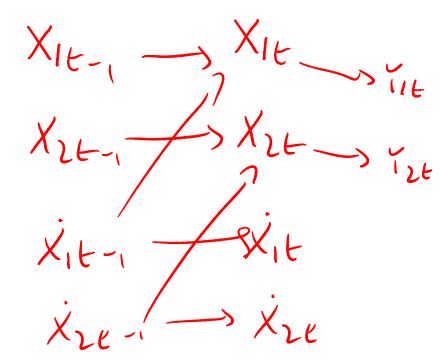


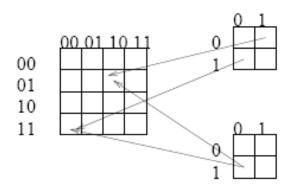
#### LDS as DGM



For linear Gaussian systems, sparse matrices = sparse graphs

#### **Dynamic Bayes Nets**





P(X1(t),X2(t) | X1(t-1),X2(t-1)

If the variables are discrete, the transition matrix of the compound model (all 4 variables) is not sparse or structured. So the graph structure is crucial.