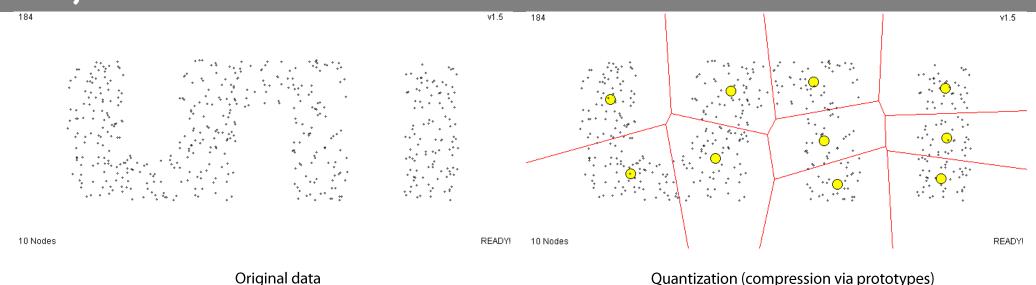
# Artificial Intelligence

#### Unsupervised Learning

Marco Piastra

An aside: The K-means algorithm (i.e. alternate optimization)

#### Vector quantization



The basic idea is to replace each real-valued vector  $x \in \mathbb{R}^d$  with a discrete symbol  $w_j \in \mathbb{R}^d$  which belongs to a codebook of k prototypes  $\theta := \{w_1, \dots, w_k\}$ 

Each data vector is encoded by using the index of the most similar prototype, where similarity is measured in terms, for instance, of Euclidean distance:

$$w(\boldsymbol{x}) := \operatorname{argmin}_{\boldsymbol{w}_j} \|\boldsymbol{x} - \boldsymbol{w}_j\|$$

For instance, part of mpeg4 and QuickTime (Apple) video compression algorithms work in this way and so does the Ogg Vorbis audio compression algorithm

Given a set  $D:=\{x_1,\ldots,x_N\}$  of observations (i.e. vectors in $\mathbb{R}^d$ ) and a set  $\theta:=\{w_1,\ldots,w_k\}$  of k prototypes (i.e. vectors in  $\mathbb{R}^d$ )

Clustering problem: find an assignment function  $w: D \to \theta$  such that the objective (loss) function:

$$J(D, \theta) := \frac{1}{2} \sum_{i=1}^{N} \| \boldsymbol{x}_i - w(\boldsymbol{x}_i) \|^2$$

is minimized.

#### *k-means* algorithm:

- 1) Position the k prototypes at random
- 2) Assign each observation to its closest prototype

$$w(\boldsymbol{x}_i) := \operatorname{argmin}_{\boldsymbol{w}_i} \|\boldsymbol{x}_i - \boldsymbol{w}_j\|$$

3) Position each prototype at the *centroid* of the observations assigned to it

$$m{w}_j = rac{1}{|D(m{w}_j)|} \sum_{D(m{w}_j)} m{x}_i \qquad ext{where} \ \ D(m{w}_j) := \{m{x}_i \in D \mid w(m{x}_i) = m{w}_j\}$$

4) Unless no prototype was moved in step 3), go back to step 2)

This algorithm converges to a <u>local</u> minimum of  $J(D, \theta)$ 

Why does the algorithm work: alternate optimization (also 'coordinate descent')

Step 2): Assign observations while keeping the k prototype fixed

$$w(\boldsymbol{x}_i) := \operatorname{argmin}_{\boldsymbol{w}_i} \|\boldsymbol{x}_i - \boldsymbol{w}_j\|$$

which minimizes each of the terms in  $J(D,\theta) := \frac{1}{2} \sum_{i=1}^N \| m{x}_i - w(m{x}_i) \|^2$ 

Step 3): Reposition the k prototypes while keeping the assignments fixed

$$J(D, \theta) := \frac{1}{2} \sum_{i=1}^{N} \|\boldsymbol{x}_i - w(\boldsymbol{x}_i)\|^2 = \frac{1}{2} \sum_{j} \sum_{D(\boldsymbol{w}_j)} (\boldsymbol{x}_i - \boldsymbol{w}_j)^2$$

$$\frac{\partial}{\partial \boldsymbol{w}_{j}} J(D, \theta) = \frac{\partial}{\partial \boldsymbol{w}_{j}} \frac{1}{2} \sum_{D(\boldsymbol{w}_{j})} (\boldsymbol{x}_{i} - \boldsymbol{w}_{j})^{2} = \frac{\partial}{\partial \boldsymbol{w}_{j}} \frac{1}{2} \sum_{D(\boldsymbol{w}_{j})} (\boldsymbol{x}_{i} - \boldsymbol{w}_{j})^{T} (\boldsymbol{x}_{i} - \boldsymbol{w}_{j})$$

$$= \frac{\partial}{\partial \boldsymbol{w}_{j}} \frac{1}{2} \sum_{D(\boldsymbol{w}_{j})} (\boldsymbol{x}_{i}^{2} + \boldsymbol{w}_{j}^{2} - 2 \boldsymbol{x}_{i}^{T} \boldsymbol{w}_{j}) = \sum_{D(\boldsymbol{w}_{j})} (\boldsymbol{w}_{j} - \boldsymbol{x}_{i})$$

then, by imposing  $\frac{\partial}{\partial w_i}J(D,\theta)=0$  we obtain

$$oldsymbol{w}_j = rac{1}{|D(oldsymbol{w}_j)|} \sum_{D(oldsymbol{w}_j)} oldsymbol{x}_i$$

#### Discussion of the *k-means algorithm*

- a) At each step of the algorithm J(D, heta) cannot increase: only decrease or stay equal
- b) The algorithm is a variant of a *gradient descent*, in which at each step the *gradient descent* is performed on one subset of variables only
- c) It must reach a *fixed point*, where both gradients vanish
- d) But the only guarantee is that the algorithm reaches a local minimum (unless it gets stuck in a saddle point)

Given a set  $D := \{x_1, \dots, x_N\}$  of observations (i.e. vectors in  $\mathbb{R}^d$ ) and a set  $\theta := \{w_1, \dots, w_k\}$  of k prototypes (i.e. vectors in  $\mathbb{R}^d$ )

#### Voronoi cell:

$$V(oldsymbol{w}_i) := \{oldsymbol{x} \in \mathbb{R}^d \mid \|oldsymbol{x} - oldsymbol{w}_i\| \leq \|oldsymbol{x} - oldsymbol{w}_l\| \ , orall \ l 
eq j \}$$

**Voronoi tesselation**: the complex of all Voronoi cells of  $\theta$ 

#### **Algorithm** (rewritten):

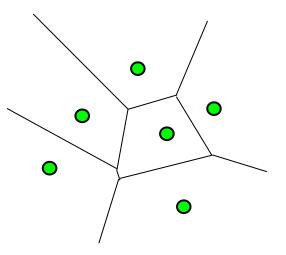
- 1) Position the k prototypes at random
- 2) Assign each observation to its Voronoi cell

$$w(\boldsymbol{x}_i) := \boldsymbol{w}_j \mid \boldsymbol{x}_i \in V(\boldsymbol{w}_j)$$



$$oldsymbol{w}_j = rac{1}{|\{oldsymbol{x}_i \in V(oldsymbol{w}_j)\}|} \sum_{\{oldsymbol{x}_i \in V(oldsymbol{w}_j)\}} oldsymbol{x}_i$$

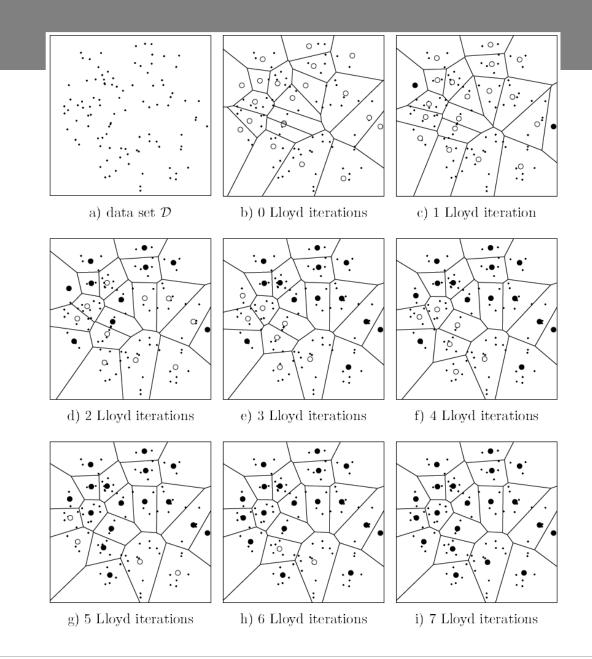
4) Unless no prototype was moved in step 3), go back to step 2)



#### k-means

*An example run of the algorithm* 

The landmarks (empty circles) become black when they cease to move



Artificial Intelligence 2022-2023

# The Expectation-Maximization (EM) algorithm

# Expected value of a random variable

(also expectation)

**Basic definition** 

$$\mathbb{E}_X[X] := \sum_{x \in \mathcal{X}} x \ P(X = x)$$

A linear operator

$$\mathbb{E}[X+Y] = \mathbb{E}[X] + \mathbb{E}[Y]$$
$$\mathbb{E}[cX] = c\mathbb{E}[X]$$

More concise notation

$$\mathbb{E}[X] := \sum_{x \in \mathcal{X}} x \ P(x)$$

Continuous case

$$\mathbb{E}[X] := \int_{x \in \mathcal{X}} x \ p(x) dx$$

Conditional expectation

$$\mathbb{E}_X[X|Y=y] = \mathbb{E}[X|Y=y] := \sum_{x \in \mathcal{X}} x \ P(X=x|Y=y)$$

Iterated expectation (see Wikipedia)

$$\mathbb{E}_X[X] = \mathbb{E}_Y[\mathbb{E}_X[X|Y]]$$

## Joint Expected Value

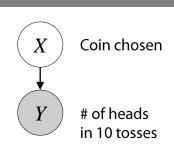
The **expected value** of a function f of a <u>set</u> of random variables is  $\{X_i\}$ 

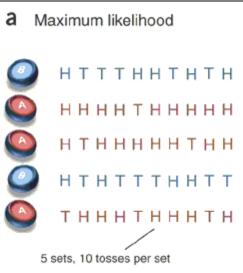
$$\mathbb{E}[f(\{X_i\})] := \sum_{\{X_i\}} f(\{X_i\}) \ P(\{X_i\})$$

the sum is over all possible combinations of values of the random variables

(Unless specified otherwise, the  $\mathbb{E}$  operator acts over *all* the random variables enclosed)

The extension to the continuous case is obvious





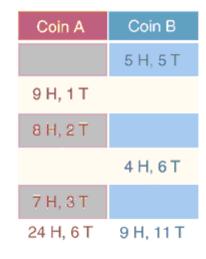


Figure from http://www.nature.com/nbt/journal/v26/n8/full/nbt1406.html

#### $\hat{\theta}_B = \frac{9}{9 + 11} = 0.45$

 $\hat{\theta}_A = \frac{24}{24 + 6} = 0.80$ 

#### An experiment with two coins

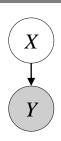
At each step, one coin is selected at random (with equal probability) and then tossed ten times

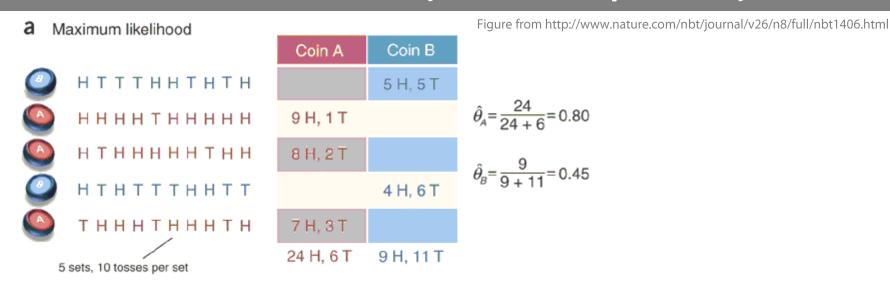
Random variables: Y number of heads, X selected coin (i.e A or B)

Parameters to be learnt:  $\theta = \{\theta_A, \theta_B\}$  probabilities of landing on head of A and B

When the results are fully observable, by MLE:

$$\theta_A^* = \frac{N_{Y=1,X=A}}{N_{X=A}}$$
  $\theta_B^* = \frac{N_{Y=1,X=B}}{N_{X=B}}$ 





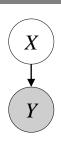
An experiment with two coins

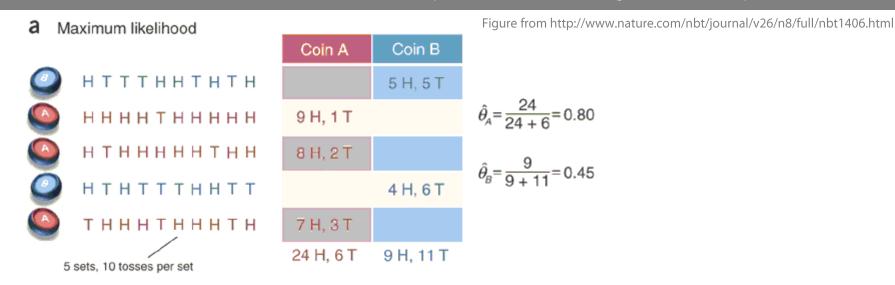
At each step, one coin is selected at random (with equal probability) and then tossed ten times

Random variables: Y number of heads, X selected coin (i.e A or B)

Parameters to be learnt:  $\theta = \{\theta_A, \ \theta_B\}$  probabilities of landing on head of A and B

• What if X is hidden (= latent, = unobserved)? The results of each sequence of coin tosses are known, but not the coin selected



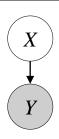


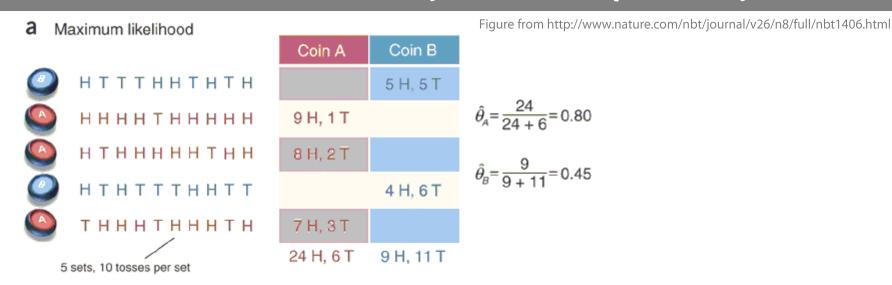
What if X is hidden (= latent, = unobserved)?
Likelihood

$$P(D \mid \theta) = P(\{Y^{(i)}\} \mid \theta) = \sum_{\{X^{(i)}\}} P(\{(Y^{(i)}, X^{(i))})\} \mid \theta)$$

MLE

$$heta^* := ext{argmax}_{ heta} \ \sum_{\{X^{(i)}\}} P(\{(Y^{(i)}, X^{(i)})\} \mid heta)$$
\* This optimization is intractable, in general





- What if X is hidden (= latent, = unobserved)?
  Intuitive idea: use expected values for unobserved variables
  - 1. Define an initial (random) guess  $\hat{\theta}^{(0)}$

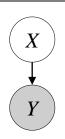
2. Compute 
$$Q_i(X^{(i)}) := P(X^{(i)} \mid Y^{(i)}; \hat{\theta}^{(t)})$$

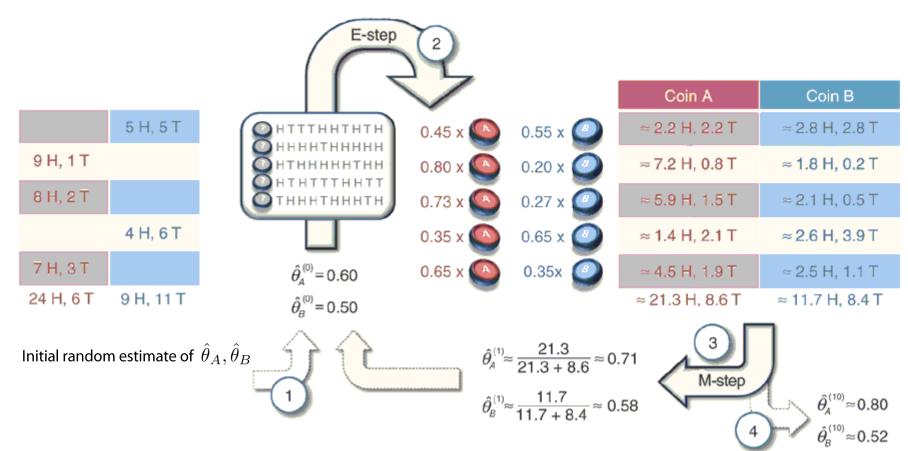
Maximize

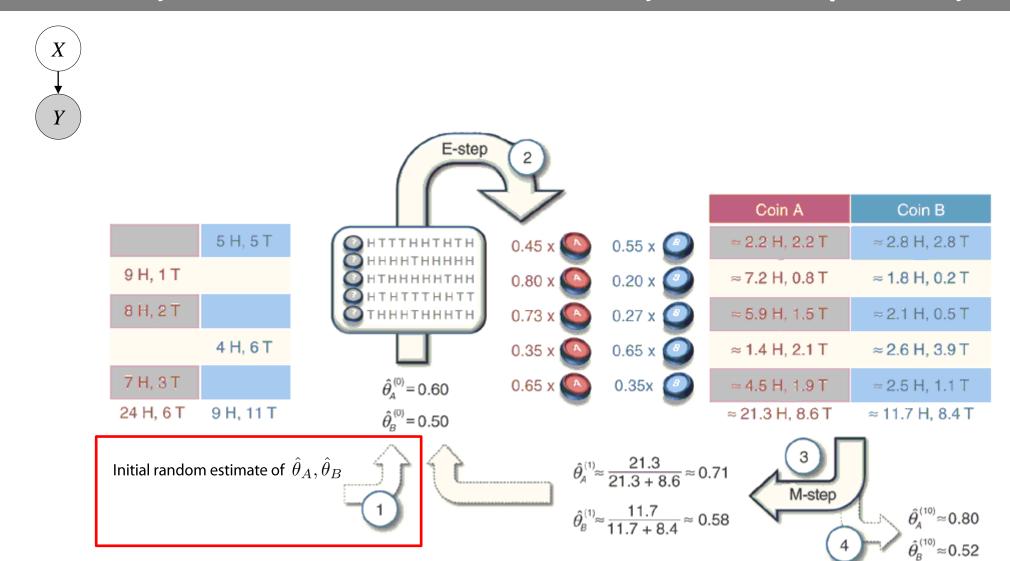
$$\hat{ heta}^{(t+1)} = \mathrm{argmax}_{ heta} \ \sum_i \mathbb{E}_{Q_i(X^{(i)})}[Y^{(i)} \mid X^{(i)}; \hat{ heta}^{(t)}]$$
 M-step

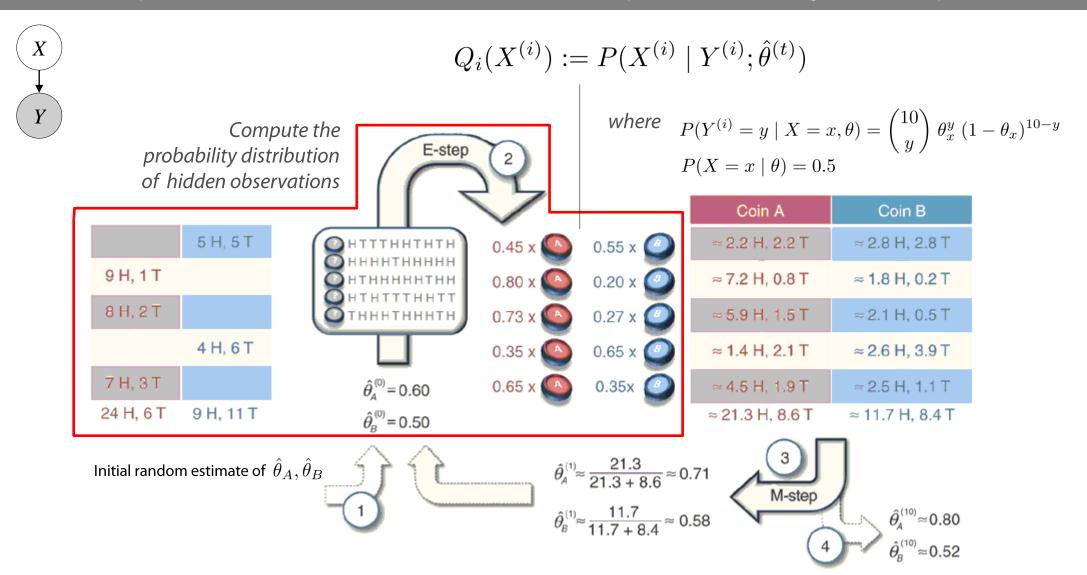
4. Unless some convergence criterion has been met, go to step 2.

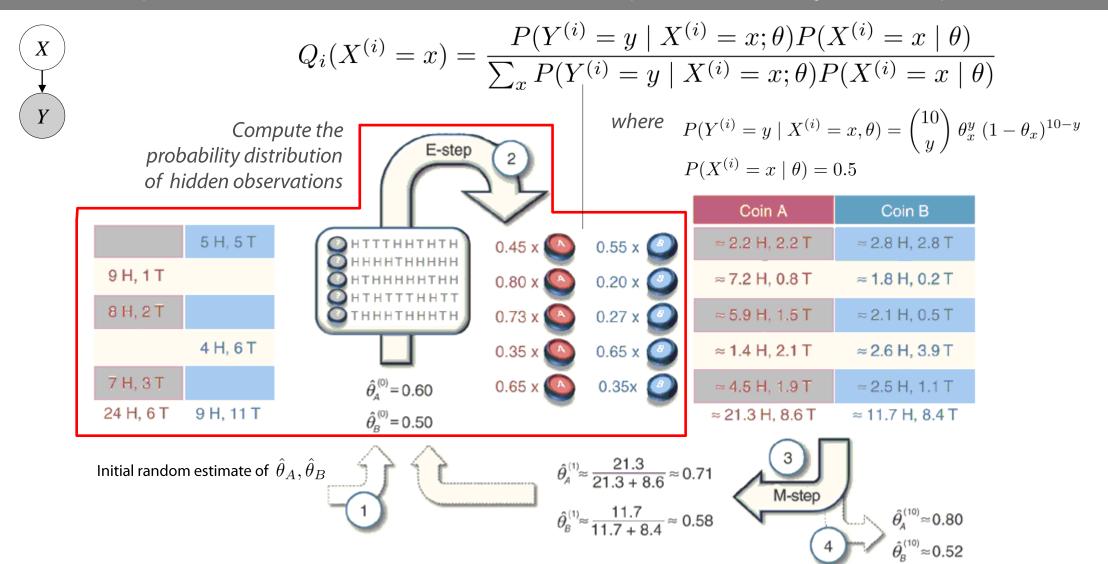
E-step











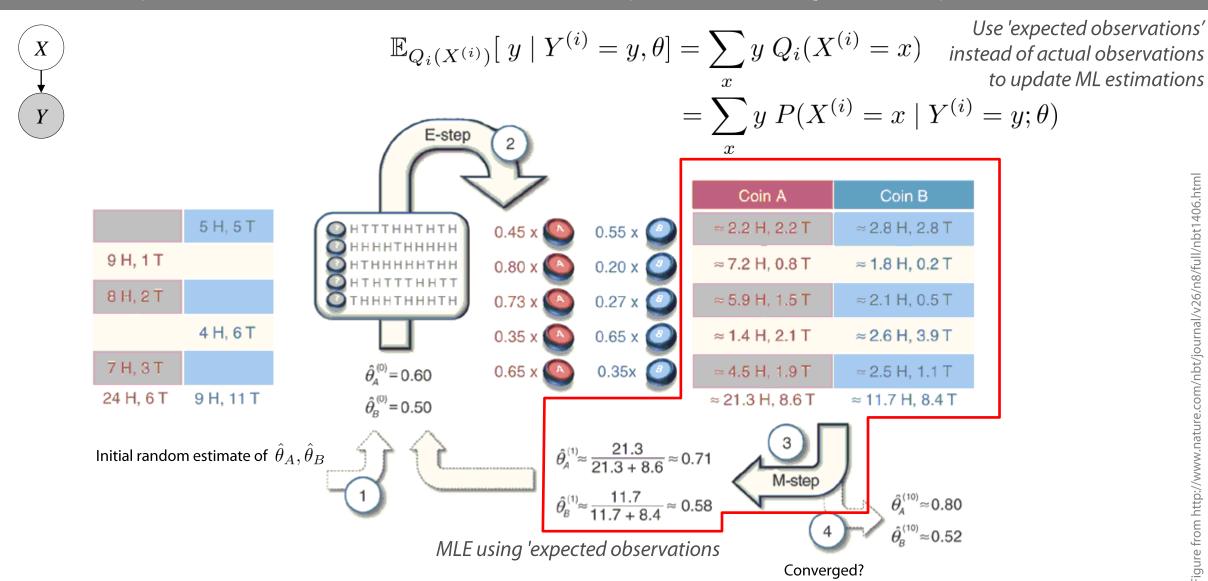


Figure from http://www.nature.com/nbt/journal/v26/n8/full/nbt1406.html

# Convergence of the EM algorithm (in the discrete case)

## An aside: Jensen's inequality

A relationship between probability and geometry

When *f* is convex function

$$f(\mathbb{E}[\{X_i\}]) \leq \mathbb{E}[f(\{X_i\})]$$

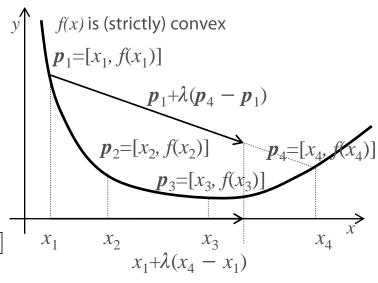
f is **convex** when for any two points  $p_i$  and  $p_j$  the segment  $(p_i - p_j)$  is not below f

That is, when

$$\lambda f(x_i) + (1 - \lambda)f(x_j) \ge f(\lambda x_i + (1 - \lambda)x_j), \, \forall \lambda \in [0, 1]$$

Furthermore, f is **strictly convex** when

$$\lambda f(x_i) + (1 - \lambda)f(x_j) \ge f(\lambda x_i + (1 - \lambda)x_j), \, \forall \lambda \in (0, 1)$$



#### Corollary:

when f is *strictly* convex, if and only if all the variables in  $\{X_i\}$  are <u>constant</u> it is true that

$$f(\mathbb{E}[\{X_i\}]) \leq \mathbb{E}[f(\{X_i\})]$$

Dual results also hold for *concave* functions

# An aside: Jensen's inequality

A relationship between probability and geometry

When *f* is convex function

$$f(\mathbb{E}[\{X_i\}]) \leq \mathbb{E}[f(\{X_i\})]$$

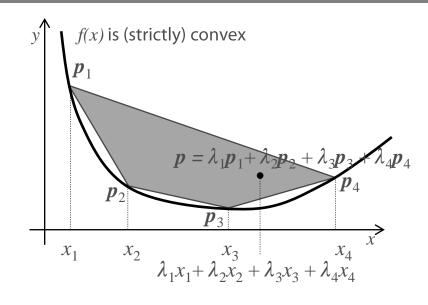
To see this, consider

$$\boldsymbol{p} = \lambda_1 \boldsymbol{p}_1 + \lambda_2 \boldsymbol{p}_2 + \lambda_3 \boldsymbol{p}_3 + \lambda_4 \boldsymbol{p}_4$$

i.e. a *linear combination* of  $p_i$  points

This is an *affine* combination if and it is a *convex* combination if also

$$\sum \lambda_i = 1$$
$$\lambda_i \ge 0, \ \forall i$$



When the  $\lambda_i$  define a probability, then p is a convex combination of  $p_i$  points

Any convex combination of  $p_i$  points lies inside their **convex hull** (see figure) and therefore above f:

$$\sum_{i} \lambda_{i} f(x_{i}) \geq f(\sum_{i} \lambda_{i} x_{i})$$

Corollary: the only way to make the convex hull be <u>on</u> f is to shrink it to a single point (i.e. the Jensen's corollary)

# Incomplete observations

Likelihood function with hidden random variables

$$\begin{split} L(\theta|D) &= P(D|\theta) = \prod_{m} P(D^{(m)}|\theta) \\ \ell(\theta|D) &= \sum_{m} \log P(D^{(m)}|\theta) = \sum_{m} \log \sum_{\{Z_i\}} P(D^{(m)}, \{X_i\}|\theta_k) \\ &= \sum_{m} \log \sum_{\{X_i\}} Q^{(m)}(\{X_i\}) \frac{P(D^{(m)}, \{X_i\}|\theta)}{Q^{(m)}(\{X_i\})} \\ &= \sum_{m} \log \mathbb{E}_{Q^{(m)}(\{X_i\})} \left[ \frac{P(D^{(m)}, \{X_i\}|\theta)}{Q^{(m)}(\{X_i\})} \right] &\geq \sum_{m} \mathbb{E}_{Q^{(m)}(\{X_i\})} \left[ \log \frac{P(D^{(m)}, \{X_i\}|\theta)}{Q^{(m)}(\{X_i\})} \right] \\ &= \sum_{m} \sum_{\{X_i\}} Q^{(m)}(\{X_i\}) \log \frac{P(D^{(m)}, \{X_i\}|\theta)}{Q^{(m)}(\{X_i\})} \end{split}$$

### Expectation-Maximization (EM) Algorithm

Alternate optimization (coordinate ascent)

Log-likelihood function:

$$\ell(\theta|D) \geq \sum_{m} \sum_{\{X_i\}} Q^{(m)}(\{X_i\}) \log \frac{P(D^{(m)}, \{X_i\}|\theta)}{Q^{(m)}(\{X_i\})}$$
This inequality becomes equality when this term is constant (see Jensen's corollary)

1) Keep  $\theta$  constant, define  $Q^{(m)}(\{Z_i\})$  so that the right side of the inequality is maximized

$$Q^{(m)}(\{X_i\}) := \frac{P(D^{(m)}, \{X_i\}|\theta)}{\sum\limits_{\{X_i\}} P(D^{(m)}, \{X_i\}|\theta)} = \frac{P(D^{(m)}, \{X_i\}|\theta)}{P(D^{(m)}|\theta)} = P(\{X_i\}|D^{(m)}, \theta) =: p_{\{X_i\}}^{(m)}$$
These numbers can be computed from the graphical model (i.e. as an inference step)

2) Then maximize the log-likelihood while keeping  $Q^{(m)}(\{Z_i\})$  constant

$$\theta^* = \operatorname{argmax}_{\theta} \sum_{m} \sum_{\{X_i\}} p_{\{X_i\}}^{(m)} \log \frac{P(D^{(m)}, \{X_i\} | \theta)}{p_{\{X_i\}}^{(m)}}$$

$$= \operatorname{argmax}_{\theta} \sum_{m} \left( \sum_{\{X_i\}} p_{\{X_i\}}^{(m)} \log P(D^{(m)}, \{X_i\} | \theta) - \sum_{\{X_i\}} p_{\{X_i\}}^{(m)} \log p_{\{X_i\}}^{(m)} \right) \right)$$

$$= \operatorname{argmax}_{\theta} \sum_{m} \sum_{\{X_i\}} p_{\{X_i\}}^{(m)} \log P(D^{(m)}, \{X_i\} | \theta)$$

$$= \operatorname{argmax}_{\theta} \sum_{m} \sum_{\{X_i\}} p_{\{X_i\}}^{(m)} \log P(D^{(m)}, \{X_i\} | \theta)$$

### Expectation-Maximization (EM) Algorithm

Alternate optimization (coordinate ascent)

Log-likelihood function and its estimator:

$$\ell(\theta|D) \geq \sum_{m} \sum_{\{X_i\}} Q^{(m)}(\{X_i\}) \log \frac{P(D^{(m)}, \{X_i\}|\theta)}{Q^{(m)}(\{X_i\})}$$

#### **Algorithm:**

- 1) Assign the  $\theta$  at random
- 2) (E-step) Compute the probabilities

$$p_{\{X_i\}}^{(m)} = Q^{(m)}(\{X_i\}) = P(\{X_i\}|D^{(m)},\theta)$$

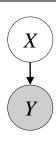
3) (*M-step*) Compute a new estimate of  $\theta$ 

$$\theta^* = \operatorname{argmax}_{\theta} \sum_{m} \sum_{\{X_i\}} p_{\{X_i\}}^{(m)} \log P(D^{(m)}, \{X_i\} | \theta)$$

4) Go back to step 2) until some convergence criterion is met

The algorithm converges to a local maximum of the log-likelihood The effectiveness of algorithm depends on the form of  $P(\{X_i\}|D^{(m)},\theta)$  (see step3) In particular, when this distribution is <u>exponential</u>... (e.g. Gaussian – see next slide) An aside: the EM algorithm in the continuous case (Mixture of Gaussians)

# EM Algorithm: mixture of Gaussians



#### **Model:**

The hidden variable X has k possible values, the observable variable Y is a point in  $\mathbb{R}^d$ 

$$P(X=k):=\phi_k$$
 Multivariate normal distribution  $P(Y=y|X=k)=\mathcal{N}(y;\mu_k,\Sigma_k):=(2\pi)^{-d/2}(\det\Sigma_k)^{-1/2}\exp\left(-rac{1}{2}(y-\mu_k)^T\Sigma_k^{-1}(y-\mu_k)
ight)$  i.e. the condition probabilities are normal distributions The observations are a set  $D=\{y^{(1)},\ldots,y^{(N)}\}$  of points in  $\mathbb{R}^d$ 

#### **Algorithm:**

- 1) For each value k, assign  $\phi_k$ ,  $\mu_k$  and  $\Sigma_k$  at random
- 2) (*E-step*) For all the  $y^{(m)}$  in D compute the probabilities  $p_k^{(m)} = P(X = k | y^{(m)}, \phi_k, \mu_k, \Sigma_k) = \phi_k \cdot \mathcal{N}(x^{(m)}; \mu_k, \Sigma_k)$
- 3) (*M-step*) Compute the new estimates for the parameters

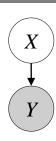
$$\phi_k = \frac{1}{n} \sum_{m} p_k^{(m)}$$

$$\mu_k = \frac{\sum_{m} p_k^{(m)} y^{(m)}}{\sum_{m} p_k^{(m)}}$$

$$\Sigma_k = \frac{\sum_{m} p_k^{(m)} (y - \mu_k) (y - \mu_k)^T}{\sum_{m} p_k^{(m)}}$$

4) Go back to step 2) until some convergence criterion is met

# EM Algorithm: mixture of Gaussians



#### **Model:**

The hidden variable X has k possible values, the variable Y is a point in  $\mathbb{R}^d$ 

$$P(X=k) := \phi_k$$

$$P(Y = y | X = k) = \mathcal{N}(y; \mu_k, \Sigma_k) := (2\pi)^{-d/2} (\det \Sigma_k)^{-1/2} \exp\left(-\frac{1}{2}(y - \mu_k)^T \Sigma_k^{-1}(y - \mu_k)\right)$$

i.e. the condition probabilities are <u>normal</u> distributions

The observations are a set  $D = \{y^{(1)}, \dots, y^{(N)}\}$  of points in  $\mathbb{R}^d$ 

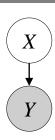
#### **Proof** (of the M-step):

$$\sum_{m} \sum_{k} p_{k}^{(m)} \log P(Y^{(m)}, X = k | \phi_{k}, \mu_{k}, \Sigma_{k})$$

$$= \sum_{m} \sum_{k} p_{k}^{(m)} \log P(Y^{(m)} | X = k, \mu_{k}, \Sigma_{k}) P(X = k | \phi_{k})$$

$$= \sum_{m} \sum_{k} p_{k}^{(m)} \left( \log \left( 2\pi^{-d/2} (\det \Sigma_{k})^{-1/2} \right) + \left( -\frac{1}{2} (y - \mu_{k})^{T} \Sigma_{k}^{-1} (y - \mu_{k}) \right) + \log \varphi_{k} \right)$$

# EM Algorithm: mixture of Gaussians



#### **Model:**

The hidden variable X has k possible values, the variable Y is a point in  $\mathbb{R}^d$ 

$$P(X=k) := \phi_k$$

$$P(Y = y | X = k) = \mathcal{N}(y; \mu_k, \Sigma_k) := (2\pi)^{-d/2} (\det \Sigma_k)^{-1/2} \exp\left(-\frac{1}{2}(y - \mu_k)^T \Sigma_k^{-1}(y - \mu_k)\right)$$

i.e. the condition probabilities are normal distributions

The observations are a set  $D = \{y^{(1)}, \dots, y^{(N)}\}\$  of points in  $\mathbb{R}^d$ 

#### **Proof** (of the M-step):

$$\begin{split} \frac{\partial}{\partial \mu_j} \sum_m \sum_k p_k^{(m)} \left( \log \left( (2\pi)^{-d/2} (\det \Sigma_k)^{-1/2} \right) + \left( -\frac{1}{2} (y^{(m)} - \mu_k)^T \Sigma_k^{-1} (y^{(m)} - \mu_k) \right) + \log \phi_k \right) \\ &= \frac{\partial}{\partial \mu_j} \sum_m \sum_k p_k^{(m)} \left( -\frac{1}{2} (y^{(m)} - \mu_k)^T \Sigma_k^{-1} (y^{(m)} - \mu_k) \right) \\ &= \frac{\partial}{\partial \mu_j} \sum_m \sum_k p_k^{(m)} \left( -\frac{1}{2} (y^{(m)}^T \Sigma_k^{-1} y^{(m)} + \mu_k^T \Sigma_k^{-1} \mu_k - 2 y^{(m)}^T \Sigma_k^{-1} \mu_k) \right) \\ &= \sum_m p_j^{(m)} \left( x^T \Sigma_j^{-1} - \mu_j^T \Sigma_j^{-1} \right) \\ &= \sum_m p_j^{(m)} \left( x^T \Sigma_j^{-1} - \mu_j^T \Sigma_j^{-1} \right) = 0 \qquad \Rightarrow \qquad \qquad \\ \text{In the web page for the derivations of other parameters} \dots \end{split}$$

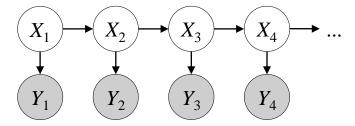
See the link in the web page for the derivations of other parameters ...

# The EM algorithm for learning with <u>missing data</u>

## Missing Values

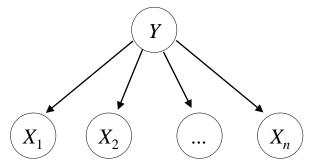
#### Hidden Variables

Some of the variables may be hidden, i.e., non observable 'by design' Example: 'Hidden Markov Model'



#### Incomplete Observations

Sometimes, however, observations may be missing 'by accident' and not 'by design' Example: 'Naïve Bayesian Classifier'



What if some classifications Y are missing, or a few features  $X_i$  are not available?

# Missing Values: Observability Model

#### Observed and Unobserved Variables

Let's consider a graphical model with a set of random variables:

$$\boldsymbol{X} := \{X_1, \dots, X_n\}$$

In each actual observation (i.e., a data item)

$$\boldsymbol{X}^{(m)} := \{X_1^{(m)}, \dots, X_n^{(m)}\}$$

each value  $X_i^{(i)}$  may be either observed or unobserved (i.e., missing) determined by a random variable  $\,O_{X_i}\in 0,1\,$ 

An **observability model** for a graphical model with random variables  $oldsymbol{X}$  is a set of random observability variables

$$O_X := \{O_{X_1}, \dots, O_{X_n}\}$$

with probability distribution

$$P_{missing}(\boldsymbol{X}, \boldsymbol{O}_{\boldsymbol{X}}) = P(\boldsymbol{X}) P_{missing}(\boldsymbol{O}_{\boldsymbol{X}} | \boldsymbol{X})$$

# Missing Values: MCAR

Missing Completely at Random (MCAR)

Fundamental assumption

$$\langle m{X} \perp m{O_X} 
angle$$

which entails that:

$$P_{missing}(\boldsymbol{X}, \boldsymbol{O}_{\boldsymbol{X}}) = P(\boldsymbol{X}) P_{missing}(\boldsymbol{O}_{\boldsymbol{X}})$$

This is tempting and it could ease all subsequent computations...

... but it is too strong, and hardly enforceable in many practical cases

Moral: we need a weaker assumption

# Missing Values: MAR

#### Missing at Random (MAR)

Consider a generic data item, with possibly missing values

$$oldsymbol{X}^{(m)} := oldsymbol{X}^{(m)}_{obs} \cup oldsymbol{X}^{(m)}_{hid}$$

Missing values need NOT be for the same variables in each data item

Fundamental assumption: in each data item

$$\langle oldsymbol{X}_{hid}^{(m)} \perp oldsymbol{O_{oldsymbol{X}}} \mid oldsymbol{X}_{obs}^{(m)} 
angle$$

Namely, the *values* of the missing variables are independent from their *observability* given the *values* of the observed variables

It is still a strong assumption, yet much more realistic...

# Missing Values: MAR

#### Missing at Random (MAR)

Fundamental assumption: in each data item

$$\langle oldsymbol{X}_{hid}^{(m)} \perp oldsymbol{O_{oldsymbol{X}}} \mid oldsymbol{X}_{obs}^{(m)} 
angle$$

This entails:

$$P_{missing}(\boldsymbol{X}^{(m)}, \boldsymbol{O}_{\boldsymbol{X}}) = P_{missing}(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)}, \boldsymbol{O}_{\boldsymbol{X}})$$

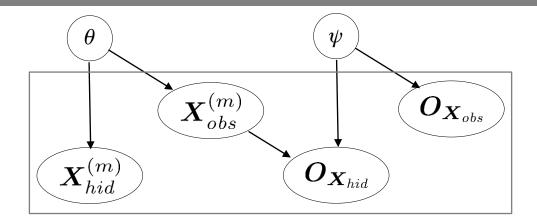
$$= P(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)}) P_{missing}(\boldsymbol{O}_{\boldsymbol{X}} | \boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)})$$

$$= P(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)}) P_{missing}(\boldsymbol{O}_{\boldsymbol{X}} | \boldsymbol{X}_{obs}^{(m)})$$

$$P_{missing}(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{O}_{\boldsymbol{X}}) = \sum_{\boldsymbol{X}_{hid}} P(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)}) P_{missing}(\boldsymbol{O}_{\boldsymbol{X}} | \boldsymbol{X}_{obs}^{(m)})$$

$$P_{missing}(m{X}_{obs}^{(m)},m{O_X}) = P(m{X}_{obs}^{(m)})\,P_{missing}(m{O_X}|m{X}_{obs}^{(m)})$$
 This is the relevant property

#### Likelihood under MAR



$$\langle oldsymbol{X}_{hid}^{(m)} \perp oldsymbol{O_{oldsymbol{X}}} \mid oldsymbol{X}_{obs}^{(m)} 
angle$$

Variables and parameters in an observability model as a graphical model

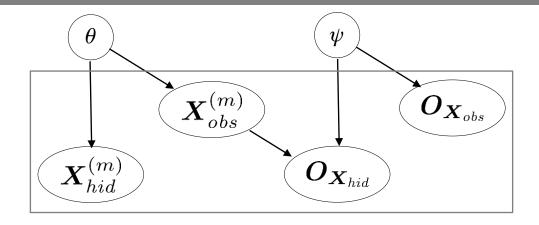
#### Likelihood

$$\begin{split} L(\theta, \psi \mid D) &= \prod_{m=1}^{N} P(D^{(m)} \mid \theta, \psi) \quad \text{where:} \quad D^{(m)} := \{\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)}\} \\ l(\theta, \psi \mid D) &= \sum_{m=1}^{N} \log P(D^{(m)} \mid \theta, \psi) \\ &= \sum_{m=1}^{N} \log \left( P(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)} \mid \theta, \psi) P_{missing}(\boldsymbol{O}_{\boldsymbol{X}} | \boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)}, \theta, \psi) \right) \\ &= \sum_{m=1}^{N} \log \left( P(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)} \mid \theta) P_{missing}(\boldsymbol{O}_{\boldsymbol{X}} | \boldsymbol{X}_{obs}^{(m)}, \psi) \right) \\ &= \sum_{m=1}^{N} \log P(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)} \mid \theta) + \sum_{m=1}^{N} \log P_{missing}(\boldsymbol{O}_{\boldsymbol{X}} | \boldsymbol{X}_{obs}^{(m)}, \psi) \end{split}$$

We are interested in optimizing heta ...

yet we have only <u>observed</u> values

#### Likelihood under MAR



$$\langle oldsymbol{X}_{hid}^{(m)} \perp oldsymbol{O_{oldsymbol{X}}} \mid oldsymbol{X}_{obs}^{(m)} 
angle$$

Variables and parameters in an observability model as a graphical model

#### Likelihood (for observed values)

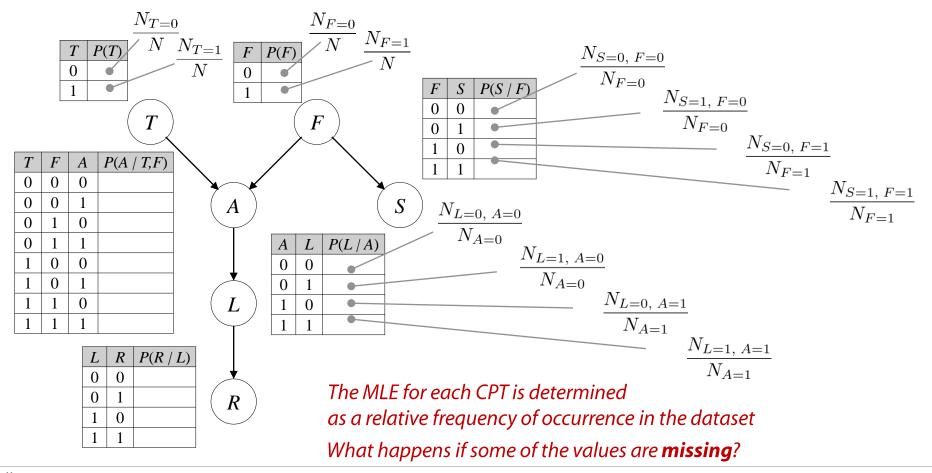
$$\begin{split} l(\theta \mid D) &:= \sum_{m=1}^{N} \log P(\boldsymbol{X}_{obs}^{(m)} \mid \theta) \\ &= \sum_{m=1}^{N} \log \sum_{\boldsymbol{X}_{hid}^{(m)}} P(\boldsymbol{X}_{obs}^{(m)}, \boldsymbol{X}_{hid}^{(m)} \mid \theta) \\ &= \sum_{m=1}^{N} \log \sum_{\boldsymbol{X}_{hid}^{(m)}} \left( P(\boldsymbol{X}_{obs}^{(m)} \mid \theta) P(\boldsymbol{X}_{hid}^{(m)} \mid \boldsymbol{X}_{obs}^{(m)}, \theta) \right) \end{split}$$

Looks promising: using probabilities instead of missing values ... but this may be very hard to optimize in general

**Model**: random variables plus the graph of dependencies

**Observations**: dataset of values, from <u>completely observed</u> outcomes

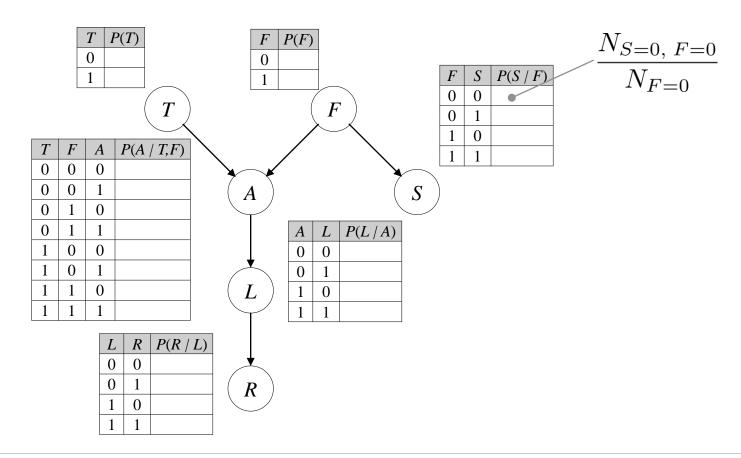
**Parameters (to be determined)**: all conditional probabilities (i.e. all CPTs)



Artificial Intelligence 2022-2023

#### Fundamental idea: using probabilities of observations

In the completely observed case: probabilities are estimated as frequencies of occurrence



Artificial Intelligence 2022-2023

#### Fundamental idea: using probabilities of missing observations

Let's consider a dataset 
$$D = \left\{ oldsymbol{X}^{(i)} 
ight\}_{i=1}^{N}$$

Each data item  $oldsymbol{X}^{(i)}$  may contain some missing data

Example: 
$$\boldsymbol{X}^{(i)} = (X_1^{(i)} = x_1, X_2^{(i)} = x_2, X_3^{(i)} = ?)$$

Define  $ilde{m{X}}^{(i)}$  as one possible **completion** of  $m{X}^{(i)}$  , namely one in which there are no missing data

Example: assuming that 
$$X_3 \in \{0,1\}$$
 ,  $\begin{array}{l} (X_1^{(i)} = x_1, X_2^{(i)} = x_2, X_3^{(i)} = 0) \\ (X_1^{(i)} = x_1, X_2^{(i)} = x_2, X_3^{(i)} = 1) \end{array}$  are the two possible completions of  $\boldsymbol{X}^{(i)}$ 

Note that there will be as many completions of  $X^{(i)}$  as there are combinations of possible values for the missing data For any <u>complete</u> observation,  $\tilde{X}^{(i)} = X^{(i)}$ : there is only one possible completion that coincides with the data item itself

Likewise,  $m{X}_{obs}^{(i)}$  is the part of  $m{X}^{(i)}$  which contain the actual observations

Example: 
$$\boldsymbol{X}_{obs}^{(i)} = (X_1^{(i)} = x_1, X_2^{(i)} = x_2)$$

#### Fundamental idea: using probabilities of observations

In the completely observed case: probabilities are estimated as frequencies of occurrence More in general:

$$\frac{N_{X_i, \mathbf{Z}}}{N_{\mathbf{Z}}}$$
 where:  $\mathbf{Z} = parents(X_i)$ 

In the EM algorithm, use *estimated* occurrences:

EM algorithm, use *estimated* occurrences: 
$$\frac{\tilde{N}_{X_i, \mathbf{Z}}}{\tilde{N}_{\mathbf{Z}}} \qquad \text{where:} \qquad \tilde{N}_{\mathbf{X}} := \sum_{i=1}^{N} \sum_{\tilde{\boldsymbol{X}}^{(i)}} P(\tilde{\boldsymbol{X}}^{(i)} \mid \boldsymbol{X}_{obs}^{(i)}, \theta)$$

*In words, any incomplete observations 'splits up' and contributes with the probabilities of possible completions Note that, when all observations are complete:* 

$$\tilde{N}_{\boldsymbol{X}} = N_{\boldsymbol{X}}$$

Fundamental idea: using probabilities of observations

#### **Algorithm:**

- 1) Assign parameters  $\theta^{(0)}$  at random
- 2) Compute  $P(oldsymbol{X} \mid heta^{(t)})$
- 3) Update all parameters using estimated occurrences:

$$heta_{X_i \,|\, m{Z}}^{(t+1)} = rac{ ilde{N}_{X_i, m{Z}}}{ ilde{N}_{m{Z}}} \quad ext{where all estimations are made using } P(m{X} \,|\, m{ heta}^{(t)})$$

3) Go back to step 2) until some convergence criterion is met