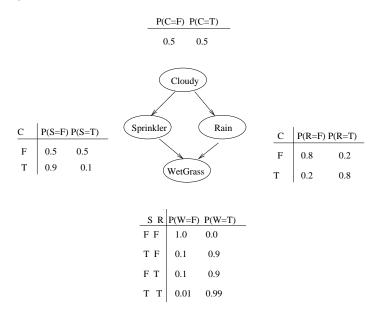
LECTURE 10:

PARAMETER LEARNING FOR BAYES NETS

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- Inference means computing $P(X_i|\theta,G)$
- Structure learning/ model selection = inferring G from data.
- Parameter learning/ estimation = inferring θ from data.



- \bullet Assume G is known and fixed and is a DAG.
- Goal: estimate θ from a dataset of M independent, identically distributed (iid) training cases $D = (x^1, \dots, x^M)$.
- In general, each training case $x^m = (x_1^m, \ldots, x_N^m)$ is a vector of values, one per node. (Think of a database with M rows and N columns.)
- We assume complete observability, i.e., every entry in the database is known (no missing values, no hidden variables).
- Initially we consider learning parameters for a single node.
- Then we consider how to learn parameters for a whole network.

• Bayesians treat the unknown parameters θ as a random variable, which can be estimated using Bayes rule:

$$p(\theta|\mathcal{D}) = \frac{p(\mathcal{D}|\theta)p(\theta)}{p(\mathcal{D})}$$

• This crucial equation can be written in words:

$$posterior = rac{likelihood imes prior}{marginal likelihood}$$

• For iid data, the likelihood is

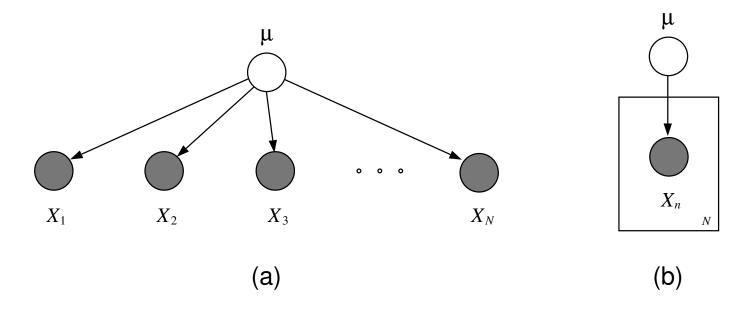
$$p(D|\theta) = \prod_{m} p(x_{m}|\theta)$$

 \bullet The prior $p(\theta)$ encodes our prior knowledge about the domain.

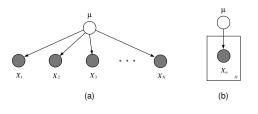
• For iid (exchangeable) data, the likelihood is

$$p(D|\theta) = \prod_{m} p(x_m|\theta)$$

- \bullet We can represent this as a Bayes net with M nodes.
- "Plates" provide a more compact representation for repetitive structure, and are very common in Bayesian models.



- "Plates" provide a compact representation for repetitive structure.
- The rules of plates are simple: repeat every structure in a box a number of times given by the integer in the corner of the box (e.g. N), updating the plate index variable (e.g. n) as you go.
- Duplicate every arrow going into the plate and every arrow leaving the plate by connecting the arrows to each copy of the structure.
- Plates are closely related to probabilistic relational models, and object oriented Bayes nets, which are forms of "syntactic sugar" for parameter tying (sharing).



- \bullet Two people with different priors $p(\theta)$ will end up with different estimates $p(\theta|D).$
- Frequentists dislike this "subjectivity".
- Frequentists think of the parameter as a fixed, unknown constant, not a random variable.
- Hence they have to come up with different estimators (ways of computing θ from data), instead of using Bayes' rule.
- These estimators have different properties, such as being "unbiased", "minimum variance", etc.
- A very popular estimator is the *maximum likelihood estimator*, which is simple and has good statistical properties.

• The log-likelihood is monotonically related to the likelihood:

$$\ell(\theta; D) = \log p(D|\theta) = \sum_{m} \log p(x^{m}|\theta)$$

 Idea of maximum likelihood estimation (MLE): pick the setting of parameters most likely to have generated the data we saw:

$$\hat{\theta}_{ML} = \operatorname{argmax}_{\theta} \ell(\theta; \mathcal{D})$$

• Often the MLE overfits the training data, so it is common to maximize a penalized log-likelihood instead:

$$\hat{\theta}_{MAP} = \operatorname{argmax}_{\theta} \ \ell(\theta; \mathcal{D}) - c(\theta)$$

• This is equivalent to picking the mode of $P(\theta|D)$, where $c(\theta) = -\log p(\theta)$, since

$$\log p(\theta|D) = \log p(D|\theta) + \log p(\theta) + c$$

- $\hat{\theta}_{MAP}$ is not Bayesian (even though it uses a prior) since it is a point estimate.
- Consider predicting the future. A Bayesian will integrate out all uncertainty:

$$p(\mathbf{x}_{\text{new}}|\mathbf{X}) = \int p(\mathbf{x}_{\text{new}}, \theta | \mathbf{X}) d\theta$$

= $\int p(\mathbf{x}_{\text{new}}|\theta, \mathbf{X}) p(\theta | \mathbf{X}) d\theta$
 $\propto \int p(\mathbf{x}_{\text{new}}|\theta) p(\mathbf{X}|\theta) p(\theta) d\theta$
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 A frequentist will typically use a "plug-in" estimator such as ML/MAP:

$$p(\mathbf{x}_{\text{new}}|\mathbf{X}) = p(\mathbf{x}_{\text{new}}|\hat{\theta}), \quad \hat{\theta} = \arg\max_{\theta} p(\mathbf{X}|\theta)$$

- This is a "theological" war.
- Advantages of Bayesian approach:
 - Mathematically elegant.
 - Works well when amount of data is much less than number of parameters (e.g., one-shot learning).
 - Easy to do incremental (sequential) learning.
 - Can be used for model selection (max likelihood will always pick the most complex model).
- Advantages of frequentist approach:
 - Mathematically/ computationally simpler.
- As $|D| \to \infty$, the two approaches become the same:

 $p(\theta|D) \to \delta(\theta, \hat{\theta}_{ML})$

- We observe M iid coin flips: $\mathcal{D}=H,H,T,H,\ldots$
- Model: $p(H) = \theta$ $p(T) = (1 \theta)$
- Likelihood:

$$\ell(\theta; \mathcal{D}) = \log p(\mathcal{D}|\theta) = \log \prod_{m} \theta^{\mathbf{x}^{m}} (1-\theta)^{1-\mathbf{x}^{m}}$$
$$= \log \theta \sum_{m} \mathbf{x}^{m} + \log(1-\theta) \sum_{m} (1-\mathbf{x}^{m})$$
$$= \log \theta N_{\mathrm{H}} + \log(1-\theta) N_{\mathrm{T}}$$

• Take derivatives and set to zero:

$$\frac{\partial \ell}{\partial \theta} = \frac{N_{\rm H}}{\theta} - \frac{N_{\rm T}}{1 - \theta}$$
$$\Rightarrow \theta_{\rm ML}^* = \frac{N_{\rm H}}{N_{\rm H} + N_{\rm T}}$$

• The counts $N_H = \sum_m x^m$ and $N_T = \sum_m (1 - x^m)$ are sufficient statistics of the data D.

 \bullet In general, T(X) is a sufficient statistic for X if $T(x^1)=T(x^2) \Rightarrow L(\theta;x^1)=L(\theta;x^2)$

- We observe M iid die rolls (K-sided): $\mathcal{D}=3,1,K,2,\ldots$
- Model: $p(k) = \theta_k$ $\sum_k \theta_k = 1$
- Likelihood (for binary indicators $[\mathbf{x}^m = k]$):

$$\ell(\theta; \mathcal{D}) = \log p(\mathcal{D}|\theta) = \sum_{m} \log \prod_{k} \theta_{1}^{[\mathbf{x}^{m}=k]}$$
$$= \sum_{m} \sum_{k} [\mathbf{x}^{m}=k] \log \theta_{k} = \sum_{k} N_{k} \log \theta_{k}$$

• We need to maximize this subject to the constraint $\sum_k \theta_k = 1$, so we use a Lagrange multiplier.

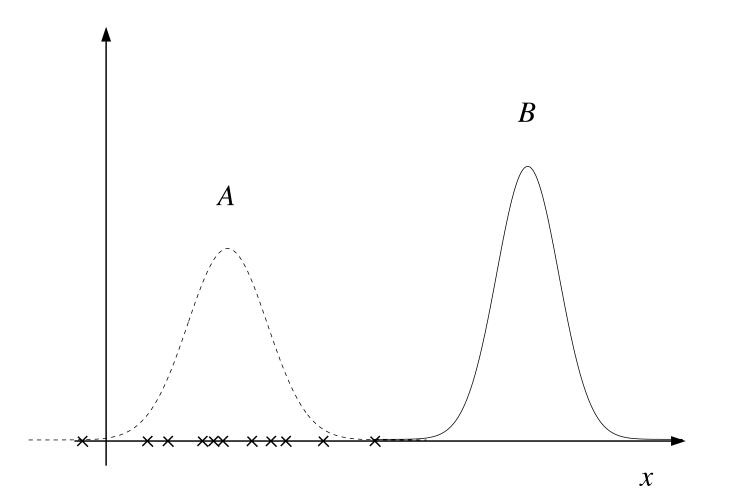
• Constrained cost function:

$$\tilde{l} = \sum_{k} N_k \log \theta_k + \lambda \left(1 - \sum_{k} \theta_k \right)$$

• Take derivatives wrt θ_k :

$$\begin{aligned} \frac{\partial \tilde{l}}{\partial \theta_k} &= \frac{N_k}{\theta_k} - \lambda = 0\\ N_k &= \lambda \theta_k\\ \sum_k N_k &= M = \lambda \sum_k \theta_k = \lambda\\ \hat{\theta}_{k,ML} &= \frac{N_k}{M} \end{aligned}$$

• $\hat{\theta}_{k,ML}$ if the fraction of times k occurs.



- We observe M iid real samples: $\mathcal{D}=1.18,-.25,.78,\ldots$
- Model: $p(x) = (2\pi\sigma^2)^{-1/2} \exp\{-(x-\mu)^2/2\sigma^2\}$
- Log likelihood:

$$\ell(\theta; \mathcal{D}) = \log p(\mathcal{D}|\theta)$$
$$= -\frac{M}{2}\log(2\pi\sigma^2) - \frac{1}{2}\sum_m \frac{(x^m - \mu)^2}{\sigma^2}$$

• Take derivatives and set to zero:

$$\frac{\partial \ell}{\partial \mu} = (1/\sigma^2) \sum_m (x_m - \mu)$$
$$\frac{\partial \ell}{\partial \sigma^2} = -\frac{M}{2\sigma^2} + \frac{1}{2\sigma^4} \sum_m (x_m - \mu)^2$$
$$\Rightarrow \mu_{\rm ML} = (1/M) \sum_m x_m$$
$$\sigma_{\rm ML}^2 = (1/M) \sum_m (x_m - \mu_{\rm ML})^2$$

 \bullet For a numeric random variable ${\bf x}$

$$p(\mathbf{x}|\eta) = h(\mathbf{x}) \exp\{\eta^{\top} T(\mathbf{x}) - A(\eta)\}$$
$$= \frac{1}{Z(\eta)} h(\mathbf{x}) \exp\{\eta^{\top} T(\mathbf{x})\}$$

is an exponential family distribution with *natural (canonical) parameter* η .

- Function $T(\mathbf{x})$ is a *sufficient statistic*.
- Function $A(\eta) = \log Z(\eta)$ is the log normalizer.
- Examples: Bernoulli, multinomial, Gaussian, Poisson, gamma,...
- A distribution p(x) has finite sufficient statistics (independent of number of data cases) iff it is in the exponential family.

• For a continuous vector random variable:

$$p(x|\mu, \Sigma) = |2\pi\Sigma|^{-1/2} \exp\left\{-\frac{1}{2}(\mathbf{x}-\mu)^{\top}\Sigma^{-1}(\mathbf{x}-\mu)\right\}$$

• Exponential family with:

$$\eta = [\Sigma^{-1}\mu; -1/2\Sigma^{-1}]$$
$$T(x) = [\mathbf{x}; \mathbf{x}\mathbf{x}^{\top}]$$
$$A(\eta) = \log |\Sigma|/2 + \mu^{\top}\Sigma^{-1}\mu/2$$
$$h(x) = (2\pi)^{-d/2}$$

 Note: a d-dimensional Gaussian is a d+d²-parameter distribution with a d+d²-component vector of sufficient statistics (but because of symmetry and positivity, parameters are constrained)

- We can easily compute moments of any exponential family distribution by taking the derivatives of the log normalizer $A(\eta)$.
- The q^{th} derivative gives the q^{th} centred moment.

$$\frac{dA(\eta)}{d\eta} = \text{mean}$$
$$\frac{d^2A(\eta)}{d\eta^2} = \text{variance}$$

• When the sufficient statistic is a vector, partial derivatives need to be considered.

$$\begin{aligned} \frac{dA}{d\eta} &= \frac{d}{d\eta} \log Z(\eta) = \frac{1}{Z(\eta)} \frac{d}{d\eta} Z(\eta) \\ &= \frac{1}{Z(\eta)} \frac{d}{d\eta} \int h(\mathbf{x}) \exp\{\eta T(\mathbf{x})\} dx \\ &= \frac{\int T(\mathbf{x}) h(\mathbf{x}) \exp\{\eta T(\mathbf{x})\}}{Z(\eta)} \\ &= ET(X) \\ \frac{d^2 A}{d\eta^2} &= VarT(X) \end{aligned}$$

• The moment parameter μ can be derived from the natural (canonical) parameter

$$\frac{dA}{d\eta} = ET(X) \stackrel{\text{def}}{=} \mu$$

 $\bullet \ \mathrm{Now} \ A(\eta)$ is convex since

$$\frac{d^2A}{d\eta^2} = VarT(X) > 0$$

• Hence we can invert the relationship and infer the canonical parameter from the moment parameter:

$$\eta \stackrel{\text{def}}{=} \psi(\mu)$$

• For iid data, the log-likelihood is

$$\ell(\eta; \mathcal{D}) = \log \prod_{m} h(x^{m}) \exp\left(\eta^{T} T(x^{m}) - A(\eta)\right)$$
$$= \left(\sum_{m} \log h(\mathbf{x}^{m})\right) - MA(\eta) + \left(\eta^{\top} \sum_{m} T(\mathbf{x}^{m})\right)$$

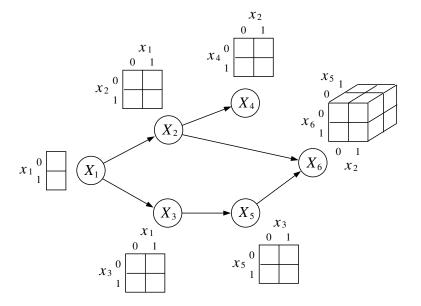
• Take derivatives and set to zero:

$$\begin{aligned} \frac{\partial \ell}{\partial \eta} &= \sum_{m} T(\mathbf{x}^{m}) - M \frac{\partial A(\eta)}{\partial \eta} = 0\\ \Rightarrow \frac{\partial A(\eta)}{\partial \eta} &= \frac{1}{M} \sum_{m} T(\mathbf{x}^{m})\\ \hat{\mu}_{\mathrm{ML}} &= \frac{1}{M} \sum_{m} T(\mathbf{x}^{m}) \end{aligned}$$

- This amounts to moment matching.
- \bullet We can infer the canonical parameters using $\hat{\eta}_{ML}=\psi(\hat{\mu}_{ML})$

• If we assume the parameters for each CPD are globally independent, then the log-likelihood function decomposes into a sum of local terms, one per node:

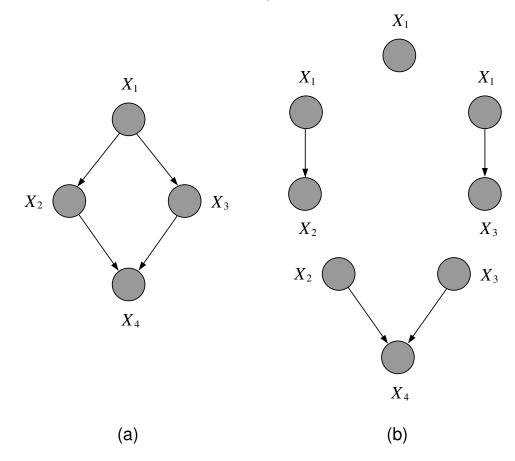
$$\log p(\mathcal{D}|\theta) = \log \prod_{m} \prod_{i} p(\mathbf{x}_{i}^{m} | \mathbf{x}_{\pi_{i}}, \theta_{i}) = \sum_{i} \sum_{m} \log p(\mathbf{x}_{i}^{m} | \mathbf{x}_{\pi_{i}}, \theta_{i})$$



• Consider the distribution defined by the DAGM:

 $p(\mathbf{x}|\theta) = p(\mathbf{x}_1|\theta_1)p(\mathbf{x}_2|\mathbf{x}_1,\theta_2)p(\mathbf{x}_3|\mathbf{x}_1,\theta_3)p(\mathbf{x}_4|\mathbf{x}_2,\mathbf{x}_3,\theta_4)$

• This is exactly like learning four separate small DAGMs, each of which consists of a node and its parents.



• Assume each CPD is represented as a table (multinomial) where

$$\theta_{ijk} \stackrel{\text{def}}{=} P(X_i = j | X_{\pi_i} = k)$$

• The sufficient statistics are just counts of family configurations

$$N_{ijk} \stackrel{\text{def}}{=} \sum_{m} I(X_i^m = j, X_{\pi_i}^m = k)$$

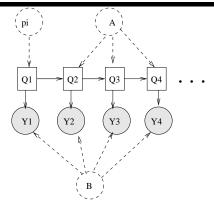
• The log-likelihood is

$$\ell = \log \prod_{m} \prod_{ijk} \theta_{ijk}^{N_{ijk}}$$
$$= \sum_{m} \sum_{ijk} N_{ijk} \log \theta_{ijk}$$

• Using a Lagrange multiplier to enforce so $\sum_{j} \theta_{ijk} = 1$ we get

$$\hat{\theta}_{ijk}^{ML} = \frac{N_{ijk}}{\sum_{j'} N_{ij'k}}$$

TIED PARAMETERS

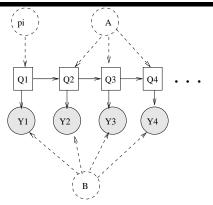


- Consider a time-invariant hidden Markov model (HMM)
 - -State transition matrix $A(i,j) \stackrel{\text{def}}{=} P(X_t = j | X_{t-1} = i)$,
 - -Discrete observation matrix $B(i,j) \stackrel{\text{def}}{=} P(Y_t = j | X_t = i)$
 - -State prior $\pi(i) \stackrel{\text{def}}{=} P(X_1 = i).$

The joint is

$$P(X_{1:T}, Y_{1:T}|\theta) = P(X_1|\pi) \prod_{t=2}^{T} P(X_t|X_{t-1}, A) \prod_{t=1}^{T} P(Y_t|X_t; B)$$

LEARNING A FULLY OBSERVED HMM



• The log-likelihood is

$$\ell(\theta; D) = \sum_{m} \log P(X_1 = x_1^m | \pi) + \sum_{t=2}^{T} P(X_t = x_t^m | X_{t-1} = x_{t-1}^m, A) + \sum_{t=1}^{T} P(Y_t = y_t^m | X_t = x_t^m, B)$$

• We can optimize each parameter (A, B, π) separately.

LEARNING A MARKOV CHAIN TRANSITION MATRIX

- Define $A(i, j) = P(X_t = j | X_{t-1} = i)$.
- A is a stochastic matrix: $\sum_j A(i,j) = 1$
- Each row of A is multinomial distribution.
- \bullet So MLE is the fraction of transitions from i to j

$$\hat{A}_{ML}(i,j) = \frac{\#i \to j}{\sum_k \#i \to k} = \frac{\sum_m \sum_{t=2}^T I(X_{t-1}^m = i, X_t^m = j)}{\sum_m \sum_{t=2}^T I(X_{t-1}^m = i)}$$

- If the states X_t represent words, this is called a *bigram language model*.
- Note that $\hat{A}_{ML}(i, j) = 0$ if the particular i, j pair did not occur in the training data; this is called the *sparse data problem*.
- We will solve this later using a prior.

- So far we have considered the case where $p(y|x, \theta)$ can be represented as a multinomial (table).
- Now we consider the case where some nodes may be continuous.

X	Y	p(Y X)
\mathbb{R}^n	\mathbb{R}^m	regression
\mathbb{R}^{n}	$\{0, 1\}$	binary classification
$\{0,1\}^n$	$\{0, 1\}$	binary classification
\mathbb{R}^{n}	$\{1,\ldots,K\}$	multiclass classification
$\{1,\ldots,K\}$	\mathbb{R}^{n}	conditional density modeling

• Consider an HMM with discrete states X_t but continuous observations $y_t \in \mathbb{R}^n$:

$$p(y_t | X_t = i) = \mathcal{N}(y_t; \mu_i, \Sigma_i)$$

• The MLE is the sample mean and sample variance of observations associated with each state (use X_t labels to partition the data):

$$\hat{\mu}_{ML}(i) = \frac{\sum_{m,t:X_t^m = i} y_t^m}{\sum_{m,t} y_t^m} = \frac{\sum_m \sum_{t=1}^T I(X_t^m = i) y_t^m}{\sum_m \sum_{t=1}^T y_t^m}$$

- Note that the MLE for Σ_i for states i with small numbers of observations is $\Sigma_i \to \infty I$.
- We will solve this later using a prior.