Lecture 21: April 8, 2015 cs 573: Probabilistic Reasoning Professor Nevatia Spring 2015

#### Review

- HW #6B posted, due 4/13/15
- Previous Lecture
  - Inference in temporal models
- Today's objective
  - Intro to learning

# Learning

- How to construct a graphical model for a problem domain?
  - Structure
    - What are the variables of interest? If they are continuous valued, should they be discretized?
    - What should be the links between the variable nodes?
      - Should the edges be directed or not?
  - Parameters
    - Specify CPDs or affinity factors (for MNs), log linear models for continuous variables
- Can the structure and parameters both be learned simply from examples?

### Motivations for Learning

- Problem domain is too complex to design by hand
  - Consider examples of medical diagnosis, computer vision, language understanding...
  - Medical knowledge, language grammar rules exist but may not be completely correct or precise
  - Human vision (and speech) understanding is not fully open to introspection...
- Distributions change over time or from site to site
- Much training data may be available in age of "big data"
  - Evidence-based medicine
- Note: this is NOT a course in Machine Learning, we confine to learning of PGMs
  - Possible overlap with cs567 and some EE courses

# **Learning Topics**

- Is structure given or to be learnt? If former, task is that of parameter learning
- Is any prior knowledge of the model, such as parameter distributions given?
- Complete data (each sample includes values of all the variables in the model) *vs* incomplete data (values of some variables are missing (*hidden*), in each instance.
- Bayesian networks vs Markov networks
  - Parameter estimation in MNs is significantly harder as the partition function entails all variables as opposed to the local normalizing constants in factored CPDs for BNs.
- We will first focus on parameter learning in BNs

### Goals of Learning

- Let P\* be the underlying probability distribution and M\* be the corresponding graphical model (BN, MN, template model...)
- Given M samples from P\*, our goal is to estimate M\*;
  - Find M<sup>∼</sup> that best approximates M<sup>\*</sup>.
- Several notions of approximations are possible:
  - Probability Density (distribution) estimation
  - Prediction/classification tasks
  - Knowledge discovery (e.g. learning a physical law)

### **Density Estimation**

- Find  $P^{\sim}$  that best approximates the *generating distribution*  $P^{*}$ .
- Use KL-divergence to measure similarity (as in variational approximations)
- $D(P^*||P^-) = E_{\xi \sim P^*} [\log ((P^*(\xi)/P^-(\xi)))]$ =  $-H_P(X) - E_{\xi \sim P^*} [\log P^-(\xi)]$ (X is the set of variables over which the distributions are defined)
- Only the second term is a function of P<sup>-</sup>, and is called *expected log-likelihood*. We want to maximize this term.
  - Assigns high probabilities to observed instances

#### Likelihood

- *likelihood* of the data, given a model M is P(D : M)
  - Note D is a set of data, probability of set D when the model is given by M
  - ":" is not the same as "|" in conditional probability
- Let D be set of M independent, identically distributed (iid) samples:

$$D = \{\xi[1], \xi[2], \dots, \xi[M]\}$$

then, 
$$P(D:M) = \prod_{m=1,M} P(\xi[m]:M)$$

•  $log-likelihood \ l(D:M) = log P(D:M)$ 

$$log P(D : M) = \sum_{m=1,M} P(log \xi[m] : M)$$

- log-loss is negative of log-likelihood = -l (D : M)
- loss ( $\xi$ : M) is loss associated with sample  $\xi$ 
  - Loss = 0 if P  $(\xi : M) = 1$
- Expected loss (or risk) is  $E_{\xi \sim P^*}$  [loss  $(\xi : M)$  ]

#### **Prediction Task**

- We may care about a specific aspect of the distribution only, say predicting P(Y|X); X and Y are specific subsets of the variables
- Most common is the task of classification; find the most likely assignment for Y.
  - Loss function could be binary: how many samples are classified correctly?
- This can be considered to be *discriminative* training
- We will focus on density estimation in this course which may be considered to be *generative* training
- Discriminative learning has shown better performance in recent years (for specific tasks) but typically requires much more training data (harder to generalize)

### Overfitting etc

- Overfitting and generalization
  - Learned model fits the training data very well but does not generalize to unseen data (e.g. new patients, new images....)

#### Bias-variance

- If we choose a simple model for approximation, it will remain sub-optimal even with large training data: this introduces a bias.
- If model is highly expressive (has many parameters), it can fit
   P\* well but requires more training data: with limited data,
   results will have high *variance*.

#### Regularization

 Model that has terms that reward good fit to the data but also terms that penalize addition of parameters

### **Evaluating Performance**

- Holdout testing: split data into training and test sets:  $D_{train}$  and  $D_{test}$ .
  - Can be used to compare performance of different learning methods
  - For one method, difference between performance on the training and test sets should not be too large; otherwise *overfitting* is indicated
  - How to split data between two sets? Trade-off between not enough data for training vs not enough for testing
- Cross-validation
  - Divide data into k sets; use one for testing, rest for training
  - Repeat *k*-times; *k*-fold cross-validation.
  - Leave one out: test on only one, use rest for testing.
  - Can also measure variance in performance
- Validation dataset
  - In cross-validation, test data is also used to select the best algorithm (or parameters)
  - Should test on a set not used in the optimization procedure
- Independent Test Set: use only once!

### Parameter Estimation (ch. 17 KF)

- Dataset D =  $\{\xi[1], \xi[2], \dots, \xi[M]\}$
- Two approaches
  - Maximum likelihood estimator: model giving highest probability to data
  - Bayesian estimator: takes prior probabilities of distributions into account
- Example: Toss a thumbtack, heads or tails

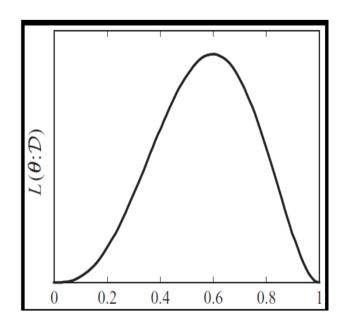


- Assume trials are iid (independent and identically distributed)
- $-\theta$  is the probability of heads
- $\Theta$ : hypothesis space (set of all parameters  $\theta$  in [0,1])

#### Parameter Estimation

- $P(\langle H, T, T, H, H \rangle) = \theta (1 \theta)(1 \theta)\theta + \theta = \theta^3(1 \theta)^2$
- *Likelihood function*, likelihood as a function of parameters  $(\theta)$ ; note reversal of order of data and  $\theta$  in the following notation:

$$L(\theta : \langle H, T, T, H, H \rangle) = P(\langle H, T, T, H, H \rangle : \theta) = \theta^{3}(1-\theta)^{2}$$



$$Log L = 3 log \theta + 2 log (1-\theta)$$

Take derivative w.r.t.  $\theta$  and set = 0 to maximize; gives  $\theta = .6$ 

Agrees with "frequency" interpretation

#### General Case

- General likelihood for M[1] heads, M[0] tails is  $L(\theta : D) = \theta^{M[1]} (1-\theta)^{M[0]}$
- log-likelihood is  $l(\theta : D) = M[1]log\theta + M[0]log(1-\theta)$
- Differentiate w.r.t.  $\theta$ , set equal to zero, we get

$$\hat{\theta} = \frac{M[1]}{M[1] + M[0]}$$

- Answer is consistent with our intuition, just measure the frequency
- Note that MLE estimate may be incorrect if number of trials is small (e.g. 1). *Confidence interval* is an estimate of the variance.

### MLE Principle

- X is the set of random variables
- Dataset D =  $\{\xi[1], \xi[2], \dots, \xi[M]\}$
- *Parametric* model:  $P(\xi; \theta)$ ;  $\xi$  is an instance,  $\sum_{\xi} P(\xi : \theta) = 1$
- $L(\boldsymbol{\theta}:D) = \prod_{m} P(\boldsymbol{\xi}[m]:\boldsymbol{\theta})$
- MLE gives  $\theta^{\wedge} = \max_{\theta \text{ in } \Theta} L(\theta : D)$
- Various Cases
  - $-\Theta_{\text{thumbtack}} = [0,1]$
  - Let X take one of k values  $x^1, \ldots, x^k$ ; then,
    - $P_{\text{multinomial}}(x; \boldsymbol{\theta}) = \theta_k \text{ if } x = x^k; \text{ note } \boldsymbol{\theta} \text{ is a k-dimensional vector; } \theta_k \text{ is the } k^{\text{th}} \text{ element.}$ 
      - $\Theta_{\text{multinomial}} = \{ \mathbf{\theta} \text{ in } [0,1]^k : \sum_i \theta_i = 1 \}$
  - $-\Theta_{Gaussian} = R \times R^+; \theta = <\mu,\sigma> (note \sigma must be positive)$

- Sufficient Statistics Informally: captures all the needed info in the samples;
- Formal Definition 17.1

A function  $\tau(\xi)$  from instances of  $\mathcal{X}$  to  $\mathbb{R}^{\ell}$  (for some  $\ell$ ) is a sufficient statistic if, for any two data sets  $\mathcal{D}$  and  $\mathcal{D}'$  and any  $\theta \in \Theta$ , we have that

$$\sum_{\xi[m]\in\mathcal{D}}\tau(\xi[m])=\sum_{\xi'[m]\in\mathcal{D}'}\tau(\xi'[m]) \quad \Longrightarrow \quad L(\theta:\mathcal{D})=L(\theta:\mathcal{D}').$$

We often refer to the tuple  $\sum_{\xi[m]\in\mathcal{D}} \tau(\xi[m])$  as the sufficient statistics of the data set  $\mathcal{D}$ .

- For thumbtack, it is counts of M[1] and M[0]
- For multinomial, it is <M[1].... M[K]>
  - $-\tau(x^k)$  is k-dimensional vector with value 1 where  $x=x^k$ ;
  - $L(\mathbf{\theta} : D) = \prod_{k} \theta_{k}^{M[k]}$
  - Max is given by  $\theta_k^{\hat{}} = M[k]/M$  (easy to derive by taking derivative of the log of the L function above and setting equal to zero)
- For Gaussian distribution,  $\tau(x) = \langle 1, x, x^2 \rangle$ 
  - μ is given by average of sample values
  - $-\sigma$  given by square terms (subtract x terms)
  - Max of likelihood is given by empirical mean and variance

### MLE for Bayesian Networks

- Structure of BN allows estimation to reduce to a set of independent parameters.
- First consider network of two binary variables  $X \rightarrow Y$
- CPD: prior probability of X given by  $\theta_{x1}$  and  $\theta_{x0}$ Conditional probabilities  $\theta_{Y|X} = \theta_{Y|x1} \cup \theta_{Y|x0}$   $\theta_{Y|x1} = \{\theta_{y1|x1}, \theta_{y0|x1}\}; \theta_{Y|x0} = \{\theta_{y1|x0}, \theta_{y0|x0}\}$
- Each training instance is a tuple  $\langle x[m], y[m] \rangle$
- Likelihood function is:  $L(\theta : D) = \prod_{m=1}^{M} P(x[m], y[m] : \theta).$
- Factorize:  $L(\boldsymbol{\theta}:\mathcal{D}) = \prod_{m} P(x[m]:\boldsymbol{\theta})P(y[m] \mid x[m]:\boldsymbol{\theta}).$
- Distribute multiply:  $L(\theta : \mathcal{D}) = \left(\prod_{m} P(x[m] : \theta)\right) \left(\prod_{m} P(y[m] \mid x[m] : \theta)\right)$
- Note that first term is probability of a single variable so can be optimized as before (from counts of 1 and 0 values)

#### MLE for Bayesian Networks

Simplify second term (from previous slide)

$$\prod_{m} P(y[m] \mid x[m] : \theta_{Y|X}) 
= \prod_{m:x[m]=x^{0}} P(y[m] \mid x[m] : \theta_{Y|X}) \cdot \prod_{m:x[m]=x^{1}} P(y[m] \mid x[m] : \theta_{Y|X}) 
= \prod_{m:x[m]=x^{0}} P(y[m] \mid x[m] : \theta_{Y|x^{0}}) \cdot \prod_{m:x[m]=x^{1}} P(y[m] \mid x[m] : \theta_{Y|x^{1}}).$$

- Consider P( y[m] | x[m] :  $\theta_{Y|x0}$ ), if y[m]=y<sup>1</sup>, this term is  $\theta_{y1|x0}$  otherwise  $\theta_{y0|x0}$ .
- Let M[x0, y¹] be number of samples where  $X = x^0$ ,  $Y = y^1$ , etc then:  $\prod_{m:x[m]=x^0} P(y[m] \mid x[m]:\theta_{Y|x^0}) = \theta_{y1|x0} M[x0,y1] \cdot \theta_{y0|x0} M[x0,y0]$
- Maximizing will give us  $\theta_{y1|x0} = M[x^0, y^1] / M[x^0]$ 
  - Again, what we would have expected
  - Similar expressions for other terms

# General likelihood Decomposition

- Bayesian network G, parameters  $\theta$
- Given Dataset D =  $\{\xi[1], \xi[2], ..., \xi[M]\}$
- $L(\theta : D) = \prod_{m} P(\xi[m] : \theta)$  $= \prod_{m} \prod_{i} P(x_{i}[m] | pa_{Xi}[m] : \theta)$   $= \prod_{i} \{\prod_{m} P(x_{i}[m] | pa_{Xi}[m] : \theta)\} \text{ (reverse product order)}$   $= \prod_{i} L_{i}(\theta_{Xi|paXi} : D) \text{; } L_{i} \text{ is the local likelihood of } Xi \text{.}$   $L_{i}(\theta_{Xi|paXi} : D) = \prod_{m} P(x_{i}[m] | pa_{Xi}[m] : \theta_{Xi|paXi})$
- Likelihood decomposes; we can maximize each L<sub>i</sub> independently; Proposition 17.1

Let  $\mathcal{D}$  be a complete data set for  $X_1, \ldots, X_n$ , let  $\mathcal{G}$  be a network structure over these variables, and suppose that the parameters  $\theta_{X_i|\text{Pa}_{X_i}}$  are disjoint from  $\theta_{X_j|\text{Pa}_{X_j}}$  for all  $j \neq i$ . Let  $\hat{\theta}_{X_i|\text{Pa}_{X_i}}$  be the parameters that maximize  $L_i(\theta_{X_i|\text{Pa}_{X_i}}:\mathcal{D})$ . Then,  $\hat{\theta} = \langle \hat{\theta}_{X_1|\text{Pa}_1}, \ldots, \hat{\theta}_{X_n|\text{Pa}_n} \rangle$  maximizes  $L(\theta:\mathcal{D})$ .

#### Table CPDs

- Variable X with parents U, CPD is P(X|U), entries are  $\theta_{x|u}$
- $L_X(\boldsymbol{\theta}_{X|U}:D) = \prod_m \theta_{x[m]|\boldsymbol{u}[m]}$ =  $\prod_{\boldsymbol{u}} \prod_x \theta_{x|\boldsymbol{u}}^{M[\boldsymbol{u},x]}$
- Maximize for each  $\theta_{x|\mathbf{u}}$  independently, subject to  $\sum \theta_{x|\mathbf{u}} = 1$
- $\theta_{x|u}^{\ \ } = M[u,x]/\sum_{x} M[u,x] = M[u,x]/M[u];$
- Note: as data will be divided into several subsets, such as shown above, it may be hard to get enough samples for each subset, possibly resulting in overfitting.

#### Gaussian BNs

• Linear Gaussian CPDs of the form:

$$- P(X|\mathbf{u}) = N (\beta_0 + \beta_1 u_1 + \dots \beta_k u_k; \sigma^2)$$

- Task is to learn  $(\beta_0, \beta_1 \dots \beta_k, \sigma)$
- Can follow s derivation similar to that for discrete case; final result is as expected:
  - Estimate means of X and U
  - Estimate covariance matrix of {X} U U
  - Can solve for parameters from the above two (Thm 7.4)

#### Next Class

• Read sections 17.3 and 17.4