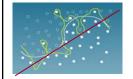
# **Machine Learning**

10-701/15-781, Spring 2008

#### **Model/Feature Selection**



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**Lecture 14, March 3, 2008** 

Reading: Chap. 1&2, CB & Chap 5,6, TM

# **Bias-variance decomposition**



• For one data set *D* and one test point *x* 

$$E_{(x,t),D}[(y(x)-t)^{2}]$$

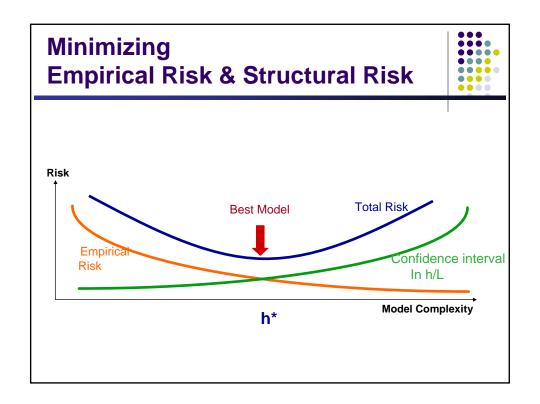
$$= \int (E_{D}[y(x;D)] - h(x))^{2} p(x) dx$$

$$+ \int E_{D}[(y(x;D) - E_{D}[y(x;D)])^{2}] p(x) dx$$

$$+ \int (h(x)-t)^{2} p(x,t) dx dt$$

- $\Rightarrow$  expected loss = (bias)<sup>2</sup> + variance + noise
- Recall the VC bound:

$$\epsilon(h) \le \hat{\epsilon}(h) + O\left(\sqrt{\frac{d}{m}\log\frac{m}{d} - \frac{1}{m}\log\delta}\right)$$



# **SRM & ERM in practice**



- There are many SRM-based strategies to build models:
- In the case of linear models

$$y = \langle w | x \rangle + b,$$

one wants to make ||w|| a controlled parameter: let us call  $H_{\rm C}$  the linear model function family satisfying the constraint:

Vapnik Major theorem:

When C decreases, d(H<sub>C</sub>) decreases

||x|| < R

# 9 =





# Regularization

- Maximum-likelihood estimates are not always the best (James and Stein showed a counter example in the early 60's)
- Alternative: we "regularize" the likelihood objective (also known as penalized likelihood, shrinkage, smoothing, etc.), by adding to it a penalty term:

$$\hat{\theta}_{\text{shrinkage}} = \arg\max_{\theta} \left[ l(\theta; D) + \lambda \|\theta\| \right]$$

where  $\lambda > 0$  and  $||\theta||$  might be the  $L_1$  or  $L_2$  norm.



- The choice of norm has an effect
  - using the  $L_2$  norm pulls directly towards the origin,
  - ullet while using the  $L_1$  norm pulls towards the coordinate axes, i.e it tries to set some of the coordinates to 0.
  - This second approach can be useful in a feature-selection setting.

# **Bayesian and Frequentist**



- Frequentist interpretation of probability
  - Probabilities are objective properties of the real world, and refer to limiting relative frequencies (e.g., number of times I have observed heads). Hence one cannot write P(Katrina could have been prevented|D), since the event will never repeat.
  - Parameters of models are *fixed, unknown constants*. Hence one cannot write  $P(\theta|D)$  since  $\theta$  does not have a probability distribution. Instead one can only write  $P(D|\theta)$ .
  - One computes point estimates of parameters using various *estimators*,  $\theta^* = f(D)$ , which are designed to have various desirable qualities when *averaged over future data D* (assumed to be drawn from the "true" distribution).
- Bayesian interpretation of probability
  - Probability describes degrees of belief, not limiting frequencies.
  - Parameters of models are *hidden variables*, so one can compute  $P(\theta|D)$  of  $P(f(\theta)|D)$  for some function f.
  - One estimates parameters by computing  $P(\theta|D)$  using Bayes rule:

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



# Bayesian interpretation of regulation



- Regularized Linear Regression
  - Recall that using squared error as the cost function results in the LMS estimate
  - And assume iid data and Gaussian noise, LMS is equivalent to MLE of  $\theta$

$$l(\theta) = n \log \frac{1}{\sqrt{2\pi\sigma}} - \frac{1}{\sigma^2} \frac{1}{2} \sum_{i=1}^n (y_i - \theta^T \mathbf{x}_i)^2$$

• Now assume that vector  $\theta$  follows a normal prior with 0-mean and a diagonal covariance matrix

$$\theta \sim N(\mathbf{0}, \tau^2 I)$$

• What is the posterior distribution of  $\theta$ ?

$$p(\theta|D) \propto p(D,\theta)$$

$$= p(D|\theta) p(\theta) = \left(2\pi\sigma^{2}\right)^{-n/2} \exp\left\{-\frac{1}{2\sigma^{2}} \sum_{i=1}^{n} \left(y_{n} - \theta^{T} x_{i}\right)^{2}\right\} \times C \exp\left\{-\left(\theta^{T} \theta / 2\tau^{2}\right)^{2}\right\}$$

# Bayesian interpretation of regulation, con'd



• The posterior distribution of  $\theta$ 

$$p(\theta|D) \propto \exp\left\{-\frac{1}{2\sigma^2} \sum_{i=1}^n \left(y_n - \theta^T x_i\right)^2\right\} \times \exp\left\{-\frac{\theta^T \theta}{2\sigma^2}\right\}$$

This leads to a now objective

$$l_{MAP}(\theta; D) = -\frac{1}{2\sigma^2} \frac{1}{2} \sum_{i=1}^{n} (y_i - \theta^T \mathbf{x}_i)^2 - \frac{1}{\tau^2} \frac{1}{2} \sum_{k=1}^{K} \theta_k^2$$
$$= l(\theta; D) + \lambda \|\theta\|$$

- This is  $L_2$  regularized LR! --- a MAP estimation of  $\theta$
- What about L<sub>1</sub> regularized LR! (homework)
- How to choose λ.
  - cross-validation!

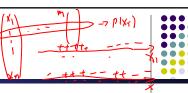
#### **Feature Selection**





- Imagine that you have a supervised learning problem where the number of features n is very large (perhaps n >>#samples), but you suspect that there is only a small number of features that are "relevant" to the learning task.
- VC-theory can tell you that this scenario is likely to lead to high generalization error – the learned model will potentially overfit unless the training set is fairly large.
- So lets get rid of useless parameters!

# How to score features

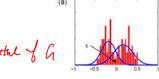


- How do you know which features can be pruned?
  - Given labeled data, we can compute some simple score S(i) that measures how informative each feature  $x_i$  is about the class labels y.
  - Ranking criteria:
    - Mutual Information: score each feature by its mutual information with respect to the class labels

 $MI(x_i, y) = \sum_{x_i \in \{0,1\}} \sum_{y \in \{0,1\}} p(x_i, y) \log \frac{p(x_i, y)}{p(x_i) p(y)}$ 



Bayes error:



(b) gene 1902

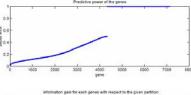


- Redundancy (Markov-blank score) ...
- We need estimate the relevant p()'s from data, e.g., using MLE

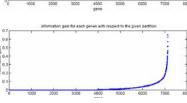
### **Feature Ranking**



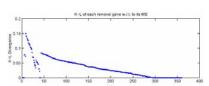
• Bayes error of each gene



 information gain for each genes with respect to the given partition



 KL of each removal gene w.r.t. to its MB



#### **Feature selection schemes**



- Given *n* features, there are 2<sup>n</sup> possible feature subsets (why?)
- Thus feature selection can be posed as a model selection problem over 2<sup>n</sup> possible models.
- For large values of n, it's usually too expensive to explicitly enumerate over and compare all  $2^n$  models. Some heuristic search procedure is used to find a good feature subset.
- Three general approaches:
  - Filter: i.e., direct feature ranking, but taking no consideration of the subsequent learning algorithm
    - add (from empty set) or remove (from the full set) features one by one based on S(i)
    - Cheap, but is subject to local optimality and may be unrobust under different classifiers
  - Wrapper: determine the (inclusion or removal of) features based on performance under the learning algorithms to be used. See next slide
  - Simultaneous learning and feature selection.
    - E.x. L₁ regularized LR, Bayesian feature selection (will not cover in this class), etc.

# Wrapper

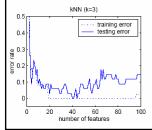


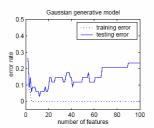
- Forward:
  - 1. Initialize  $\mathcal{F} = \emptyset$
  - 2. Repeat
    - For i = 1, ..., n
       if i ∉ F, let F₁ = F ∪ {i}, and use some version of cross validation to evaluate features F₁. (l.e., train your learning algorithm using only the features in F₁, and estimate its generalization error.)
    - Set  $\mathcal F$  to be the best feature subset found on the last step step.
  - 3. Select and output the best feature subset that was evaluated during the entire search procedure.
- Backward search
  - 1. Initialize  $\mathcal{F}$ = full set
  - 2. ..

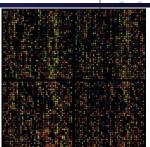
# Case study [Xing et al, 2001]

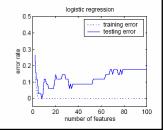


- The case:
  - 7130 genes from a microarray dataset
  - 72 samples
  - 47 type I Leukemias (called ALL)
     and 25 type II Leukemias (called AML)
- Three classifier:
  - kNN
  - Gaussian classifier
  - Logistic regression





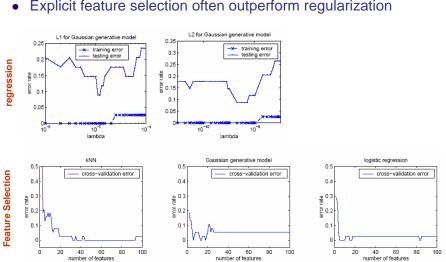




# Regularization vs. Feature **Selection**



• Explicit feature selection often outperform regularization



# **Model Selection**



- Suppose we are trying select among several different models for a learning problem.
- Examples:
  - 1. polynomial regression

$$h(x;\theta) = g(\theta_0 + \theta_1 x + \theta_2 x^2 + \dots + \theta_k x^k)$$

- Model selection: we wish to **automatically** and **objectively** decide if k should be, say, 0,
- 2. locally weighted regression,
- Model selection: we want to automatically choose the bandwidth parameter  $\ensuremath{ au}$ .
- 3. Mixture models and hidden Markov model,
- Model selection: we want to decide the number of hidden states
- The Problem:
  - $\bullet \quad \text{Given model family } \boldsymbol{\mathcal{F}} = \left\{ \boldsymbol{M}_1, \boldsymbol{M}_2, \dots, \boldsymbol{M}_I \right\}, \text{ find } \boldsymbol{M}_i \in \boldsymbol{\mathcal{F}} \quad \text{ s.t. }$  $M_i = \arg\max_{M \in \mathcal{F}} J(D, M)$

# **Model Selection via Information Criteria**



- How can we compare the closeness of a learned hypothesis and the true model?
- The relative entropy (also known as the <u>Kullback-Leibler</u> <u>divergence</u>) is a measure of how different two probability distributions (over the same event space) are.
  - For 2 pdfs, p(x) and q(x), their **KL-devergence** is:

$$D(p || q) = \sum_{x \in X} p(x) \log \frac{p(x)}{q(x)}$$

 The KL divergence between p and q can also be seen as the average number of bits that are wasted by encoding events from a distribution p with a code based on a not-quite-right distribution q.

#### An information criterion



- Let f(x) denote the truth, the underlying distribution of the data
- Let  $g(x, \theta)$  denote the model family we are evaluating
  - f(x) does not necessarily reside in the model family
  - $\theta_{ML}(y)$  denote the MLE of model parameter from data y
- Among early attempts to move beyond Fisher's Maliximum Likelihood framework, Akaike proposed the following information criterion:

$$E_{y} \Big[ D \Big( f \, \big\| \, g(x \, | \, \theta_{ML}(y) \Big) \Big]$$

which is, of course, intractable (because f(x) is unknown)

### **AIC and TIC**



• AIC (A information criterion, not Akaike information criterion)

$$A = \log g(x \mid \hat{\theta}(y)) - k$$

where k is the number of parameters in the model

• TIC (Takeuchi information criterion)

$$A = \log g(x \mid \hat{\theta}(y)) - \operatorname{tr}(I(\theta_0)\Sigma)$$

where

$$\theta_{0} = \arg\min D(f \parallel g(\cdot \mid \theta)) \qquad I(\theta_{0}) = -E_{x} \left[ \frac{\partial^{2} \log g(x \mid \theta)}{\partial \theta \partial \theta^{T}} \right] \bigg|_{\theta = \theta_{0}} \qquad \Sigma = E_{y} \left( \hat{\theta}(y) - \theta_{0} \right) \left( \hat{\theta}(y) - \theta_{0} \right)^{T} d\theta d\theta$$

- We can approximate these terms in various ways (e.g., using the bootstrap)
- $\operatorname{tr}(I(\theta_0)\Sigma) \approx k$

# **Bayesian Model Selection**



• Recall the Bayesian Theory: (e.g., for date *D* and model *M*)

$$P(M|D) = P(D|M)P(M)/P(D)$$

- the posterior equals to the likelihood times the prior, up to a constant.
- Assume that P(M) is uniform and notice that P(D) is constant, we have the following criteria:

$$P(D \mid M) = \int_{\theta} P(D \mid \theta, M) P(\theta \mid M) d\theta$$

 A few steps of approximations (you will see this in advanced ML class in later semesters) give you this:

$$P(D \mid M) \approx \log P(D \mid \hat{\theta}_{ML}) - \frac{k}{2} \log N$$

where N is the number of data points in D.

# **Summary**



- Bias-variance decomposition
- The battle against overfitting:
  - Cross validation
  - Regularization
  - Model selection --- Occam's razor
  - Model averaging
    - The Bayesian-frequentist debate
    - Bayesian learning (weight models by their posterior probabilities)

Review								
Method	Input.	Output	Loss	hypothesis	Op+ procedure	Discriminative		
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