## Lecture 9: Learning Theory

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January 5, 2022

## Why Learning Theory?

- 2 Bias, Variance and Model Complexity
- 3 Bias-Variance Decomposition
- 4 The Gap Between Training Error and Generalization Error
- 5 Selecting Right Model and Features

- How can we tell if your learning algorithm will do a good job in future (test time)?
  - Experimental results
  - Theoretical analysis
- Why theory?
  - Can only run a limited number of experiments..
  - Experiments rarely tell us what will go wrong
- Using learning theory, we can make formal statements/give guarantees on
  - Expected performance ("generalization") of a learning algorithm on test data
  - Number of examples required to attain a certain level of test accuracy
  - Hardness of learning problems in general

• Bias is a learner's tendency to consistently learn the same wrong thing

- The bias is error from erroneous assumptions in the learning algorithm
- High bias can cause an algorithm to miss the relevant relations between features and target outputs (underfitting)
- Variance is the tendency to learn random things irrespective of the real signal
  - The variance is error from sensitivity to small fluctuations in the training set
  - High variance can cause an algorithm to model the random noise in the training data, rather than the intended outputs (overfitting)

- The loss function for measuring errors between Y and  $\hat{f}(X)$

$$L(Y, \hat{f}(X)) = egin{cases} (Y - \hat{f}(X))^2, & ext{squared error} \ |Y - \hat{f}(X)|, & ext{absolute error} \end{cases}$$

• Test error (or generalization error)

$$\operatorname{Err}_{\mathcal{D}} = \mathbb{E}[L(Y, \hat{f}(X)) \mid \mathcal{D}]$$

• Expected prediction (or test) error

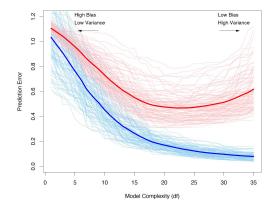
$$\operatorname{Err} = \mathbb{E}[L(Y, \hat{f}(X))] = \mathbb{E}[\operatorname{Err}_{\mathcal{D}}]$$

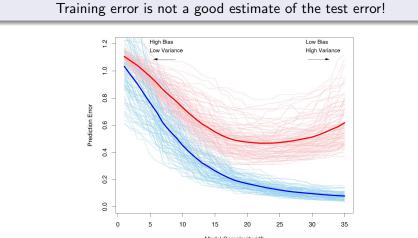
which averages over everything that is random, including the randomness in the training set that produced  $\hat{f}$ 

Training error

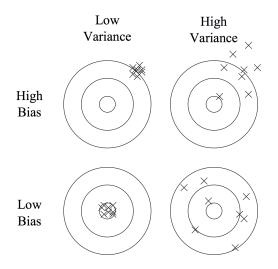
$$\overline{\operatorname{err}} = \frac{1}{m} \sum_{i=1}^{m} L(y_i, \hat{f}(x_i))$$

Behavior of test sample and training sample error as the model complexity is varied. The light blue curves show the training error  $\overline{\text{Err}}$ , while the light red curves show the conditional test error  $\overline{\text{Err}}_{\mathcal{D}}$  for 100 training sets of size 50 each, as the model complexity is increased. The solid curves show the expected test error  $\overline{\text{Err}}$  and the expected training error  $\mathbb{E}[\overline{\text{Err}}]$ .

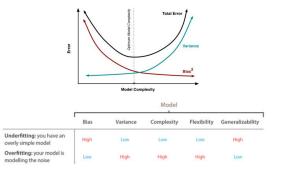




Model Complexity (df)

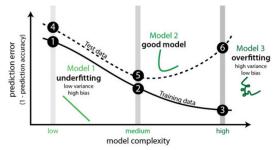


• Simple model have high bias and small variance, complex models have small bias and high variance



 If you modified a model to reduce its bias (e.g., by increasing the model's complexity), you are likely to increase its variance, and viceversa (if, however, both increase then you might be doing it wrong!)

- The bad performance (low accuracy on test data) could be due to either high bias (underfitting) or high variance (overfitting)
- Looking at the training and test error can tell which of the two is the case



- High bias: Both training and test error are large
- High variance: Small training error, large test error (and huge gap)

- **Model section**: Estimating the performance of different models in order to choose the best one
- Model assessment: Having chosen a final model, estimating its prediction error (generalization error) on new data.
- If we are in a data-rich situation, the best approach for both problems is to randomly divide the dataset into three parts: a *training* set, a *validation* set, and a *test* set.
  - The training set is used to fit the models
  - The validation set is used to estimate prediction error for model selection
  - The test set is used for assessment of the generalization error of the final chosen model
- A typical split might be 50% for training, and 25% each for validation and testing

• For a model 
$$Y = f(X) + \epsilon$$
 with  $\mathbb{E}(\epsilon) = 0$  and  $Var(\epsilon) = \sigma_{\epsilon}^2$ 

$$\begin{aligned} \mathsf{Err}(x_0) &= \mathbb{E}[(y_0 - \hat{f}(x_0))^2] \\ &= \mathbb{E}[y_0^2 - 2y_0\hat{f}(x_0) + \hat{f}^2(x_0)] \\ &= \mathbb{E}[y_0^2] + \mathbb{E}[\hat{f}^2(x_0)] - \mathbb{E}[2y_0\hat{f}(x_0)] \\ &= \mathsf{Var}[y_0] + \mathbb{E}^2[y_0] + \mathsf{Var}[\hat{f}(x_0)] + \mathbb{E}^2[\hat{f}(x_0)] - \mathbb{E}[2y_0\hat{f}(x_0)] \\ &= \mathsf{Var}[f(x_0) + \epsilon] + \mathbb{E}^2[f(x_0) + \epsilon] + \mathsf{Var}[\hat{f}(x_0)] \\ &\quad + \mathbb{E}^2[\hat{f}(x_0)] - \mathbb{E}[2(f(x_0) + \epsilon)\hat{f}(x_0)] \\ &= \sigma_{\epsilon}^2 + f^2(x_0) + \mathsf{Var}[\hat{f}(x_0)] + \mathbb{E}^2[\hat{f}(x_0)] - 2f(x_0)\mathbb{E}[\hat{f}(x_0)] \\ &= \sigma_{\epsilon}^2 + (f(x_0) - \mathbb{E}[\hat{f}(x_0)])^2 + \mathsf{Var}[\hat{f}(x_0)] \\ &= \mathsf{Irreducible Error} + \mathsf{Bias}^2 + \mathsf{Variance} \end{aligned}$$

## Bias-Variance Decomposition (Contd.)

• For a model  $Y = f(X) + \epsilon$  with  $\mathbb{E}(\epsilon) = 0$  and  $Var(\epsilon) = \sigma_{\epsilon}^2$ 

$$\begin{aligned} \mathsf{Err}(x_0) &= \mathbb{E}[(Y - \hat{f}(x_0))^2] \\ &= \sigma_{\epsilon}^2 + (f(x_0) - E[\hat{f}(x_0)])^2 + \mathsf{Var}[\hat{f}(x_0)] \\ &= \mathsf{Irreducible \ Error} + \mathsf{Bias}^2 + \mathsf{Variance} \end{aligned}$$

- The first term is the variance of the target around its true mean  $f(x_0)$ , and cannot be avoided no matter how well we estimate  $f(x_0)$ , unless  $\sigma_{\epsilon}^2 = 0$
- The second term is the squared bias, i.e., the amount by which the average of our estimate differs from the true mean
- The last term is the variance, i.e., the expected squared deviation of  $\hat{f}(x_0)$  around its mean

• For k-nearest neighbor regression,

$$\operatorname{Err}(x_0) = \mathbb{E}[(Y - \hat{f}_k(x_0))^2]$$
  
=  $\sigma_{\epsilon}^2 + \left[f(x_0) - \frac{1}{k}\sum_{\ell=1}^k f(x_{(\ell)})\right]^2 + \frac{\sigma_{\epsilon}^2}{k}$ 

• For linear regression model  $Y = X\theta + \epsilon$ ,

$$\begin{aligned} \mathsf{Bias}(x_0) &= f(x_0) - \mathbb{E}[\hat{f}(x_0)] \\ &= x_0^T \theta - \mathbb{E}[x_0^T \hat{\theta}] \\ &= x_0^T \theta - \mathbb{E}[x_0^T (X^T X)^{-1} X^T Y] \\ &= x_0^T \theta - \mathbb{E}[x_0^T (X^T X)^{-1} X^T (X \theta + \epsilon)] \\ &= x_0^T \theta - \mathbb{E}[x_0^T (X^T X)^{-1} X^T X \theta + x_0^T (X^T X)^{-1} X^T \epsilon] \\ &= x_0^T \theta - \mathbb{E}[x_0^T \theta + x_0^T (X^T X)^{-1} X^T \epsilon] \\ &= \mathbb{E}[x_0^T \theta - x_0^T \theta + x_0^T (X^T X)^{-1} X^T \epsilon] \\ &= 0 \end{aligned}$$

## Bias-Variance Decomposition (Contd.)

• For linear regression model  $Y = X\theta + \epsilon$ ,

$$\begin{aligned} \mathsf{Var}(\hat{f}(x_{0})) &= & \mathbb{E}[(f(x_{0}) - \mathbb{E}[\hat{f}(x_{0}))^{2}] \\ &= & \mathbb{E}[(x_{0}^{T}(X^{T}X)^{-1}X^{T}Y - X_{0}^{T}\theta)^{2}] \\ &= & \mathbb{E}[(x_{0}^{T}(X^{T}X)^{-1}X^{T}(X\theta + \epsilon) - X_{0}^{T}\theta)^{2}] \\ &= & \mathbb{E}[(x_{0}^{T}(X^{T}X)^{-1}X^{T}\epsilon)^{2}] \\ &= & \mathbb{E}[(x_{0}^{T}(X^{T}X)^{-1}X^{T}\epsilon)(x_{0}^{T}(X^{T}X)^{-1}X^{T}\epsilon)^{T}] \\ &= & \mathbb{E}[x_{0}^{T}(X^{T}X)^{-1}X^{T}\epsilon\epsilon^{T}(x_{0}^{T}(X^{T}X)^{-1}X^{T})^{T}] \\ &= & x_{0}^{T}(X^{T}X)^{-1}X^{T}\mathbb{E}[\epsilon\epsilon^{T}](x_{0}^{T}(X^{T}X)^{-1}X^{T})^{T} \\ &= & x_{0}^{T}(X^{T}X)^{-1}X^{T}\sigma_{\epsilon}^{2}I(x_{0}^{T}(X^{T}X)^{-1}X^{T})^{T} \\ &= & \sigma_{\epsilon}^{2}x_{0}^{T}(X^{T}X)^{-1}X^{T}(x_{0}^{T}(X^{T}X)^{-1}X^{T})^{T} \\ &= & \sigma_{\epsilon}^{2}x_{0}^{T}(X^{T}X)^{-1}x_{0} \\ &\approx & \sigma_{\epsilon}^{2}\frac{n}{m} \end{aligned}$$

#### The union bound

Assume  $A_1, A_2, \dots, A_k$  be k different events (that may not be independent),

$$p(A_1 \bigcup A_2 \cdots \bigcup A_k) \leq p(A_1) + \cdots + p(A_k)$$

## • Hoeffding inequality (Chernoff bound)

Let  $Z_1, \dots, Z_m$  be *m* independent and identically distributed (iid) random variables drawn from a Bernoulli( $\phi$ ) distribution (i.e.,  $p(Z_i = 1) = \phi$  and  $p(Z_i = 0) = 1 - \phi$ ). Let  $\hat{\phi} = \frac{1}{m} \sum_{i=1}^{m} Z_i$  be the mean of these random variables, and let any  $\gamma > 0$  be fixed. Then

$$p(|\phi - \hat{\phi}| > \gamma) \le 2 \exp(-2\gamma^2 m)$$

- $\bullet$  A hypothesis class  $\mathcal{H}:$  a set of all classifiers considered by a learning algorithm
- A training set  $S = \{(x^{(i)}, y^{(i)})\}_{i=1, \cdots, m}$  with  $y^{(i)} \in \{0, 1\}$  are drawn i.i.d. from some probability distribution D
- The learning algorithm, given training data, learns a hypothesis  $h \in \mathcal{H}$

• The training error (or empirical risk, empirical error) is

$$\overline{\operatorname{Err}}(h) = \frac{1}{m} \sum_{i=1}^{m} \mathbf{1}\{h(x^{(i)}) \neq y^{(i)}\}$$

i.e., the fraction of the misclassified training examples

• The generalization is

$$\operatorname{Err}_{\mathcal{D}}(h) = \mathbb{P}_{(x,y)\sim\mathcal{D}}(h(x)\neq y)$$

i.e., the probability that, if we now draw a new example (x, y) from the distribution  $\mathcal{D}$ , h will misclassify it

• Empirical Risk Minimization (ERM)

- Consider a linear classification  $h_{\theta}(x) = \mathbf{1}(\theta^T x \ge 0)$
- Minimize the training error

$$heta^* = rg\min_{ heta} \overline{\operatorname{Err}}(h_{ heta})$$

• Optimal hypothesis  $h^* = h_{ heta^*}$ 

• ERM can also be thought of a minimization over the class

$$h^* = \arg\min_{h \in \mathcal{H}} \overline{\operatorname{Err}}(h)$$

- A finite hypothesis class  $\mathcal{H} = \{h_1, \cdots, h_k\}$
- *h*<sup>\*</sup> ∈ *H* denotes the optimal hypothesis function with the training error minimized by ERM
- Does there exist a guarantee on the generalization error of  $\hat{h}$ ?
  - $\operatorname{Err}_{\mathcal{D}}(h)$  is a reliable estimate of  $\operatorname{Err}(h)$  for  $\forall h$
  - This implies an upper-bound on the generalization error of  $h^*$

# Finite $\mathcal{H}$ (Contd.)

- Assume  $(x, y) \sim \mathcal{D}$
- For  $h_i \in \mathcal{H}$ , define Bernoulli random variables

$$Z = \mathbf{1}(h_i(x) \neq y)$$
  
 $Z_j = \mathbf{1}\{h_i(x^{(j)}) \neq y^{(j)}\}$ 

• The generalization error

$$\operatorname{Err}_{\mathcal{D}}(h_i) = \mathbb{E}[Z] = \mathbb{E}[Z_j]$$

• The training error

$$\overline{\mathsf{Err}}(h_i) = \frac{1}{m} \sum_{j=1}^m Z_j$$

# Finite $\mathcal{H}$ (Contd.)

- Assume  $(x, y) \sim \mathcal{D}$
- For  $h_i \in \mathcal{H}$ , define Bernoulli random variables

$$Z = \mathbf{1}(h_i(x) \neq y)$$
  
 $Z_j = \mathbf{1}\{h_i(x^{(j)}) \neq y^{(j)}\}$ 

- The generalization error  $\operatorname{Err}_{\mathcal{D}}(h_i) = \mathbb{E}[Z] = \mathbb{E}[Z_j]$
- The training error  $\overline{\operatorname{Err}}(h_i) = \frac{1}{m} \sum_{j=1}^m Z_j$
- By applying Hoeffding inequality, we have

$$P(|\overline{\operatorname{Err}}(h_i) - \operatorname{Err}_{\mathcal{D}}(h_i)| > \gamma) \le 2 \exp(-2\gamma^2 m)$$

• For a particular  $h_i$ , training error will be close to generalization error with high probability, assuming m is large

• Let  $A_i$  denote the event that  $|\overline{\operatorname{Err}}(h_i) - \operatorname{Err}_{\mathcal{D}}(h_i)| > \gamma$ , then

$$\mathbb{P}(A_i) \leq 2\exp(-2\gamma^2 m)$$

• By using the union bound, we have

$$\mathbb{P}(|\overline{\operatorname{Err}}(h_i) - \operatorname{Err}_{\mathcal{D}}(h_i)| > \gamma)$$

$$= \mathbb{P}(A_1 \bigcup \cdots \bigcup A_k) \le \sum_{i=1}^k P(A_i)$$

$$\le \sum_{i=1}^k 2 \exp(-2\gamma^2 m) = 2k \exp(-2\gamma^2 m)$$

• Then, we have the following result

$$\mathbb{P}(\neg \exists h \in \mathcal{H} : |\overline{\mathsf{Err}}(h) - \mathsf{Err}_{\mathcal{D}}(h)| > \gamma)$$
  
=  $\mathbb{P}(\forall h \in \mathcal{H} : |\overline{\mathsf{Err}}(h) - \mathsf{Err}_{\mathcal{D}}(h)| > \gamma)$   
≥  $1 - 2k \exp(-2\gamma^2 m)$ 

• With probability at least  $1 - 2k \exp(-2\gamma^2 m)$ , we have

$$|\overline{\mathsf{Err}}(h) - \mathsf{Err}_{\mathcal{D}}(h)| \leq \gamma$$

for  $\forall h \in \mathcal{H}$ 

Given  $\gamma$  and  $\delta > 0$ , how large should *m* be such that we can guarantee

 $|\overline{\operatorname{Err}}(h) - \operatorname{Err}_{\mathcal{D}}(h)| \leq \gamma$ 

with probability  $\geq 1 - \delta$ ?

Solution

$$1-2k\exp(-2\gamma^2m)\geq 1-\delta \Rightarrow m\geq rac{1}{2\gamma^2}\lograc{2k}{\delta}$$

- The training set size *m* that a certain method or algorithm requires in order to achieve a certain level of performance is so-called the algorithm's sample complexity
- The number of training examples needed to make this guarantee is only logarithmic in the number of hypotheses in  $\mathcal{H}$  (i.e., k)

• Fixing *m* and  $\delta$ , solving for  $\gamma$  gives

$$1 - 2k \exp(-2\gamma^2 m) \ge 1 - \delta \Rightarrow |\overline{\mathsf{Err}}(h) - \mathsf{Err}_{\mathcal{D}}(h)| \le \sqrt{\frac{1}{2m} \log \frac{2k}{\delta}}$$

Given m and  $\delta > 0$ , with probability at least  $1 - \delta$ ,

$$|\overline{\mathsf{Err}}(h) - \mathsf{Err}_{\mathcal{D}}(h)| \leq \sqrt{rac{1}{2m}\lograc{2k}{\delta}}$$

• Assume 
$$\hat{h} = \arg\min_{h \in \mathcal{H}} \operatorname{Err}_{\mathcal{D}}(h)$$

$$\begin{aligned} \mathsf{Err}_{\mathcal{D}}(h^*) &\leq \quad \overline{\mathsf{Err}}(h^*) + \gamma \\ &\leq \quad \overline{\mathsf{Err}}(\hat{h}) + \gamma \\ &\leq \quad \mathsf{Err}_{\mathcal{D}}(\hat{h}) + 2\gamma \end{aligned}$$

• If uniform convergence occurs, then the generalization error of  $h^*$  is at most  $2\gamma$  worse than the best possible hypothesis in  $\mathcal{H}$ 

#### Theorem

Let  $|\mathcal{H}| = k$  and let any m and  $\delta$  be fixed. With probability at least  $1 - \delta$ , we have

$$\operatorname{Err}_{\mathcal{D}}(h^*) \leq \left(\min_{h \in \mathcal{H}} \operatorname{Err}_{\mathcal{D}}(h)\right) + 2\sqrt{\frac{1}{2m}\log \frac{2k}{\delta}}$$

- $\bullet$  If we take a larger hypothesis set  $\mathcal{H}'$  such that  $\mathcal{H}\subseteq \mathcal{H}'$ 
  - the first term is decreased (the bias is decreased)
  - the second term is increased (the variance is increased)

### Corollary

Let  $|\mathcal{H}| = k$  and let any  $\delta$ ,  $\gamma$  be fixed. For

$$\mathsf{Err}_{\mathcal{D}}(h^*) \leq \min_{h \in \mathcal{H}} \mathsf{Err}_{\mathcal{D}}(h) + 2\gamma$$

to hold with probability at least  $1 - \delta$ , it suffices that

$$egin{array}{rcl} m & \geq & rac{1}{2\gamma^2}\lograc{2k}{\delta} \ & = & O(rac{1}{\gamma^2}\lograc{k}{\delta}) \end{array}$$

- What happens when the hypothesis class size  $|\mathcal{H}|$  is infinite?
  - Example: The set of all linear classifiers
- The above bound does not apply (it just becomes trivial)
- $\bullet$  We need some other way of measuring the size of  ${\cal H}$ 
  - $\bullet\,$  One way: use the complexity  ${\cal H}$  as a measure of its size
  - Vapnik-Chervonenkis dimension (VC dimension)
  - VC dimension: a measure of the complexity of a hypothesis class

## Shattering

A set of points (in a given configuration) is shattered by a hypothesis class *H*, if, no mater how the points are labeled, there exists some *h* ∈ *H* that can separate the points

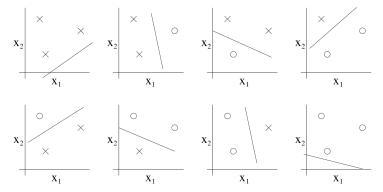


Figure: 3 points in 2D (locations fixed, only labeling varies),  $\mathcal{H}_{:}$  set of linear classifier

### Definition (VC Dimension)

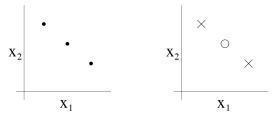
Given a hypothesis class  $\mathcal{H}$ , we then define its Vapnik-Chervonenkis dimension, VC( $\mathcal{H}$ ), to be the size of the largest set that is shattered by  $\mathcal{H}$ 

• Consider the following shattering game between us and an adversary

- We choose d points in an input space, positioned however we want
- Adversary labels these d points
- We define a hypothesis  $h \in \mathcal{H}$  that separates the points
- Note: Shattering just one configuration of d points is enough to win
- The VC dimension of  $\mathcal{H}$ , in that input space, is the maximum d we can choose so that we always succeed in the game

## VC Dimension (Contd.)

• Even when VC(H) = 3, there exist sets of size 3 that cannot be classified correctly



 In order words, under the definition of the VC dimension, in order to prove that VC(H) is at least d, we need to show only that there's at least one set of size d that H can shatter. • A measure of the "power" or the "complexity" of the hypothesis space

- Higher VC dimension implies a more "expressive" hypothesis space)
- Shattering: A set of N points is shattered if there exists a hypothesis that is consistent with every classification of the N points
- VC Dimension: The maximum number of data points that can be "shattered"
- If VC Dimension = d, then:
  - There exists a set of *d* points that can be shattered
  - There does not exist a set of d + 1 points that can be shattered

#### Theorem

Let  $\mathcal{H}$  be given, and let  $d = VC(\mathcal{H})$ . Then, with probability at least  $1 - \delta$ , we have that for all  $h \in \mathcal{H}$ 

$$|\mathsf{Err}_{\mathcal{D}}(h) - \overline{\mathsf{Err}}(h)| \le O\left(\sqrt{rac{d}{m}\lograc{m}{d} + rac{1}{m}\lograc{1}{\delta}}
ight)$$

and thus

$$\operatorname{Err}_{\mathcal{D}}(h^*) \leq \overline{\operatorname{Err}}(\hat{h}) + O\left(\sqrt{\frac{d}{m}\log{\frac{m}{d}} + \frac{1}{m}\log{\frac{1}{\delta}}}\right)$$

• Recall for finite hypothesis space

$$\operatorname{\mathsf{Err}}_{\mathcal{D}}(h^*) \leq \left(\min_{h \in \mathcal{H}} \operatorname{\mathsf{Err}}_{\mathcal{D}}(h)\right) + 2\sqrt{\frac{1}{2m}\log \frac{2k}{\delta}}$$

• VC(H) is like a substitute for  $k = |\mathcal{H}|$ 

- Given a set of models  $M = \{M_1, M_2, ..., M_R\}$ , choose the model that is expected to do the best on the test data
- *M* may consist of:
  - Same learning model with different complexities or hyperparameters
    - Nonlinear Regression: Polynomials with different degrees
    - K-Nearest Neighbors: Different choices of K
    - Decision Trees: Different choices of the number of levels/leaves
    - SVM: Different choices of the misclassification penalty parameter C
    - Regularized Models: Different choices of the regularization parameter
    - Kernel based Methods: Different choices of kernels
    - ... and almost any learning problem
  - Different learning models (e.g., SVM, KNN, DT, etc.)

- Given a training set S, do the following
  - Randomly split S into  $S_{train}$  (say, 70% of the data) and  $S_{cv}$  (the remaining 30% called the hold-out cross validation set)
  - Train each model  $M_i$  on  $S_{train}$  only, to get some hypothesis  $h_i$ .
  - Select and output the hypothesis  $h_i$  that had the smallest error  $\overline{\operatorname{Err}}_{S_{cv}}(h_i)$  on the hold-out cross validation set
- Option: After selecting  $M^* \in \mathcal{M}$  such that  $h^* = \arg \min_i \overline{\operatorname{Err}}_{S_{cv}}(h_i)$ , retrain  $M^*$  on the entire training set S
- Weakness: It seems we are trying to select the best model based on only part of the training set

- Randomly split S into k disjoint subsets  $S_1, \dots, S_k$ , each of which involves m/k training examples
- For each model  $M_i$ , we evaluate it as follows:
  - For j = 1, · · · , k, train the model M<sub>i</sub> on S<sub>1</sub> ∪ · · · ∪ S<sub>j-1</sub> ∪ S<sub>j+1</sub> ∪ · · · ∪ S<sub>k</sub> (i.e., train on all the data except S<sub>j</sub>) to get some hypothesis h<sub>ij</sub>, and then test the hypothesis h<sub>ij</sub> on S<sub>j</sub>, to get Err<sub>Si</sub>(h<sub>ij</sub>).
  - The estimated generalization error of model  $M_i$  is then calculated as the average of the  $\overline{\operatorname{Err}}_{S_i}(h_{ij})$ 's (averaged over j).
- Pick the model  $M_i$  with the lowest estimated generalization error, and retrain that model on the entire training set S

- Given *n* features resulting in 2<sup>*n*</sup> possible feature subsets, which one is the optimal?
- Forward search:
  - Initialize  $\mathcal{F} = \emptyset$
  - Until  $|\mathcal{F}| = \epsilon$  or  $|\mathcal{F}| = n$ , repeat
    - (a) For i = 1, · · · , n, if i ∉ F, let F<sub>i</sub> = F ∪{i}, and use cross validation to evaluate F<sub>i</sub>
    - (b) Set  ${\mathcal F}$  to be the best feature subset found in (a)
- Backward search: Start with  $\mathcal{F} = \{1, \cdots, n\}$ , and repeatedly deletes features one at a time until  $|\mathcal{F}| = \epsilon$
- The above two methods are so-called wrapper model, which is a procedure that "wraps" around your learning algorithm
- Wrapper feature selection algorithms usually have considerable computational cost
  - $O(n^2)$  calls to the learning algorithm

- Heuristic but computationally efficient
- Basic idea: Compute a score S(i) to measure how informative each feature  $x_i$  is about the class labels y; then, select the k features with the largest scores S(i)
- Mutual information  $MI(x_i, y)$  between  $x_i$  and y

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

with  $p(x_i, y)$ ,  $p(x_i)$  and p(y) estimated according their empirical distributions on the training set

- How to choose a right k?
  - Use cross validation

# Thanks!

Q & A