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#### **Announcements**

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- So far, we've been focused only on algorithms for finding the best hypothesis in the hypothesis space
  - How do we know that the learned hypothesis will perform well on the test set?
  - How many samples do we need to make sure that we learn a good hypothesis?
  - In what situations is learning possible?



- If the training data was linearly separable, we saw that perceptron/SVMs will always perfectly classify the training data
  - This does not mean that it will perfectly classify the test data
  - Intuitively, if the true distribution of samples is linearly separable, then seeing more data should help us do better



## **Problem Complexity**

- Complexity of a learning problem depends on
  - Size/expressiveness of the hypothesis space
  - Accuracy to which a target concept must be approximated
  - Probability with which the learner must produce a successful hypothesis
  - Manner in which training examples are presented, e.g. randomly or by query to an oracle



#### **Problem Complexity**

- Measures of complexity
  - Sample complexity
    - How much data you need in order to (with high probability)
       learn a good hypothesis
  - Computational complexity
    - Amount of time and space required to accurately solve (with high probability) the learning problem
    - Higher sample complexity means higher computational complexity



- Probably approximately correct (PAC)
  - Developed by Leslie Valiant
  - The only reasonable expectation of a learner is that with high probability it learns a close approximation to the target concept
  - Specify two small parameters,  $\epsilon$  and  $\delta$ , and require that with probability at least  $(1-\delta)$  a system learn a concept with error at most  $\epsilon$



#### **Consistent Learners**

- Imagine a simple setting
  - The hypothesis space is finite (i.e., |H| = c)
  - The true distribution of the data is  $p(\vec{x})$ , no noisy labels
  - We learned a perfect classifier on the training set, let's call it  $h \in H$ 
    - A learner is said to be consistent if it always outputs a perfect classifier on the training data assuming that one exists
  - Want to compute the error of the classifier



#### **Notions of Error**

- Training error of  $h \in H$ 
  - The error on the training data
  - Number of samples incorrectly classified divided by the total number of samples
- True error of  $h \in H$ 
  - The error over all possible future random samples
  - Probability that h misclassifies a random data point

$$p(h(x) \neq y)$$



- Let  $(x^{(1)}, y_1), \dots, (x^{(m)}, y_m)$  be m labelled data points sampled independently according to p
- Let  $C_i^h$  be a random variable that indicates whether or not the  $i^{th}$  data point is correctly classified
- The probability that h misclassifies the  $i^{th}$  data point is

$$p(C_i^h = 0) = \sum_{(x,y)} p(x,y) \, 1_{h(x) \neq y} = \epsilon_h$$



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This is the true error of *h* 



Probability that all data points classified correctly?

$$p(C_1^h = 1, ..., C_m^h = 1) = \prod_{i=1}^m p(C_i^h = 1) = (1 - \epsilon_h)^m$$

• Probability that a hypothesis  $h \in H$  whose true error is at least  $\epsilon$  correctly classifies the m data points is then

$$p(C_1^h = 1, ..., C_m^h = 1) \le (1 - \epsilon)^m \le e^{-\epsilon m}$$

for 
$$\epsilon \leq 1$$



- The version space (set of consistent hypotheses) is said to be  $\epsilon$ -exhausted if and only if every consistent hypothesis has true error less than  $\epsilon$ 
  - Enough samples to guarantee that every consistent hypothesis has error at most  $\epsilon$
- We'll show that w.h.p. every hypothesis with true error at least  $\epsilon$  is not consistent with the data



#### The Union Bound

- Let  $H_{BAD} \subseteq H$  be the set of all hypotheses that have true error at least  $\epsilon$
- From before for each  $h \in H_{BAD}$ ,

 $p(h \text{ correctly classifies all } m \text{ data points}) \leq e^{-\epsilon m}$ 

• So, the probability that *some*  $h \in H_{BAD}$  correctly classifies all of the data points is

$$p\left(\bigvee_{h\in H_{BAD}}\left(C_{1}^{h}=1,\ldots,C_{m}^{h}=1\right)\right)\leq\sum_{h\in H_{BAD}}p\left(C_{1}^{h}=1,\ldots,C_{m}^{h}=1\right)$$

$$\leq\left|H_{BAD}\right|e^{-\epsilon m}$$

$$\leq\left|H\right|e^{-\epsilon m}$$



### Haussler, 1988

What we just proved:

**Theorem:** For a finite hypothesis space, H, with m i.i.d. samples, and  $0 < \epsilon < 1$ , the probability that the version space is not  $\epsilon$ -exhausted is at most  $|H|e^{-\epsilon m}$ 

We can turn this into a sample complexity bound



## **Sample Complexity**

- Let  $\delta$  be an upper bound on the desired probability of not  $\epsilon$ -exhausting the sample space
  - The probability that the version space is not  $\epsilon$ exhausted is at most  $|H|e^{-\epsilon m} \leq \delta$
  - Solving for m yields

$$m \ge -\frac{1}{\epsilon} \log \frac{\delta}{|H|}$$
$$= \left(\log |H| + \log \frac{1}{\delta}\right) / \epsilon$$



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This is sufficient, but not necessary (union bound is quite loose)



#### **Decision Trees**

- Suppose that we want to learn an arbitrary Boolean function given n Boolean features
- Hypothesis space consists of all decision trees
  - Size of this space = ?
- How many samples are sufficient?



#### **Decision Trees**

- Suppose that we want to learn an arbitrary Boolean function given n Boolean features
- Hypothesis space consists of all decision trees
  - Size of this space =  $2^{2^n}$  = number of Boolean functions on n inputs
- How many samples are sufficient?

$$m \ge \left(\log 2^{2^n} + \log \frac{1}{\delta}\right)/\epsilon$$



#### Generalizations

- How do we handle the case the there is no perfect classifier?
  - Pick the hypothesis with the lowest error on the training set
- What do we do if the hypothesis space isn't finite?
  - Infinite sample complexity?
  - Next time...



#### **Chernoff Bounds**

• Chernoff bound: Suppose  $Y_1, \ldots, Y_m$  are i.i.d. random variables taking values in  $\{0,1\}$  such that  $E_p[Y_i]=y$ . For  $\epsilon>0$ ,

$$p\left(y - \frac{1}{m}\sum_{i} Y_{i} \ge \epsilon\right) \le e^{-2m\epsilon^{2}}$$

• Applying this to  $1-C_1^h$ , ...,  $1-C_m^h$  gives

$$p\left(\epsilon_h - \frac{1}{m}\sum_{i}(1 - C_i^h) \ge \epsilon\right) \le e^{-2m\epsilon^2}$$



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#### **PAC Bounds**

- **Theorem:** For a finite hypothesis space H finite, m i.i.d. samples, and  $0 < \epsilon < 1$ , the probability that true error of any of the best classifiers (i.e., lowest training error) is larger than its training error plus  $\epsilon$  is at most  $|H|e^{-2m\epsilon^2}$ 
  - Sample complexity (for desired  $\delta \geq |H|e^{-2m\epsilon^2}$ )

$$m \ge \left(\log|H| + \log\frac{1}{\delta}\right)/2\epsilon^2$$



#### **PAC Bounds**

• If we require that the previous error is bounded above by  $\delta$ , then with probability  $(1 - \delta)$ , for all  $h \in H$ 

$$\epsilon_h \leq \epsilon_h^{train} + \sqrt{\frac{1}{2m} \left( \log |H| + \log \frac{1}{\delta} \right)}$$
 "bias" "variance"

- For small |H|
  - High bias (may not be enough hypotheses to choose from)
  - Low variance



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 "bias" "variance"

- For large |H|
  - Low bias (lots of good hypotheses)
  - High variance



- Given:
  - Set of data X
  - Hypothesis space H
  - Set of target concepts C
  - Training instances from unknown probability distribution over X of the form (x, c(x))
- Goal:
  - Learn the target concept  $c \in C$



- Given:
  - A concept class C over n instances from the set X
  - A learner L with hypothesis space H
  - Two constants,  $\epsilon, \delta \in (0, \frac{1}{2})$
- C is said to be PAC learnable by L using H iff for all distributions over X, learner L by sampling n instances, will with probability at least  $1-\delta$  output a hypothesis  $h\in H$  such that
  - $-\epsilon_h \le \epsilon$
  - Running time is polynomial in  $\frac{1}{\epsilon}$ ,  $\frac{1}{\delta}$ , n, size(c)



- PAC concerned about computational resources required for learning
  - In practice, we are often only concerned about the number of training examples required
  - The two are related
    - The computational limitation also imposes a polynomial constraint on the training set size, since a learner can process at most polynomial data in polynomial time
    - The learner must visit each example at least once

