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Based on the slides of Vibhav Gogate and David Sontag



- 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 is 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

## PAC Learning



- Probably approximately correct (PAC)
  - 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  $h \in \mathbf{H}$ 
    - A learner is said to be consistent if it always outputs a perfect classifier (assuming that one exists)
  - Want to compute the (expected) 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, with respect to the data generating distribution, that h misclassifies a random data point

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



- Assume that there exists a hypothesis in *H* that perfectly classifies all data points and that |*H*| is finite
- The version space (set of consistent hypotheses) is said to be 
   *exhausted* if and only if every consistent hypothesis has true
   error less than 
   *e*
  - Want enough samples to guarantee that every consistent hypothesis has error at most  $\epsilon$
- We'll show that, given enough samples, w.h.p. every hypothesis with true error at least ε is not consistent with the data



- Let (x<sup>(1)</sup>, y<sup>(1)</sup>), ..., (x<sup>(M)</sup>, y<sup>(M)</sup>) be M labeled data points sampled independently according to p
- Let C<sup>h</sup><sub>m</sub> be a random variable that indicates whether or not the m<sup>th</sup> data point is correctly classified
- The probability that h misclassifies the  $m^{th}$  data point is

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



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Probability that a randomly
sampled pair (x,y) is
incorrectly classified by h



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This is the true error of hypothesis h



• Probability that all data points classified correctly?

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



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

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

$$p(C_1^h = 1, \dots, C_M^h = 1) = (1 - \epsilon_h)^M \le (1 - \epsilon)^M \le e^{-\epsilon M}$$

for  $\epsilon \leq 1$ 



- 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

$$p\left(\bigvee_{h\in H_{BAD}} \left(C_1^h = 1, \dots, C_M^h = 1\right)\right) \leq \sum_{h\in H_{BAD}} p\left(C_1^h = 1, \dots, C_M^h = 1\right)$$
$$\leq |H|_{BAD} |e^{-\epsilon M}$$
$$\leq |H| |e^{-\epsilon M}$$



- 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**



- What we just proved:
  - Theorem: For a finite hypothesis space, H, with M i.i.d. samples, and 0 < ε < 1, the probability that there exists a hypothesis in H that is consistent with the data but has true error larger than ε is at most |H|e<sup>-εM</sup>
- 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
  - That is, the probability that the version space is not  $\epsilon$ -exhausted is at most  $|H|e^{-\epsilon M} \leq \delta$
- Solving for *M* yields

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



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



- 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?



- 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?

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



- How do we handle situations with 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?
  - Coming soon...

#### 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  $\epsilon > 0$ ,

$$p\left(\left|y-\frac{1}{M}\sum_{m}Y_{m}\right| \geq \epsilon\right) \leq 2e^{-2M\epsilon^{2}}$$

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

$$p\left(\left|\epsilon_{h} - \frac{1}{M}\sum_{m}(1 - C_{m}^{h})\right| \ge \epsilon\right) \le 2e^{-2M\epsilon^{2}}$$

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• Applying this to  $1 - C_1^h, \dots, 1 - C_M^h$  gives

$$p\left(\epsilon_h - \frac{1}{M}\sum_m (1 - C_m^h) \ge \epsilon\right) \le e^{-2M\epsilon^2}$$

This is the training error

#### PAC Bounds



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

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

#### PAC Bounds



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

$$\epsilon_{h} \leq \epsilon_{h}^{train} + \sqrt{\frac{1}{2M} \left( \ln |H| + \ln \frac{1}{\delta} \right)}$$
  
"bias" "variance"

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

#### PAC Bounds



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$$\epsilon_{h} \leq \epsilon_{h}^{train} + \sqrt{\frac{1}{2M} \left( \ln |H| + \ln \frac{1}{\delta} \right)}$$
  
"bias" "variance"

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