



# Learning Theory

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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

- 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)
  - 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  $\epsilon$

- 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 (assuming that one exists)
  - Want to compute the (expected) error of the classifier

- 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  **$\epsilon$ -exhausted** if and only if every consistent hypothesis has true error less than  $\epsilon$ 
  - 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  $\epsilon$  is not consistent with the data**

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

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

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

for  $\epsilon \leq 1$

# 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

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



- 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  $0 < \epsilon < 1$ , the probability that **there exists a hypothesis in  $H$  that is consistent with the data but has true error larger than  $\epsilon$**  is at most  $|H|e^{-\epsilon M}$
- We can turn this into a **sample complexity bound**

- 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

$$\begin{aligned} M &\geq -\frac{1}{\epsilon} \ln \frac{\delta}{|H|} \\ &= \left( \ln |H| + \ln \frac{1}{\delta} \right) / \epsilon \end{aligned}$$

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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?

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

- 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 bound: Suppose  $Y_1, \dots, Y_M$  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  $1 - C_1^h, \dots, 1 - C_M^h$  gives

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This is the training error

- **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  $\delta \geq |H|e^{-2M\epsilon^2}$ )

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

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

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

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

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