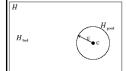


10-701/15-781, Fall 2011

Computational Learning Theory



Eric Xing



Lecture 6, September 28, 2011

Reading: Chap. 5 CB
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Generalizability of Learning



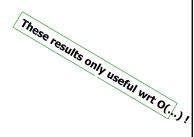
- In machine learning it's really the generalization error that we care about, but most learning algorithms fit their models to the training set.
- Why should doing well on the training set tell us anything about generalization error? Specifically, can we relate error on to training set to generalization error?
- Are there conditions under which we can actually prove that learning algorithms will work well?

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What General Laws constrain Inductive Learning?



- Sample Complexity
 - How many training examples are sufficient to learn target concept?
- Computational Complexity
 - Resources required to learn target concept?
- Want theory to relate:
 - Training examples
 - Quantity
 - Quality
 - How presented
 - Complexity of hypothesis/concept space H
 - Accuracy of approx to target concept
 - Probability of successful learning



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Two Basic Competing Models



PAC framework

Sample labels are consistent with some *h* in *H*

Learner's hypothesis required to meet absolute upper bound on its error

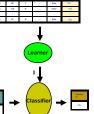
Agnostic framework

No prior restriction on the sample labels

The required upper bound on the hypothesis error is only relative (to the best hypothesis in the class)

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Protocol



- Given:
 - set of examples X
 - fixed (unknown) distribution D over X
 - set of hypotheses H
 - set of possible target concepts C
- Learner observes sample S = { ⟨ x_i, c(x_i) ⟩ }
 - instances x_i drawn from distr. D
 - labeled by target concept $c \in C$ (Learner does NOT know c(.), D)
- Learner outputs h ∈ H estimating c
 - h is evaluated by performance on subsequent instances drawn from D
- For now:
 - $C = H (so c \in H)$
 - Noise-free data

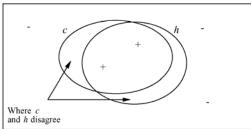
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True error of a hypothesis



Instance space X



• Definition: The *true error* (denoted $\varepsilon_D(h)$) of hypothesis h with respect to target concept c and distribution \mathcal{D} is the probability that h will misclassify an instance drawn at random according to \mathcal{D} .

$$\epsilon_{\mathcal{D}}(h) \equiv Pr_{x \in \mathcal{D}}[c(x) \neq h(x)]$$

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Two notions of error



- Training error (a.k.a., empirical risk or empirical error) of hypothesis h with respect to target concept c
 - How often $h(x) \neq c(x)$ over training instance from S

$$\hat{\epsilon}_{\mathcal{S}}(h) \equiv Pr_{x \in \mathcal{S}}[c(x) \neq h(x)] \equiv \frac{\sum_{x \in \mathcal{S}} \delta(c(x) \neq h(x))}{|\mathcal{S}|}$$

- True error of (a.k.a., generalization error, test error) hypothesis h with respect to c
 - How often $h(x) \neq c(x)$ over future random instances drew iid from \mathcal{D}

 $\epsilon_{\mathcal{D}}(h) \equiv Pr_{x \in \mathcal{D}}[c(x) \neq h(x)]$

Can we bound $\hat{\epsilon}_{\mathcal{D}}(h)$ in terms of $\hat{\epsilon}_{\mathcal{S}}(h)$

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The Union Bound



• Lemma. (The union bound). Let $A_1; A_2, \ldots, A_k$ be k different events (that may not be independent). Then

$$P(A_1 \cup A_2 \cup ... \cup A_k) \le P(A_1) + P(A_2) + ... + P(A_k)$$

In probability theory, the union bound is usually stated as an axiom (and thus we
won't try to prove it), but it also makes intuitive sense: The probability of any one
of k events happening is at most the sums of the probabilities of the k different
events.

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Hoeffding inequality



• Lemma. (Hoeding inequality) Let $Z_1,...,Z_m$ be m independent and identically distributed (iid) random variables drawn from a Bernoulli(ϕ) distribution, i.e., $P(Z_i = 1) = \phi$, and $P(Z_i = 0) = 1 - \phi$.

Let $\hat{\phi} = (1/m) \sum_{i=1}^{m} Z_i$ be the mean of these random variables, and let any $\gamma > 0$ be fixed. Then

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

• This lemma (which in learning theory is also called the Chernoff bound) says that if we take $\hat{\phi}$ — the average of m Bernoulli(ϕ) random variables — to be our estimate of ϕ , then the probability of our being far from the true value is small, so long as m is large.

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Version Space



A hypothesis h is consistent with a set of training examples S of target concept c if and only if h(x)=c(x) for each training example \(\lambda_{ij}, c(x_{ij}) \) in S

$$Consistent(h, S) \models h(x) = c(x), \forall \langle x, c(x) \rangle \in S$$

The version space, VS_{H,S}, with respect to hypothesis space H
and training examples S is the subset of hypotheses from H
consistent with all training examples in S.

$$VS_{H,S} \equiv \{h \in H|Consistent(h,S)\}$$

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Consistent Learner



- A learner is *consistent* if it outputs hypothesis that perfectly fits the training data
 - This is a quite reasonable learning strategy
- Every consistent learning outputs a hypothesis belonging to the version space
- We want to know how such hypothesis generalizes

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Probably Approximately Correct



Goal:

PAC-Learner produces hypothesis \hat{h} that is approximately correct, $err_D(\hat{h}) \approx 0$ with high probability $P\big(err_D(\hat{h}) \approx 0\big) \approx 1$

- Double "hedging"
 - approximately
 - probably

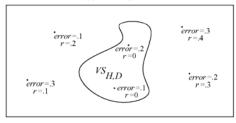
Need both!

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Hypothesis space H



(r = training error, error = true error)

 Definition: The version space VS_{H,S} is said to be ε-exhausted with respect to c and S, if every hypothesis h in VS_{H,S} has true error less than ε with respect to c and D.

$$\forall h \in VS_{H,S}, \quad \hat{\epsilon}_{\mathcal{D}}(h) < \epsilon$$

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How many examples will ϵ -exhaust the VS



Theorem: [Haussler, 1988].

If the hypothesis space H is finite, and S is a sequence of m ≥ 1 independent random examples of some target concept c, then for ANY 0 ≤ ε ≤ 1/2, the probability that the version space with respect to H and S is not ε-exhausted (with respect to c) is less than

$$|H|e^{-\epsilon m}$$

• This bounds the probability that any consistent learner will output a hypothesis h with $\varepsilon(h) \ge \varepsilon$

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Proof



h G V Sh.s h
$$\rightarrow$$
 men. enov. lex v. p. $506 \times G$

h G V Sh.s h \rightarrow no enov on lex v. p. $1-G$.

M.

h G V Sh.s h \rightarrow No ever on m example ? (1-G)^m.

 $P(\mp h.GV \le 0 \le 1. h \Rightarrow no ever on) \leftarrow (1-G)^m$

m liple

 $P(\forall h GV \le 0 \le 1. h \Rightarrow no ever on) \leftarrow (1-G)^m$
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 $P(\forall h GV \le 0$

What it means



• [Haussler, 1988]: probability that the version space is not ε -exhausted after m training examples is at most $|H|e^{-\varepsilon m}$

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$$Pr(\exists h \in H, \ s.t. \ (error_{train}(h) = 0) \land (error_{true}(h) > \epsilon) \) \le |H|e^{-\epsilon m}$$

Suppose we want this probability to be at most δ

$$|H|e^{-\epsilon m} \le \delta$$

1. How many training examples suffice?

$$m \ge \frac{1}{\epsilon} (\ln|H| + \ln(1/\delta))$$

2. If $error_{train}(h) = 0$ then with probability at least (1- δ):

$$error_{true} \leq \frac{1}{m} (\ln |H| + \ln(1/\delta))$$

PAC Learnability



A learning algorithm is PAC learnable if it

- Requires no more than polynomial computation per training example, and
- no more than polynomial number of samples

Theorem: conjunctions of Boolean literals is PAC learnable

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PAC-Learning



• Learner L can draw labeled instance $\langle x, c(x) \rangle$ in unit time, $x \in X$ of length n drawn from distribution \mathcal{D} , labeled by target concept $c \in C$

```
Def'n: Learner L PAC-learns class C using hypothesis space H if 
1. for any target concept c \in C, any distribution \mathscr{D}, any \epsilon such that 0 < \epsilon < 1/2, \delta such that 0 < \delta < 1/2, L returns h \in H s.t. w/ \text{prob.} \ge 1 - \delta, \text{err}_D(h) < \epsilon 
2. L's run-time (and hence, sample complexity) is \text{poly}(|x|, \text{size}(c), 1/\epsilon, 1/\delta)
```

• Sufficient:

```
Sufficient:

1. Only poly(...) training instances - |H| = 2^{\text{poly()}} m \ge \frac{1}{\varepsilon} (\ln |H| + \ln(1/\delta))

2. Only poly time / instance ...

Often C = H

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Agnostic Learning



So far, assumed $c \in H$

Agnostic learning setting: don't assume $c \in H$

- What do we want then?
 - The hypothesis h that makes fewest errors on training data
- What is sample complexity in this case?

$$m \ge \frac{1}{2\varepsilon^2} (\ln |H| + \ln(1/\delta))$$

derived from Hoeffding bounds:

$$\Pr[error_D(h) > error_S(h) + \varepsilon] \le e^{-2m\varepsilon^2}$$

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Empirical Risk Minimization Paradigm



- Choose a *Hypothesis Class H* of subsets of *X*.
- For an input sample S, find some h in H that fits S "well".
- For a new point x, predict a label according to its membership in h.

$$\hat{h} = \arg\min_{h \in H} \hat{\epsilon}_{\mathcal{S}}(h)$$

- Example:
 - Consider linear classification, and let $h_{\theta}(x) = 1\{\theta^T x \geq 0\}$ Then $H = \{h_{\theta} : h_{\theta}(x) = 1\{\theta^T x \geq 0\}, \ \theta \in R^{n+1}\}$

$$\hat{\theta} = \arg\min_{\theta} \hat{\epsilon}_{\mathcal{S}}(h_{\theta})$$

- We think of ERM as the most "basic" learning algorithm, and it will be this algorithm that we focus on in the remaining.
- In our study of learning theory, it will be useful to abstract away from the specific parameterization of hypotheses and from issues such as whether we're using a linear classier or an ANN

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The Case of Finite H



- $H = \{h_1, ..., h_k\}$ consisting of k hypotheses.
- We would like to give guarantees on the generalization error of \hat{h} .
- First, we will show that $\hat{\epsilon}(h)$ is a reliable estimate of $\epsilon(h)$ for all h.
- Second, we will show that this implies an upper-bound on the generalization error of \hat{h} .

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Misclassification Probability



- The outcome of a binary classifier can be viewed as a Bernoulli random variable $Z: Z = 1\{h_i(x) \neq c(x)\}$
- $\bullet \quad \text{For each sample:} \ \ Z_j = 1\{h_i(x_j) \neq c(x_j)\}$

$$\hat{\epsilon}(h_i) = \frac{1}{m} \sum_{j=1}^{m} Z_j$$

Hoeffding inequality

$$P(|\epsilon(h_i) - \hat{\epsilon}(h_i)| > \gamma) \le 2 \exp(-2\gamma^2 m)$$

• This shows that, for our particular h_i training error will be close to generalization error with high probability, assuming m is large.

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Uniform Convergence



- But we don't just want to guarantee that $\hat{\epsilon}(h_i)$ will be close $\epsilon(h_i)$ (with high probability) for just only one particular h_i . We want to prove that this will be true simultaneously for all $h_i \in H$
- For *k* hypothesis:

$$P(\exists h \in H, |\epsilon(h_i) - \hat{\epsilon}(h_i)| > \gamma) = P(A_1 \cup \ldots \cup A_k)$$

$$< \sum_{i=1}^k P(A_i)$$

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

• This means:

$$P(\neg \exists h \in H, |\epsilon(h_i) - \hat{\epsilon}(h_i)| > \gamma) = P(\forall h \in H, |\epsilon(h_i) - \hat{\epsilon}(h_i)| \le \gamma)$$
$$= 1 - 2k \exp(-2\gamma^2 m)$$

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• In the discussion above, what we did was, for particular values of m and γ , given a bound on the probability that:

for some $h_i \in H$

$$|\epsilon(h_i) - \hat{\epsilon}(h_i)| > \gamma$$

• There are three quantities of interest here: m and γ , and probability of error; we can bound either one in terms of the other two.

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Sample Complexity



• How many training examples we need in order make a guarantee?

$$P(\exists h \in H, |\epsilon(h) - \hat{\epsilon}(h)| > \gamma) = 2k \exp(-2\gamma^2 m)$$

• We find that if

$$m \ge \frac{1}{2\gamma^2} \log \frac{2k}{\delta}$$

then with probability at least 1- δ , we have that $|\epsilon(h_i) - \hat{\epsilon}(h_i)| \le \gamma$ for all $h_i \in H$

• The key property of the bound above is that the number of training examples needed to make this guarantee is only logarithmic in *k*, the number of hypotheses in H. This will be important later.

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Generalization Error Bound



• Similarly, we can also hold m and δ fixed and solve for γ in the previous equation, and show [again, convince yourself that this is right!] that with probability 1- δ , we have that for all $h_i \in H$

$$|\hat{\epsilon}(h) - \epsilon(h)| \le \sqrt{\frac{1}{m} \log \frac{2k}{\delta}}$$

• Define $h^* = \arg\min_{h \in H} \epsilon(h)$ to be the best possible hypothesis in H.

$$\begin{array}{lcl} \epsilon(\hat{h}) & \leq & \hat{\epsilon}(\hat{h}) + \gamma \\ & \leq & \hat{\epsilon}(\hat{h}^*) + \gamma \\ & \leq & \epsilon(\hat{h}^*) + 2\gamma \end{array}$$

If uniform convergence occurs, then the generalization error of $\epsilon(\hat{h})$ is at most 2γ worse than the best possible hypothesis in H!

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Summary



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

$$\varepsilon(\hat{h}) \leq \left(\min_{h \in \mathcal{H}} \varepsilon(h)\right) + 2\sqrt{\frac{1}{2m}\log\frac{2k}{\delta}}.$$

Corollary. Let $|\mathcal{H}| = k$, and let any δ, γ be fixed. Then for $\varepsilon(\hat{h}) \leq \min_{h \in \mathcal{H}} \varepsilon(h) + 2\gamma$ to hold with probability at least $1 - \delta$, it suffices that

$$\begin{array}{rcl} m & \geq & \frac{1}{2\gamma^2}\log\frac{2k}{\delta} \\ & = & O\left(\frac{1}{\gamma^2}\log\frac{k}{\delta}\right), \end{array}$$

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What if H is not finite?



- Can't use our result for infinite H
- Need some other measure of complexity for H
 - Vapnik-Chervonenkis (VC) dimension!

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How do we characterize "power"?



- Different machines have different amounts of "power".
- Tradeoff between:
 - More power: Can model more complex classifiers but might overfit.
 - Less power: Not going to overfit, but restricted in what it can model
- How do we characterize the amount of power?

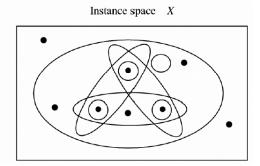
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The Vapnik-Chervonenkis Dimension



 Definition: The Vapnik-Chervonenkis dimension, VC(H), of hypothesis space H defined over instance space X is the size of the largest finite subset of X shattered by H. If arbitrarily large finite sets of X can be shattered by H, then VC(H) = ∞.



Definition:

Given a set $S = \{x(1), \dots, x(d)\}$ of points $x(\hat{p})$ \hat{I} X, we say that H shatters S if H can realize any labeling on S.

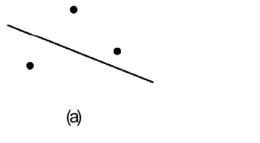
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VC dimension: examples



Consider $X = \mathbb{R}^2$, want to learn c: $X \rightarrow \{0,1\}$

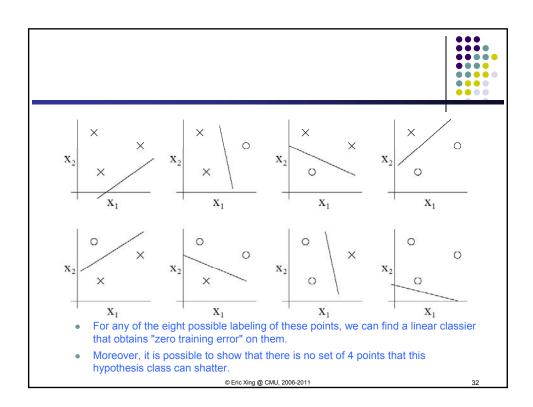
What is VC dimension of lines in a plane?
 H= { ((wx+b)>0 → y=1) }

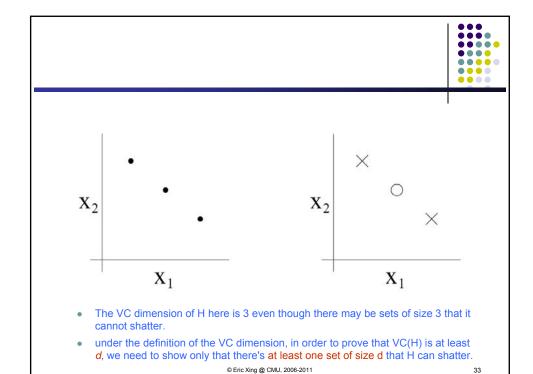


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(b)







- **Theorem** Consider some set of m points in \mathbb{R}^n . Choose any one of the points as origin. Then the m points can be shattered by oriented hyperplanes if and only if the position vectors of the remaining points are linearly independent.
- **Corollary**: The VC dimension of the set of oriented hyperplanes in \mathbb{R}^n is n+1.

Proof: we can always choose n+1 points, and then choose one of the points as origin, such that the position vectors of the remaining n points are linearly independent, but can never choose n+2 such points (since no n+1 vectors in \mathbb{R}^n can be linearly independent).

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The VC Dimension and the Number of Parameters

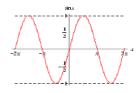


- The VC dimension thus gives concreteness to the notion of the capacity of a given set of *h*.
- Is it true that learning machines with many parameters would have high VC dimension, while learning machines with few parameters would have low VC dimension?

An infinite-VC function with just one parameter!

$$f(x,\alpha) \equiv \theta(\sin(\alpha x)), \quad x,\alpha \in R$$

where θ is an indicator function



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An infinite-VC function with just one parameter



• You choose some number *I*, and present me with the task of finding *I* points that can be shattered. I choose them to be

$$x_i = 10^{-i}$$
 $i = 1, \dots, l.$

• You specify any labels you like:

$$y_1, y_2, \dots, y_l, \quad y_i \in \{-1, 1\}$$

• Then $f(\alpha)$ gives this labeling if I choose α to be

$$\alpha = \pi (1 + \sum_{i=1}^{l} \frac{(1 - y_i)10^i}{2})$$

• Thus the VC dimension of this machine is infinite.

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Sample Complexity from VC Dimension



• How many randomly drawn examples suffice to ε -exhaust VS_{HS} with probability at least $(1 - \delta)$?

ie., to guarantee that any hypothesis that perfectly fits the training data is probably $(1-\delta)$ approximately (ϵ) correct on testing data from the same distribution

$$m \ge \frac{1}{\varepsilon} (4 \log_2(2/\delta) + 8VC(H) \log_2(13/\varepsilon))$$

Compare to our earlier results based on |H|:

$$m \ge \frac{1}{2\varepsilon^2} (\ln |H| + \ln(1/\delta))$$

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What You Should Know



- Sample complexity varies with the learning setting
- Learner actively queries trainer
 - Examples provided at random
- Within the PAC learning setting, we can bound the probability that learner will output hypothesis with given error
 - For ANY consistent learner (case where c in H)
 - For ANY "best fit" hypothesis (agnostic learning, where perhaps c not in H)
- VC dimension as measure of complexity of H

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