

Recap

For a single hypothesis, we have

$$\mathbb{P} \left[\left| \widehat{R}_n(h) - R(h) \right| > \epsilon \right] \leq 2e^{-2\epsilon^2 n}$$

For $m = |\mathcal{H}|$ hypotheses, and $h^* \in \mathcal{H}$, we have

$$\mathbb{P} \left[\left| \widehat{R}_n(h^*) - R(h^*) \right| > \epsilon \right] \leq 2me^{-2\epsilon^2 n}$$

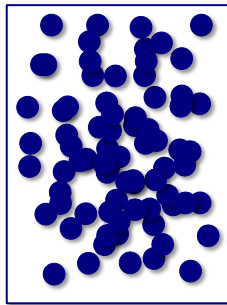
or equivalently, that with probability at least $1 - \delta$

$$R(h^*) \leq \widehat{R}_n(h^*) + \sqrt{\frac{1}{2n} \log \frac{2m}{\delta}}$$

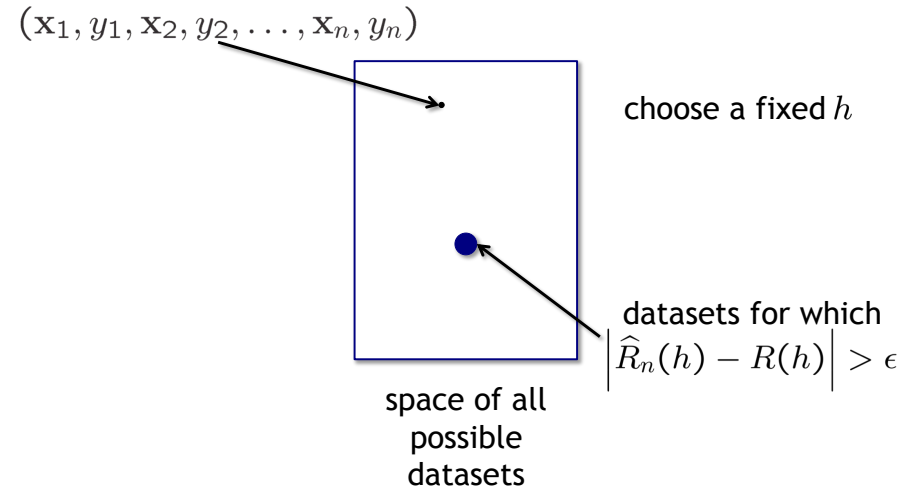
Bound becomes meaningless when $|\mathcal{H}| = \infty$

Union bound intuition

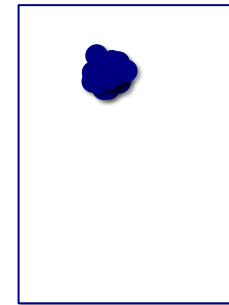
Consider many different h at once



Hoeffding's inequality



An alternative picture



If all the “bad” datasets overlap, maybe we can handle much bigger \mathcal{H} than the union bound suggests

Big idea from last time

Rather than measuring the “size” of \mathcal{H} with $|\mathcal{H}|$, we can instead think about:

Using \mathcal{H} , how many ways can we label a dataset?

We call a particular labeling of x_1, \dots, x_n a **dichotomy**

Using this language, we can answer our question via the growth function $m_{\mathcal{H}}(n)$, which counts the **most** dichotomies that \mathcal{H} could ever generate on n points

It is easy to see that $m_{\mathcal{H}}(n) \leq 2^n$

- If $m_{\mathcal{H}}(n) = 2^n$, we say that \mathcal{H} can **shatter** a set of size n
- If no set of size k can be shattered by \mathcal{H} ($m_{\mathcal{H}}(k) < 2^k$) then k is a **break point**

Bottom line

For a given \mathcal{H} , all we need is for a break point to exist

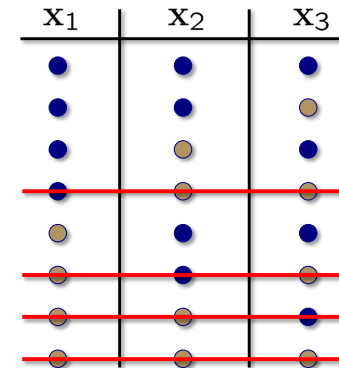
$$m_{\mathcal{H}}(n) \leq \underbrace{\sum_{i=0}^{k-1} \binom{n}{i}}_{\text{polynomial with dominant term } n^{k-1}}$$

All that remains is to argue that we can actually replace $|\mathcal{H}|$ with $m_{\mathcal{H}}(n)$ to obtain an inequality along the lines of

$$R(h^*) \leq \hat{R}_n(h^*) + \sqrt{\frac{1}{2n} \log \frac{2m_{\mathcal{H}}(n)}{\delta}}$$

How many dichotomies?

You are given a hypothesis set which has a break point of 2
How many dichotomies can you get on 3 data points?



VC generalization bound

We won't be able to quite show

$$R(h^*) \leq \hat{R}_n(h^*) + \sqrt{\frac{1}{2n} \log \frac{2m_{\mathcal{H}}(n)}{\delta}}$$

For technical reasons (which we will soon see), we will only be able to show that with probability $\geq 1 - \delta$

$$R(h^*) \leq \hat{R}_n(h^*) + \sqrt{\frac{8}{n} \log \frac{4m_{\mathcal{H}}(2n)}{\delta}}$$

This is called the **VC generalization bound**

Named after Vapnik and Chervonenkis, who proved it in 1971

Mathematical statement

Using Hoeffding's inequality together with a union bound, we were able to show that

$$\mathbb{P} \left[\max_{h \in \mathcal{H}} |\hat{R}_n(h) - R(h)| > \epsilon \right] \leq |\mathcal{H}| \cdot 2e^{-2\epsilon^2 n}$$

What the VC bound gives us is a generalization of the form

$$\mathbb{P} \left[\sup_{h \in \mathcal{H}} |\hat{R}_n(h) - R(h)| > \epsilon \right] \leq 2 \cdot m_{\mathcal{H}}(2n) \cdot 2e^{-\frac{1}{8}\epsilon^2 n}$$

supremum: maximum over an infinite set

Role of the growth function

We aim to get a bound on $\mathbb{P}[|\hat{R}_n(h) - R(h)| > \epsilon]$ that holds for any $h \in \mathcal{H}$, i.e., a bound on

$$\mathbb{P} \left[\sup_{h \in \mathcal{H}} |\hat{R}_n(h) - R(h)| > \epsilon \right]$$

Perhaps it is not surprising that we can understand $\hat{R}_n(h)$ using the growth function...

There may be infinitely many $h \in \mathcal{H}$, but \mathcal{H} can only generate $m_{\mathcal{H}}(n)$ **unique dichotomies**

Thus, $\hat{R}_n(h)$ can only take at most $m_{\mathcal{H}}(n)$ different values

Unfortunately, $R(h)$ can still take infinitely many different values, and so there are infinitely many $|\hat{R}_n(h) - R(h)|$

Supremum

The **supremum** of a set $S \subset T$ is the least element of T that is greater than or equal to all elements of S

Sometimes called the **least upper bound**

Examples

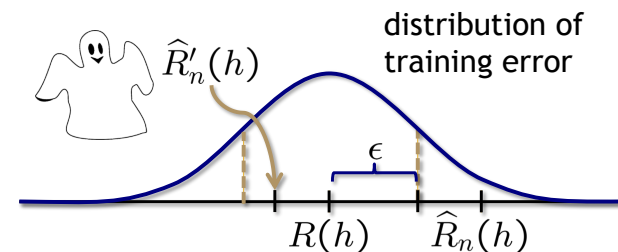
- $\sup\{1, 2, 3\} = 3$
- $\sup\{x : 0 \leq x \leq 1\} = 1$
- $\sup\{x : 0 < x < 1\} = 1$
- $\sup\{1 - 1/n : n > 0\} = 1$

The magic in the proof of the VC bound is to realize that we can relate the **supremum** over **all** $h \in \mathcal{H}$ to the **maximum** over a finite number of $h \in \mathcal{H}$ using a really cool trick!

Fundamental insight

The key insight (or trick) is to consider **two** datasets!

We will imagine that in addition to our training data, we have access to a second independent dataset (of size n), which we call the **ghost dataset**



Can we relate $\mathbb{P}[|\hat{R}_n(h) - R(h)| > \epsilon]$ to something like $\mathbb{P}[|\hat{R}_n(h) - \hat{R}'_n(h)| > \epsilon]$?

Using the ghost dataset

Suppose (for the moment) that the empirical estimates $\widehat{R}_n(h)$ and $\widehat{R}'_n(h)$ are random variables that are drawn from a *symmetric* distribution with mean (and median) $R(h)$

Consider the following events:

- A : the event that $|\widehat{R}_n(h) - R(h)| > \epsilon$
- B : the event that $|\widehat{R}_n(h) - \widehat{R}'_n(h)| > \epsilon$

Claim: $\mathbb{P}[B|A] \geq \frac{1}{2}$

Thus $\mathbb{P}[B] = \mathbb{P}[B|A] \cdot \mathbb{P}[A] \geq \frac{1}{2}\mathbb{P}[A]$

$\rightarrow \mathbb{P}[|\widehat{R}_n(h) - R(h)| > \epsilon] \leq 2\mathbb{P}[|\widehat{R}_n(h) - \widehat{R}'_n(h)| > \epsilon]$

Bounding the worst-case deviation

Lemma 2 (Where the magic happens)

Let $S = \{(x_i, y_i), i = 1, \dots, 2n\}$. Then

$$\begin{aligned} & \mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \right] \\ & \leq m_{\mathcal{H}}(2n) \cdot \sup_S \sup_{h \in \mathcal{H}} \underbrace{\mathbb{P} \left[|\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \middle| S \right]} \end{aligned}$$

here, the dataset S is fixed
the probability is with respect
to a random partition of S into
two training sets of size n

Using the ghost dataset

Unfortunately, the distribution of $\widehat{R}_n(h)$ and $\widehat{R}'_n(h)$ is binomial (not symmetric) so this exact statement doesn't hold in general, but the intuition is valid

Instead, we have the following bound:

Lemma 1 (Ghost dataset)



$$\begin{aligned} & \mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - R(h)| > \epsilon \right] \\ & \leq 2\mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \right] \end{aligned}$$

Proof of Lemma 2

It is straightforward to show that

$$\begin{aligned} & \mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \right] \\ & \leq \sup_S \mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \middle| S \right] \end{aligned}$$

Note that in the probability on the right-hand side, the dataset S is fixed.

Thus, there are only a finite number of dichotomies that \mathcal{H} can generate on S . Call this number $m_{\mathcal{H}}(S)$.

Let $h_1, \dots, h_{m_{\mathcal{H}}(S)}$ be the classifiers giving rise to these dichotomies

Proof of Lemma 2

Using this observation, we have

$$\begin{aligned}
 & \mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \middle| S \right] \\
 &= \mathbb{P} \left[\max_{h_1, \dots, h_{m_{\mathcal{H}}(S)}} |\widehat{R}_n(h_i) - \widehat{R}'_n(h_i)| > \frac{\epsilon}{2} \middle| S \right] \\
 &\leq \sum_{i=1}^{m_{\mathcal{H}}(S)} \mathbb{P} \left[|\widehat{R}_n(h_i) - \widehat{R}'_n(h_i)| > \frac{\epsilon}{2} \middle| S \right] \\
 &\leq m_{\mathcal{H}}(S) \max_{h_1, \dots, h_{m_{\mathcal{H}}(S)}} \mathbb{P} \left[|\widehat{R}_n(h_i) - \widehat{R}'_n(h_i)| > \frac{\epsilon}{2} \middle| S \right] \\
 &\leq m_{\mathcal{H}}(2n) \cdot \sup_{h \in \mathcal{H}} \mathbb{P} \left[|\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \middle| S \right]
 \end{aligned}$$

Putting it all together

$$\begin{aligned}
 & \mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - R(h)| > \epsilon \right] \\
 &\leq 2 \mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \right] \\
 &\leq 2 \cdot m_{\mathcal{H}}(2n) \cdot \sup_S \sup_{h \in \mathcal{H}} \mathbb{P} \left[|\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \middle| S \right] \\
 &\leq 2 \cdot m_{\mathcal{H}}(2n) \cdot 2e^{-\frac{1}{8}\epsilon^2 n}
 \end{aligned}$$

Thus, for any $h \in \mathcal{H}$, we have that with probability $\geq 1 - \delta$

$$R(h) \leq \widehat{R}_n(h) + \sqrt{\frac{8}{n} \log \frac{4m_{\mathcal{H}}(2n)}{\delta}}$$

Final step

At this point, we have shown

$$\begin{aligned}
 & \mathbb{P} \left[\sup_{h \in \mathcal{H}} |\widehat{R}_n(h) - R(h)| > \epsilon \right] \\
 &\leq 2 \cdot m_{\mathcal{H}}(2n) \cdot \sup_S \sup_{h \in \mathcal{H}} \mathbb{P} \left[|\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \middle| S \right]
 \end{aligned}$$

Lemma 3 (Random partitions)

For **any** h and **any** S ,

$$\mathbb{P} \left[|\widehat{R}_n(h) - \widehat{R}'_n(h)| > \frac{\epsilon}{2} \middle| S \right] \leq 2e^{-\frac{1}{8}\epsilon^2 n}$$

Proof follows from a simple lemma (also by Hoeffding)

Using the VC bound: The VC dimension

We went to a lot of trouble to show that if k is a break point for \mathcal{H} , then $m_{\mathcal{H}}(n) \leq \sum_{i=0}^{k-1} \binom{n}{i} \leq n^{k-1} + 1$

$$\begin{aligned}
 \Rightarrow R(h) &\leq \widehat{R}_n(h) + \sqrt{\frac{8}{n} \log \frac{4((2n)^{k-1} + 1)}{\delta}} \\
 &\lesssim \widehat{R}_n(h) + \sqrt{\frac{8(k-1)}{n} \log \frac{8n}{\delta}} \quad \leftarrow \text{True for } k \geq 3
 \end{aligned}$$

The **VC dimension** of a hypothesis set \mathcal{H} , denoted $d_{\text{VC}}(\mathcal{H})$, is the largest d for which $m_{\mathcal{H}}(n) = 2^n$

- $d_{\text{VC}}(\mathcal{H})$ is the most points that \mathcal{H} can shatter
- $d_{\text{VC}}(\mathcal{H})$ is 1 less than the smallest break point

$$\Rightarrow R(h) \lesssim \widehat{R}_n(h) + \sqrt{\frac{8d_{\text{VC}} \log \frac{8n}{\delta}}{n}}$$

Examples

- Positive rays:

$$d_{VC} = 1$$

- Positive intervals:

$$d_{VC} = 2$$

- Convex sets:

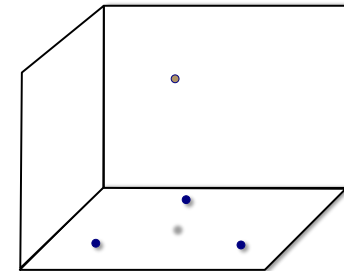
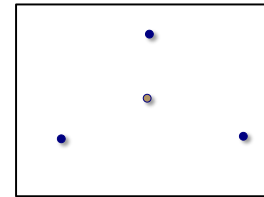
$$d_{VC} = \infty$$

- Linear classifiers in \mathbb{R}^2 :

$$d_{VC} = 3$$

VC dimension of general linear classifiers

For $d = 2$, $d_{VC} = 3$



In general $d_{VC} = d + 1$

We will prove this by showing that $d_{VC} \leq d + 1$ and $d_{VC} \geq d + 1$

One direction

Lets first show that there exists a set of $d + 1$ points in \mathbb{R}^d that are shattered

$$\mathbf{X} = \underbrace{\begin{bmatrix} -\tilde{\mathbf{x}}_1^T \\ -\tilde{\mathbf{x}}_2^T \\ \vdots \\ -\tilde{\mathbf{x}}_{d+1}^T \end{bmatrix}}_{d+1} = \begin{bmatrix} 1 & 0 & 0 & \cdots & 0 \\ 1 & 1 & 0 & \cdots & 0 \\ 1 & 0 & 1 & \cdots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & 0 & 0 & \cdots & 1 \end{bmatrix}$$

One can show that \mathbf{X} is invertible

Can we shatter this data set?

For any $\mathbf{y} = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_{d+1} \end{bmatrix} = \begin{bmatrix} \pm 1 \\ \pm 1 \\ \vdots \\ \pm 1 \end{bmatrix}$, can we find a vector

satisfying $\text{sign}(\mathbf{X}\boldsymbol{\theta}) = \mathbf{y}$?

Easy! Just make $\boldsymbol{\theta} = \mathbf{X}^{-1}\mathbf{y}$ and we have

$$\text{sign}(\mathbf{X}\boldsymbol{\theta}) = \text{sign}(\mathbf{y}) = \mathbf{y}$$

We can shatter a set of $d + 1$ points

What does this prove?

- a) $d_{VC} = d + 1$
- b) $d_{VC} \geq d + 1$ ✓
- c) $d_{VC} \leq d + 1$
- d) None of the above

The other direction

Take any $d + 2$ points $\tilde{\mathbf{x}}_1, \dots, \tilde{\mathbf{x}}_{d+2}$

More points than dimensions, so there must be some j for which

$$\tilde{\mathbf{x}}_j = \sum_{i \neq j} \alpha_i \tilde{\mathbf{x}}_i$$

where not all $\alpha_i = 0$

Consider the dichotomy where the $\tilde{\mathbf{x}}_i$ with $\alpha_i \neq 0$ are labeled $y_i = \text{sign}(\alpha_i)$, and $y_j = -1$

No linear classifier can implement such a dichotomy!

To finish the proof

In order to show that $d_{VC} \leq d + 1$, we need to show

- a) There are $d + 1$ points we cannot shatter
- b) There are $d + 2$ points we cannot shatter
- c) We cannot shatter any set of $d + 1$ points
- d) We cannot shatter any set of $d + 2$ points ✓

Why not?

$$\tilde{\mathbf{x}}_j = \sum_{i \neq j} \alpha_i \tilde{\mathbf{x}}_i \rightarrow \boldsymbol{\theta}^T \tilde{\mathbf{x}}_j = \sum_{i \neq j} \alpha_i \boldsymbol{\theta}^T \tilde{\mathbf{x}}_i$$

If $y_i = \text{sign}(\boldsymbol{\theta}^T \tilde{\mathbf{x}}_i) = \text{sign}(\alpha_i)$, then $\alpha_i \boldsymbol{\theta}^T \tilde{\mathbf{x}}_i > 0$

This means that $\boldsymbol{\theta}^T \tilde{\mathbf{x}}_j = \sum_{i \neq j} \alpha_i \boldsymbol{\theta}^T \tilde{\mathbf{x}}_i > 0$

Thus $y_j = \text{sign}(\boldsymbol{\theta}^T \tilde{\mathbf{x}}_j) = +1$

Interpreting the VC dimension

We have just shown that for a linear classifier in \mathbb{R}^d

$$\begin{array}{l} d_{VC} \geq d + 1 \\ d_{VC} \leq d + 1 \end{array} \quad \longrightarrow \quad d_{VC} = d + 1$$

How many parameters does a linear classifier in \mathbb{R}^d have?

$$\begin{array}{l} \mathbf{w} \in \mathbb{R}^d \\ b \in \mathbb{R} \end{array} \quad \longrightarrow \quad d + 1$$

Effective number of parameters

Additional parameters do not always contribute additional degrees of freedom

Example

Take the output of a linear classifier, and then feed this into another linear classifier

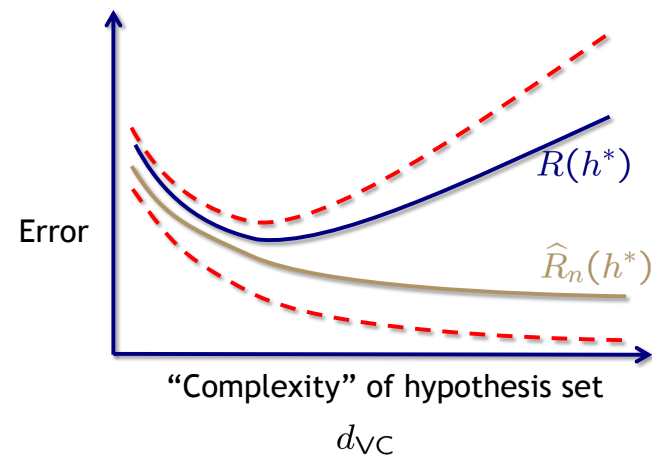
$$y_i = \text{sign}(w'(\text{sign}(\theta^T \mathbf{x}_i)) + b')$$

The parameters w' and b' are totally redundant (they do not allow us to create any new classifiers/dichotomies)

The usual examples

- Positive rays
 - $d_{VC} = 1$
 - 1 parameter
- Positive intervals
 - $d_{VC} = 2$
 - 2 parameters
- Convex sets
 - $d_{VC} = \infty$
 - as many parameters as you want

Interpreting the VC bound



VC bound in action

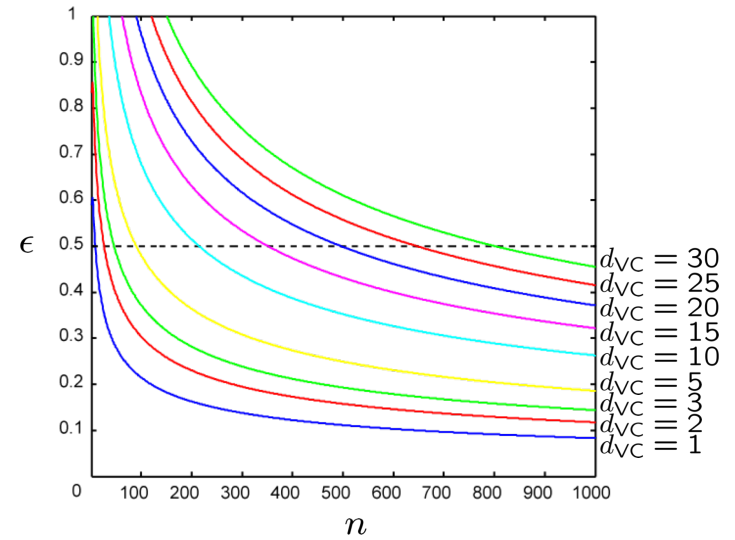
How big does our training set need to be?

$$R(h) \lesssim \hat{R}_n(h) + \underbrace{\sqrt{\frac{8d_{VC}}{n} \log \frac{8n}{\delta}}}_{\epsilon}$$

Just to see how this behaves, let's ignore the constants and suppose that

$$\epsilon \sim \sqrt{\frac{d_{VC}}{n} \log n}$$

VC bound in action



RULE OF THUMB: $n \geq 10d_{VC}$