

Our question: Does $E_{in}(h)$ say anything about $E_{out}(h)$?

Probability of a “bad” event (fixed h)

(Hoeffding)

$$P\left(|E_{in}(h) - E_{out}(h)| > \epsilon\right) \leq 2e^{-2\epsilon^2 N}$$

Probability of a “bad” event (g selected from a set of M hypothesis)

$$P\left(|E_{in}(g) - E_{out}(g)| > \epsilon\right) \leq 2Me^{-2\epsilon^2 N}$$

Compare the experiment of **tossing one coin N times** with the experiment of **tossing M coins, N times each**. The chance of a coin resulting in N heads is much larger for the second case.

$$P\left(|E_{in}(g) - E_{out}(g)| > \epsilon\right) \leq 2Me^{-2\epsilon^2 N}$$

M appears because of the **union bound**, which does not take overlaps among “bad” events into consideration

Can we find another bound that takes the overlaps into consideration ?

(and also works for infinite Hypothesis set?)

Let

- $X = \{\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N\}$ (N points)
- \mathcal{H} : a hypothesis space

Dichotomies generated by \mathcal{H} :

any bipartition of X as $X_{-1} \cup X_{+1}$ that agrees with a hypothesis $h \in \mathcal{H}$

$$\mathcal{H}(\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N) = \left\{ (h(\mathbf{x}_1), h(\mathbf{x}_2), \dots, h(\mathbf{x}_N)) \mid h \in \mathcal{H} \right\}$$

We know that $|\mathcal{H}(\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N)| \leq 2^N$

$$m_{\mathcal{H}}(N) = \max_{\mathbf{x}_1, \dots, \mathbf{x}_N \in \mathcal{X}} |\mathcal{H}(\mathbf{x}_1, \dots, \mathbf{x}_N)|$$

Perceptron 2D: $m_{\mathcal{H}}(3) = 8 = 2^3$, $m_{\mathcal{H}}(4) = 14 < 2^4$

Positive rays: $m_{\mathcal{H}}(N) = N + 1$

Positive intervals: $m_{\mathcal{H}}(N) = \frac{1}{2}N^2 + \frac{1}{2}N + 1$

Convex sets: $m_{\mathcal{H}}(N) = 2^N$

Break point

If no dataset of size k can be shattered by \mathcal{H} then k is a break point for \mathcal{H}

Perceptron 2D: $k = 4$ is a break point

Positive rays: $m_{\mathcal{H}}(N) = N + 1$, break point $k = 2$

Positive intervals: $m_{\mathcal{H}}(N) = \frac{1}{2}N^2 + \frac{1}{2}N + 1$, break point $k = 3$

Convex sets: $m_{\mathcal{H}}(N) = 2^N$, break point $k = +\infty$

Outline

- Proof that $m_{\mathcal{H}}(N)$ is polynomial
- Proof that $m_{\mathcal{H}}(N)$ can replace M

Bounding $m_{\mathcal{H}}(N)$

To show: $m_{\mathcal{H}}(N)$ is polynomial

We show: $m_{\mathcal{H}}(N) \leq \dots \leq \dots \leq$ a polynomial

Key quantity:

$B(N, k)$: Maximum number of dichotomies on N points, with break point k

$B(N, k)$: Maximum number of dichotomies on N points, with break point k

Example of last meeting: Supposing $k = 2$ is a break point, we computed $B(3, k) = 4$

x_1	x_2	x_3
○	○	○
○	○	●
○	●	○
●	○	○

Recursive bound on $B(N, k)$

Consider the following table:

$$B(N, k) = \alpha + 2\beta$$

	# of rows	x_1	x_2	...	x_{N-1}	x_N
S_1	α	+1	+1	...	+1	+1
		-1	+1	...	+1	-1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	-1	-1
		-1	+1	...	-1	+1
S_2^+	β	+1	-1	...	+1	+1
		-1	-1	...	+1	+1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	+1	+1
		-1	-1	...	-1	+1
S_2^-	β	+1	-1	...	+1	-1
		-1	-1	...	+1	-1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	+1	-1
		-1	-1	...	-1	-1

Estimating α and β

Focus on $\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_{N-1}$ columns:

$$\alpha + \beta \leq B(N-1, k)$$

	# of rows	\mathbf{x}_1	\mathbf{x}_2	...	\mathbf{x}_{N-1}	\mathbf{x}_N
S_1	α	+1	+1	...	+1	+1
		-1	+1	...	+1	-1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	-1	-1
		-1	+1	...	-1	+1
S_2^+	β	+1	-1	...	+1	+1
		-1	-1	...	+1	+1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	+1	+1
		-1	-1	...	-1	+1
S_2^-	β	+1	-1	...	+1	-1
		-1	-1	...	+1	-1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	+1	-1
		-1	-1	...	-1	-1

Estimating β by itself

Now, focus on the $S_2 = S_2^+ \cup S_2^-$ rows:

$$\beta \leq B(N-1, k-1)$$

	# of rows	x_1	x_2	...	x_{N-1}	x_N
S_1	α	+1	+1	...	+1	+1
		-1	+1	...	+1	-1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	-1	-1
		-1	+1	...	-1	+1
S_2	β	+1	-1	...	+1	+1
		-1	-1	...	+1	+1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	+1	+1
		-1	-1	...	-1	+1
S_2^-	β	+1	-1	...	+1	-1
		-1	-1	...	+1	-1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	+1	-1
		-1	-1	...	-1	-1

Putting it together

$$B(N, k) = \alpha + 2\beta$$

$$\alpha + \beta \leq B(N - 1, k)$$

$$\beta \leq B(N - 1, k - 1)$$

$$B(N, k) \leq$$

$$B(N - 1, k) + B(N - 1, k - 1)$$

	# of rows	x_1	x_2	...	x_{N-1}	x_N
S_1	α	+1	+1	...	+1	+1
		-1	+1	...	+1	-1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	-1	-1
		-1	+1	...	-1	+1
S_2^+	β	+1	-1	...	+1	+1
		-1	-1	...	+1	+1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	+1	+1
		-1	-1	...	-1	+1
S_2^-	β	+1	-1	...	+1	-1
		-1	-1	...	+1	-1
		⋮	⋮	⋮	⋮	⋮
		+1	-1	...	+1	-1
		-1	-1	...	-1	-1

Numerical computation of $B(N, k)$ bound

$$B(N, k) \leq B(N - 1, k) + B(N - 1, k - 1)$$



		k						
		1	2	3	4	5	6	..
N	1	1	2	2	2	2	2	..
	2	1	3	4	4	4	4	..
	3	1	4	7	8	8	8	..
	4	1	5	11
	5	1	6	:	.			
	6	1	7	:		.		
	:	:	:	:				.

Analytic solution for $B(N, k)$ bound

$$B(N, k) \leq B(N - 1, k) + B(N - 1, k - 1)$$

Theorem:

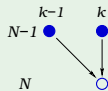
$$B(N, k) \leq \sum_{i=0}^{k-1} \binom{N}{i}$$

1. Boundary conditions: easy

	k						
	1	2	3	4	5	6	..
1	1	2	2	2	2	2	..
2	1						
3	1						
N	1			●	●		
5	1				○		
6	1						
:	:						

2. The induction step

$$\begin{aligned}\sum_{i=0}^{k-1} \binom{N}{i} &= \sum_{i=0}^{k-1} \binom{N-1}{i} + \sum_{i=0}^{k-2} \binom{N-1}{i} \text{ ?} \\ &= 1 + \sum_{i=1}^{k-1} \binom{N-1}{i} + \sum_{i=1}^{k-1} \binom{N-1}{i-1} \\ &= 1 + \sum_{i=1}^{k-1} \left[\binom{N-1}{i} + \binom{N-1}{i-1} \right] \\ &= 1 + \sum_{i=1}^{k-1} \binom{N}{i} = \sum_{i=0}^{k-1} \binom{N}{i} \checkmark\end{aligned}$$



It is polynomial!

For a given \mathcal{H} , the break point k is fixed

$$m_{\mathcal{H}}(N) \leq \underbrace{\sum_{i=0}^{k-1} \binom{N}{i}}_{\text{maximum power is } N^{k-1}}$$

Three examples

$$\sum_{i=0}^{k-1} \binom{N}{i}$$

- \mathcal{H} is positive rays: (break point $k = 2$)

$$m_{\mathcal{H}}(N) = N + 1 \leq N + 1$$

- \mathcal{H} is positive intervals: (break point $k = 3$)

$$m_{\mathcal{H}}(N) = \frac{1}{2}N^2 + \frac{1}{2}N + 1 \leq \frac{1}{2}N^2 + \frac{1}{2}N + 1$$

- \mathcal{H} is 2D perceptrons: (break point $k = 4$)

$$m_{\mathcal{H}}(N) = ? \leq \frac{1}{6}N^3 + \frac{5}{6}N + 1$$

Outline

- Proof that $m_{\mathcal{H}}(N)$ is polynomial
- Proof that $m_{\mathcal{H}}(N)$ can replace M

What we want

Instead of:

$$\mathbb{P}[|E_{\text{in}}(g) - E_{\text{out}}(g)| > \epsilon] \leq 2 M e^{-2\epsilon^2 N}$$

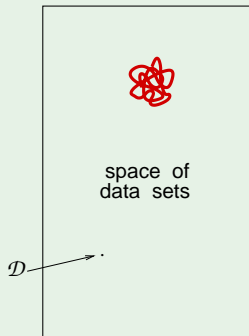
We want:

$$\mathbb{P}[|E_{\text{in}}(g) - E_{\text{out}}(g)| > \epsilon] \leq 2 m_{\mathcal{H}}(N) e^{-2\epsilon^2 N}$$

Pictorial proof ☺

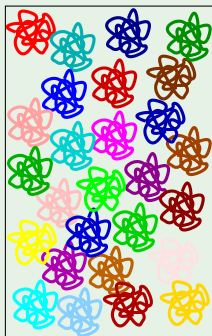
- How does $m_{\mathcal{H}}(N)$ relate to overlaps?
- What to do about E_{out} ?
- Putting it together

Hoeffding Inequality



(a)

Union Bound

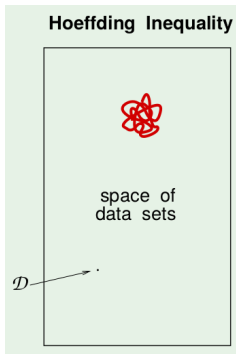


(b)

VC Bound

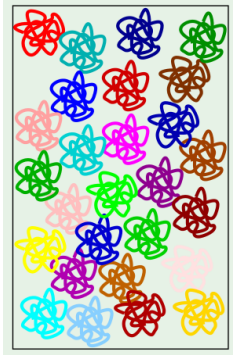


(c)



- The canvas is the space of all possible datasets of size N
- Each point in the canvas is a dataset of size N
- Given a hypothesis h , one can compute $E_{in}(h)$ with respect to each dataset
- The red points are the “bad” events for h (i.e., $|E_{in}(h) - E_{out}(h)| > \epsilon$)
- According to Hoeffding the probability of “bad” event of h is bounded (thus only a small area of the canvas is painted red)

Union Bound

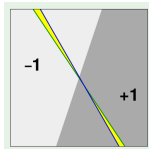


- When we have multiple hypothesis, we should consider the probability of “bad” events associated to all of them
- Each color in the canvas correspond to points that are the “bad” events for a specific h (i.e., $|E_{in}(h) - E_{out}(h)| > \epsilon$)
- Since we are considering the union bound (no overlaps between “bad” events) a large area of the canvas is colored (as “bad” events)

VC Bound



- It is very reasonable to think that one dataset corresponds to “bad” event for multiple hypothesis
- For instance, the two separating lines could have $E_{in} = 0$ and both have large error (the same dataset corresponds to a “bad” event for both)



- Considering the overlaps, the canvas painting should look like the one at the left, suggesting a bound larger than the original Hoeffding bound but much smaller than the union bound

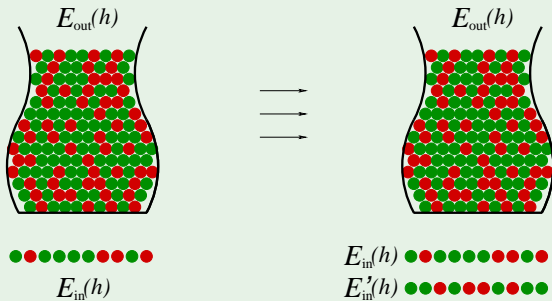
Many hypotheses share the same dichotomy on a given \mathcal{D} , since there are finitely many dichotomies even with an infinite number of hypotheses. Any statement based on \mathcal{D} alone will be simultaneously true or simultaneously false for all the hypotheses that look the same on that particular \mathcal{D} . What

The growth function groups hypotheses based on their behavior on D

The event $|E_{in}(h) - E_{out}(h)| > \epsilon$ depends not only on D , but also on entire \mathcal{X}

Now we need to consider the event $|E_{in}(h) - E_{out}(h)| > \epsilon$ in terms of a group of hypothesis ...

What to do about E_{out}



Putting it together

Not quite:

$$\mathbb{P}[|E_{\text{in}}(g) - E_{\text{out}}(g)| > \epsilon] \leq 2 m_{\mathcal{H}}(N) e^{-2\epsilon^2 N}$$

but rather:

$$\mathbb{P}[|E_{\text{in}}(g) - E_{\text{out}}(g)| > \epsilon] \leq 4 m_{\mathcal{H}}(2N) e^{-\frac{1}{8}\epsilon^2 N}$$

The Vapnik-Chervonenkis Inequality

$$P\left(|E_{in}(g) - E_{out}(g)| > \epsilon\right) \leq 2 m_{\mathcal{H}}(N) e^{-2\epsilon^2 N}$$

$$P\left(|E_{in}(g) - E_{out}(g)| > \epsilon\right) \leq 4 m_{\mathcal{H}}(2N) e^{-\frac{1}{8}\epsilon^2 N}$$

$2N$:

- hypotheses are grouped based on their behavior on D , but their behavior outside D is not the same
- to track $|E_{in}(h) - E_{out}(h)| > \epsilon$, we track $|E_{in}(h) - E'_{in}(h)| > \epsilon$ (relative to D and D' , both of size N)

4 and $\frac{1}{8}$:

- these are factors to account for the uncertainties added when we replace $|E_{in}(h) - E_{out}(h)| > \epsilon$ with $|E_{in}(h) - E'_{in}(h)| > \epsilon$

Summary

- The growth function (counts number of dichotomies) is polynomially bounded if \mathcal{H} has a break point
- The growth function can replace M
- Main result: VC inequality

$$P\left(|E_{in}(g) - E_{out}(g)| > \varepsilon\right) \leq 4 m_{\mathcal{H}}(2N) e^{-\frac{1}{8}\varepsilon^2 N}$$

Again, we have a bound that can be made small enough by taking a sufficiently large N

- Next meeting: (i) Do we need to have the growth function ?
(ii) Sample complexity