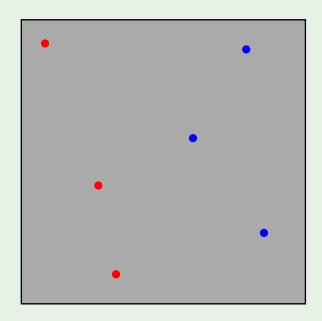
Review of Lecture 5

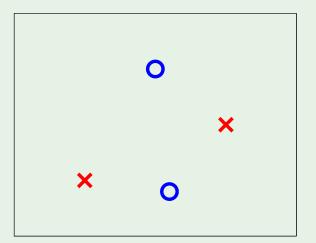
• Dichotomies = hypotheses restricted to a finite set of points



• Growth function

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

Break point



Maximum # of dichotomies

resulting from the constraint of the break point (here k=2)

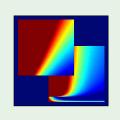
| \mathbf{x}_1 | \mathbf{x}_2 | \mathbf{x}_3 |
|----------------|----------------|----------------|
| 0 | 0 | 0 |
| 0 | 0 | • |
| 0 | • | 0 |
| • | 0 | 0 |

Learning From Data

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Lecture 6: Theory of Generalization





Outline

ullet Proof that $m_{\mathcal{H}}(N)$ is polynomial (with a break point)

ullet Proof that $m_{\mathcal{H}}(N)$ can replace M

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

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

We show: $m_{\mathcal{H}}(N) \leq \cdots \leq m \leq n$ polynomial

We want to bound m, and we do this with B(N, k) - the maximum number of dichotomies you can possibly have with no other constraints (other than N, k). This is purely combinatorial, meaning we can avoid any consideration of input space or correlation between events etc.

Key quantity:

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

i.e max number of dichotomies on N points so that no k columns are shattered.

(non-specific to X or H, although H does determine k)

Recursive bound on B(N, k)

Consider the following table:

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

B(N,k) is the maximum number of patterns we can get of N points such that no k columns have all possible patterns (are shattered).

S1 contains rows which appear only once as far as x1 to xN-1 are concerned - the prefix (x1-xN-1) happens once and only has one extension (xN=+1 OR xN=-1)

S2 contains prefixes with both xN=+1 AND xN=-1 - we split each of these into subgroups S2+ and S2-

| | | # of rows | \mathbf{x}_1 | \mathbf{x}_2 | | \mathbf{x}_{N-1} | \mathbf{x}_N |
|-----------------------------|---------|-----------|----------------|----------------|-------|--------------------|----------------|
| | | | +1 | +1 | • • • | +1 | +1 |
| | | | -1 | +1 | | +1 | -1 |
| | S_1 | lpha | : | : | : | : | : |
| | | | +1 | -1 | | -1 | -1 |
| | | | -1 | +1 | | -1 | +1 |
| | | | +1 | -1 | | +1 | +1 |
| | S_2^+ | eta | -1 | -1 | | +1 | +1 |
| | | | ÷ | ÷ | ÷ | : | : |
| | | | +1 | -1 | | +1 | +1 |
| S_2 | | | -1 | -1 | | -1 | +1 |
| $\mathcal{O}_{\mathcal{L}}$ | S_2^- | eta | +1 | -1 | | +1 | -1 |
| | | | -1 | -1 | | +1 | -1 |
| | | | ÷ | ÷ | ÷ | : | ÷ |
| | | | +1 | -1 | | +1 | -1 |
| | | | -1 | -1 | | -1 | -1 |
| | | | | | | | |

Estimating α and β

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

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

All rows (total alpha+beta) highlighted are different, (note S2+ and S2- have equal prefixes, so not different).

Also, on the original matrix we could not find all possible patterns on any k columns, so we also cannot on the highlighted matrix. If we could, then S_2 these k columns would feature all possible patterns in the original matrix, but we do not. So the smaller matrix has the same break point k.

| | # of rows | \mathbf{x}_1 | \mathbf{x}_2 | | \mathbf{x}_{N-1} | \mathbf{x}_N |
|---------|-----------|----------------|--|--|--|--|
| | | +1 | +1 | | +1 | +1 |
| | | -1 | +1 | | +1 | -1 |
| $ S_1 $ | α | : | : | : | ŧ | : |
| | | +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 |
| | | +1 | -1 | | +1 | -1 |
| S_2^- | | -1 | -1 | | +1 | -1 |
| | β | : | : | : | : | : |
| | | +1 | -1 | | +1 | -1 |
| | | -1 | -1 | | -1 | -1 |
| | S_2^+ | S_2^+ eta | $egin{array}{cccccccccccccccccccccccccccccccccccc$ | $egin{array}{cccccccccccccccccccccccccccccccccccc$ | $egin{array}{cccccccccccccccccccccccccccccccccccc$ | S_1 α $\begin{array}{c} +1 & +1 & \dots & +1 \\ -1 & +1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ -1 & +1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & -1 \\ -1 & +1 & \dots & -1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ -1 & -1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ -1 & -1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ -1 & -1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ \end{array}$ $\begin{array}{c} +1 & -1 & \dots & +1 \\ \end{array}$ |

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 | \mathbf{x}_1 | \mathbf{x}_2 | | \mathbf{x}_{N-1} | $ \mathbf{x}_N $ |
|-------|---------|-----------|----------------|----------------|---|--------------------|------------------|
| | | | +1 | +1 | | +1 | +1 |
| | | | -1 | +1 | | +1 | -1 |
| | S_1 | α | : | : | : | : | : |
| | | | +1 | -1 | | -1 | -1 |
| | | | -1 | +1 | | -1 | +1 |
| | S_2^+ | eta | +1 | -1 | | +1 | +1 |
| | | | -1 | -1 | | +1 | +1 |
| | | | i | ÷ | ÷ | : | : |
| | | | +1 | -1 | | +1 | +1 |
| S_2 | | | -1 | -1 | | -1 | +1 |
| | | | +1 | -1 | | +1 | -1 |
| | | | -1 | -1 | | +1 | -1 |
| | S_2^- | eta | : | : | : | : | : |
| | | | +1 | -1 | | +1 | -1 |
| | | | -1 | -1 | | -1 | -1 |

Putting it together

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

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

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

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

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

| | | # 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 | S_2^+ | β | +1 -1 : +1 -1 | -1 -1 : -1 -1 | | +1 +1 : +1 -1 | +1 +1 :: +1 +1 |
| | S_2^- | eta | +1 -1 : +1 -1 | -1 -1 : -1 -1 | · · · · · · · · · · · · · · · · · · · | +1 +1 : +1 -1 | -1 -1 : -1 -1 |

Numerical computation of B(N, k) bound

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Analytic solution for B(N, k) bound

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

Theorem:

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

1. Boundary conditions: easy

| | | k | | | | | | |
|---|---|---|---|---|---|---|---|-----|
| | | 1 | 2 | 3 | 4 | 5 | 6 | • • |
| | 1 | 1 | 2 | 2 | 2 | 2 | 2 | |
| | 2 | | | | | | | |
| | 3 | 1 | | | | | | |
| N | 4 | 1 | | | | | | |
| | 5 | 1 | | | | | | |
| | 6 | 1 | | | | | | |
| | • | • | | | | | | |

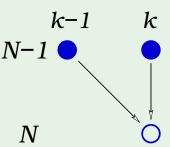
2. The induction step

$$\sum_{i=0}^{k-1} \binom{N}{i} = \sum_{i=0}^{k-1} \binom{N-1}{i} + \sum_{i=0}^{k-2} \binom{N-1}{i}?$$

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

$$= 1 + \sum_{i=1}^{k-1} \left[\binom{N-1}{i} + \binom{N-1}{i-1} \right]$$

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



It is polynomial!

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

$$m_{\mathcal{H}}(N) \leq \sum_{i=0}^{k-1} \binom{N}{i}$$
 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)

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

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

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

Outline

ullet Proof that $m_{\mathcal{H}}(N)$ is polynomial

ullet Proof that $m_{\mathcal{H}}(N)$ can replace M

What we want

Instead of:

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

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 ©

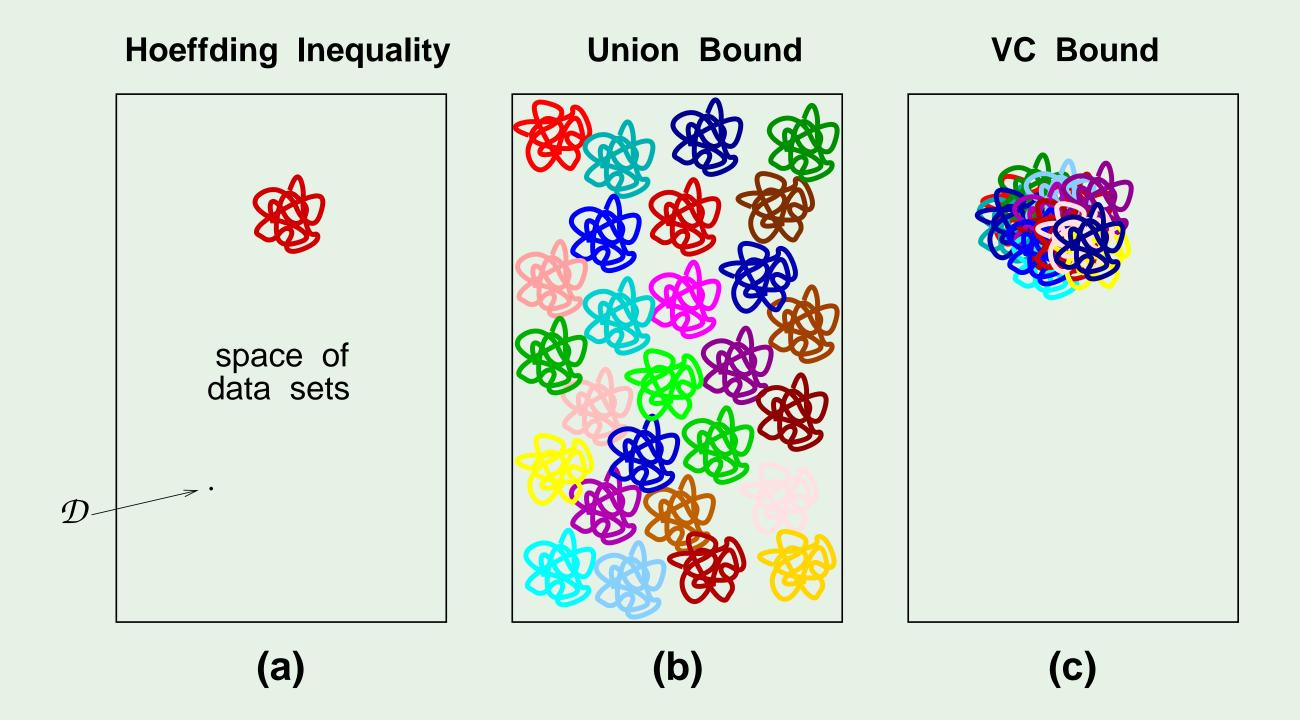
ullet How does $m_{\mathcal{H}}(N)$ relate to overlaps? since M is created from the union bound which assumes disjoint hypotheses

ullet What to do about $E_{
m out}$?

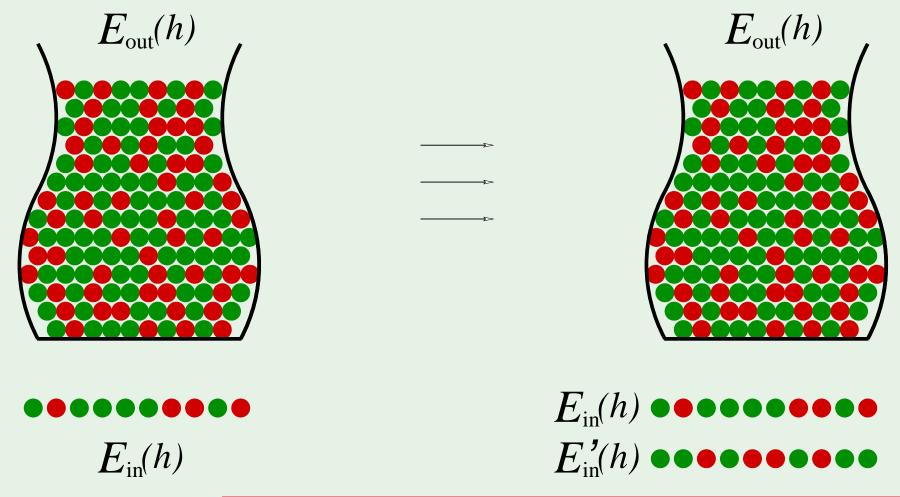
since the growth function relies on a finite sample (and the subsequent dichotomies), so it will handle the Ein aspect of Hoeffding. However Eout relates to the performance over the entire input space X and so we are dealing with full hypotheses, not dichotomies, so we lose the benefit of the growth function m.

Putting it together

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What to do about E_{out}



Ein and Ein' track eachother since they both track Eout (even if their tracking is looser) - e.g. you expect two polls of equal N to have close results to eachother. Like how the tracking of Eout and Ein become looser as the number of hypotheses increased (from M in Hoeffding), it also happens with Ein and Ein'. If we characterize this using the two samples only, no longer appealing to Eout, we are completely in the realm of dichotomies and, although the sample is bigger (2N), we can define a growth function on them - see next slide.

Putting it together

Not quite:

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

but rather:

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

The Vapnik-Chervonenkis Inequality