# COMP5212: Machine Learning

Lecture 7

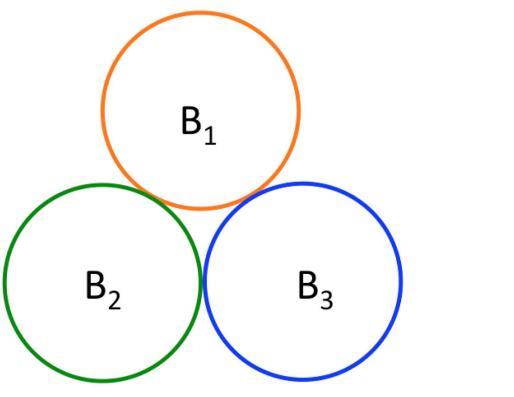
## Where did the | \mathcal{H} | come from?

- The Bad events  $\mathcal{B}_m$ :
  - $|E_{\mathsf{tr}}(h_m) E(h_m)| > \epsilon$  with probability  $\leq 2e^{-2\epsilon^2 N}$

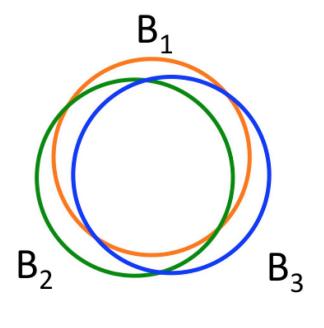
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  - $|E_{\mathsf{tr}}(h_m) E(h_m)| > \epsilon$  with probability  $\leq 2e^{-2\epsilon^2 N}$
- The union bound:

$$\mathbb{P}[\mathscr{B}_1 \text{ or } \mathscr{B}_2 \text{ or } \dots \text{ or } \mathscr{B}_M] \leq \mathbb{P}[\mathscr{B}_1] + \mathbb{P}[\mathscr{B}_2] + \dots + \mathbb{P}[\mathscr{B}_M] \leq 2 \, |\mathscr{H}| \, e^{-2\epsilon^2 N}$$
 consider worst case: no overlaps



No overlap: bound is tight



Large overlap

## A simple solution

- For each particular h,
  - $P[|E_{tr}(h) E(h)| > \epsilon] \le 2e^{-2\epsilon^2 N}$
- If we have a hypothesis set  $\mathscr{H}$ , we want to derive the bound for  $P[\sup_{h\in\mathscr{H}}|E_{tr}(h)-E(h)|>\epsilon]$ 
  - $P[|E_{tr}(h_1) E(h_1)| > \epsilon]$  or ... or  $P[|E_{tr}(h_{|\mathcal{H}|}) E(h_{|\mathcal{H}|})| > \epsilon]$
  - $\leq \sum_{m=1}^{\mathcal{H}} P[|E_{tr}(h_m) E(h_m)|] \leq 2|\mathcal{H}|e^{-2\epsilon^2 N}$ 
    - Because of union bound inequality  $P(\bigcup_{i=1}^{\infty} A_i) \leq \sum_{i=1}^{\infty} P(A_i)$

## Uniform convergence

- When our learning algorithm  $\mathcal{A}$  picks the hypothesis g:
  - $P[\exists h \in \mathcal{H} \mid E_{tr}(h) E(h) \mid > \epsilon] \le 2 \mid \mathcal{H} \mid e^{-2\epsilon^2 N}$
- Subtract both sides from 1

$$P[\neg \exists h \in \mathcal{H} | E_{tr}(h) - E(h) | > \epsilon] = P[\forall h \in \mathcal{H} | E_{tr}(h) - E(h) | \le \epsilon]$$
$$\ge 1 - 2 |\mathcal{H}| e^{-2\epsilon^2 N}$$

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- Given  $\epsilon$  and some  $\delta > 0$ , how large must N be before we can guarantee that with probability at least  $1 \delta$ , training error will be within  $\epsilon$  of generalization error?
  - Set  $\delta = 2 |\mathcal{H}| e^{-2\epsilon^2 N}$ , solve N

$$N \ge \frac{1}{2\epsilon^2} \log \frac{2|\mathcal{H}|}{\delta}$$

 The training set size N that a certain method or algorithm requires in order to achieve a certain level of performance is also called the algorithm's sample complexity

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• Given N and some  $\delta$ , we have

$$|E_{tr}(h) - E(h)| \le \sqrt{\frac{1}{2N}} \log \frac{2|\mathcal{H}|}{\delta}$$

• i.e  $|E_{tr}(h) - E(h)| \le \gamma$  for all  $h \in \mathcal{H}$ 

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- . What about the best hypothesis in training data?  $\hat{h} = \arg\min_{h \in \mathcal{H}} E_{tr}(h)$
- . Define the best hypothesis as  $h^* = \arg\min_{h \in \mathcal{H}} E(h)$
- We have  $E(\hat{h}) \leq E_{tr}(\hat{h}) + \gamma \leq E_{tr}(h^*) + \gamma \leq E(h^*) + 2\gamma$

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  - So we have

$$E(\hat{h}) \le (\min_{h \in \mathcal{H}} E(h)) + 2\sqrt{\frac{1}{2N}} \log \frac{2|\mathcal{H}|}{\delta}$$

Connection with bias/variance tradeoff

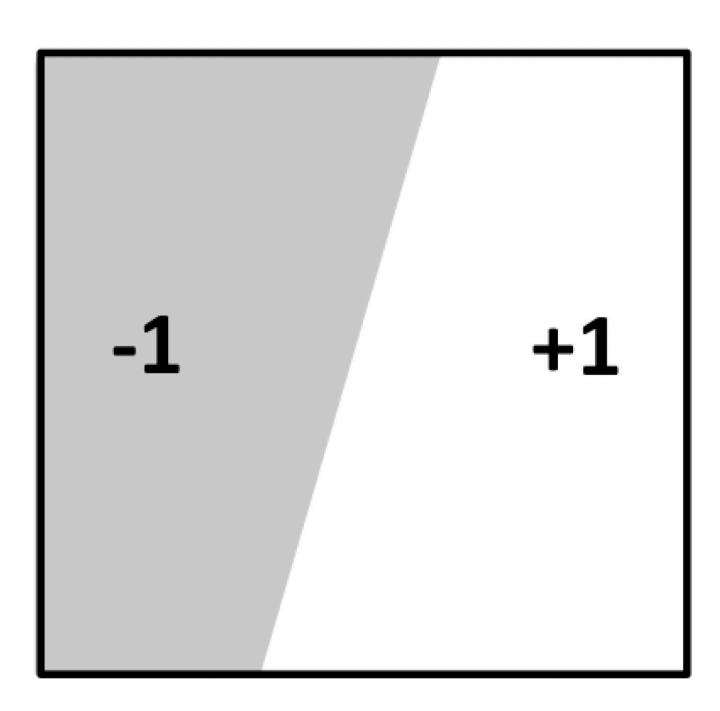
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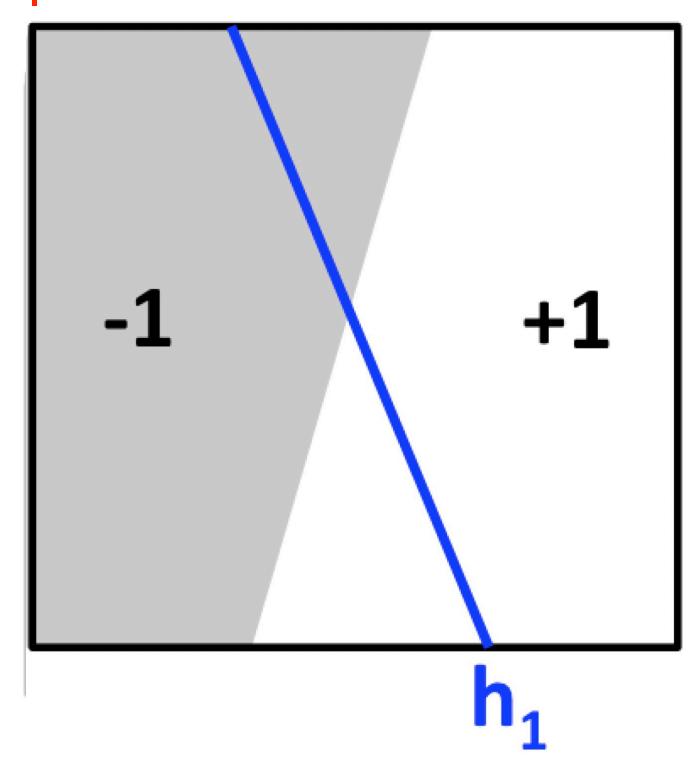
- Connection with bias/variance tradeoff
- Further, given  $\epsilon$  and some  $\delta > 0$ , is suffices that

$$N \ge \frac{1}{2\epsilon^2} \log \frac{2|\mathcal{H}|}{\delta} = O(\frac{1}{\epsilon^2} \log \frac{|\mathcal{H}|}{\delta})$$

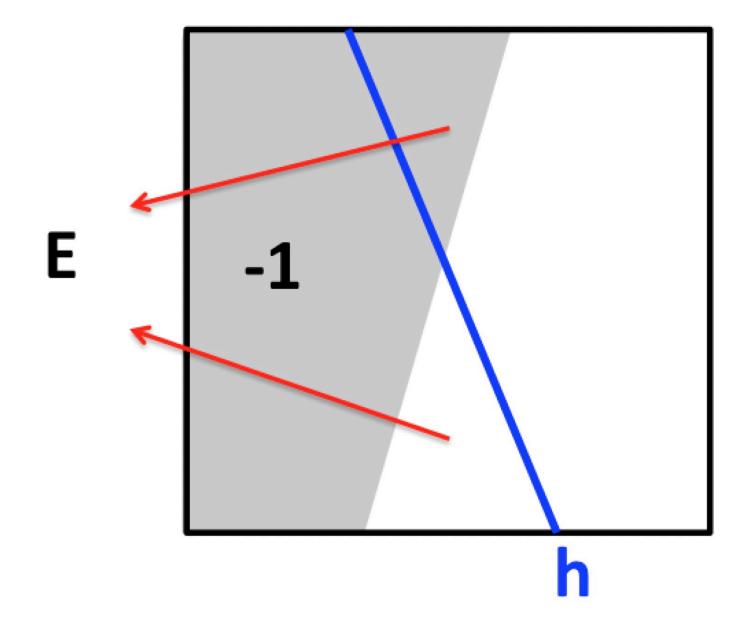
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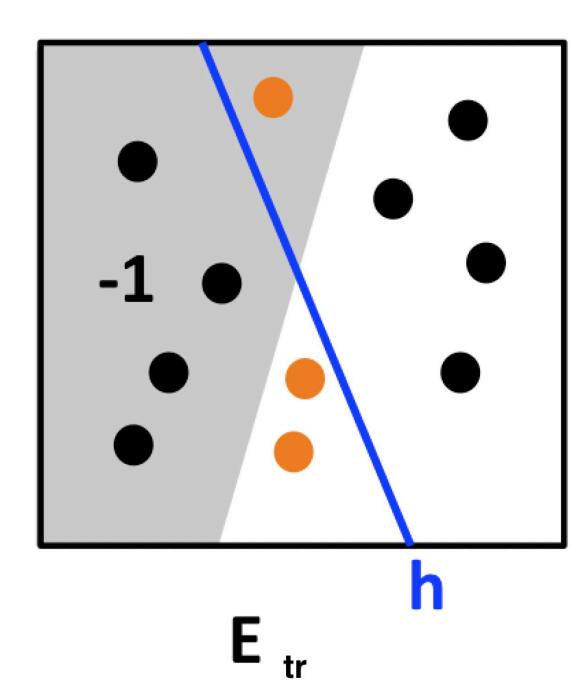


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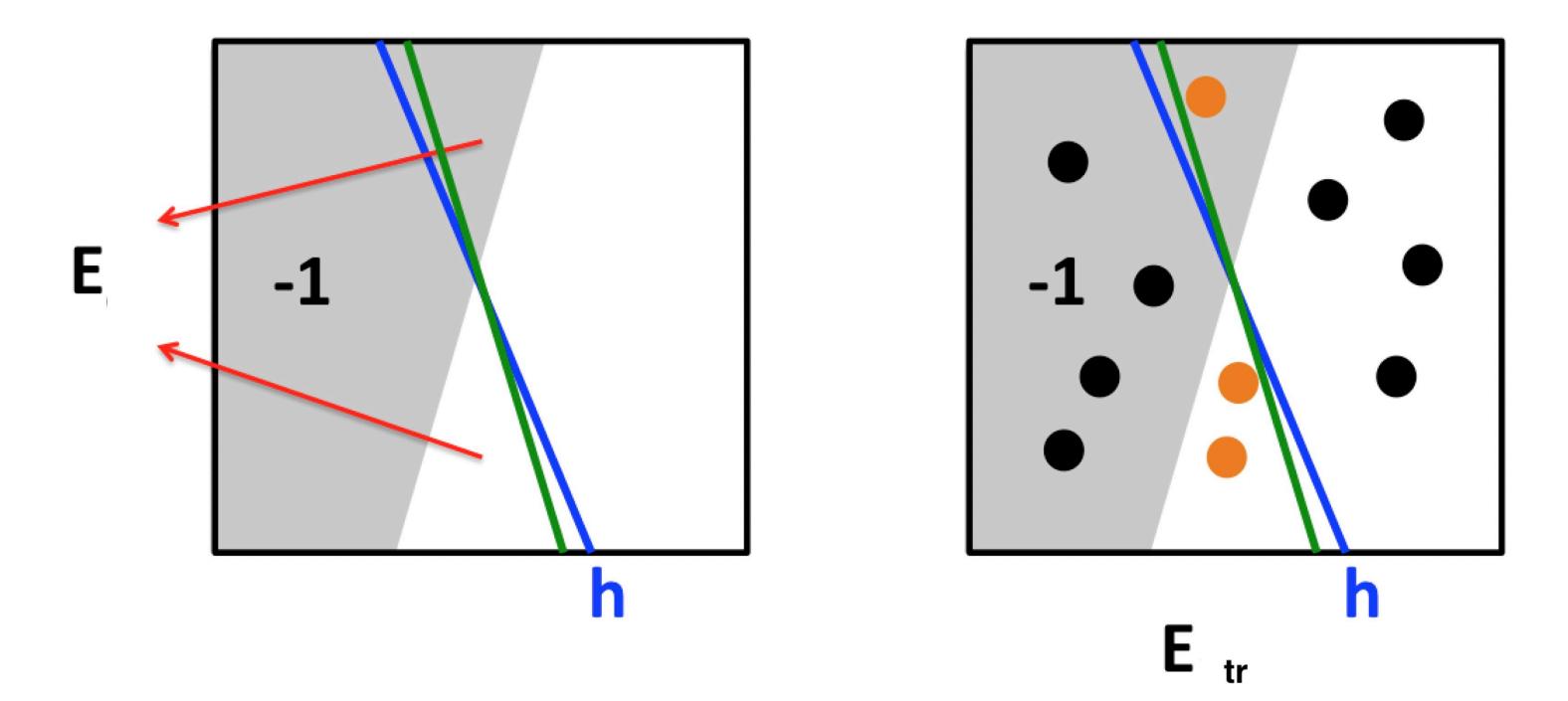


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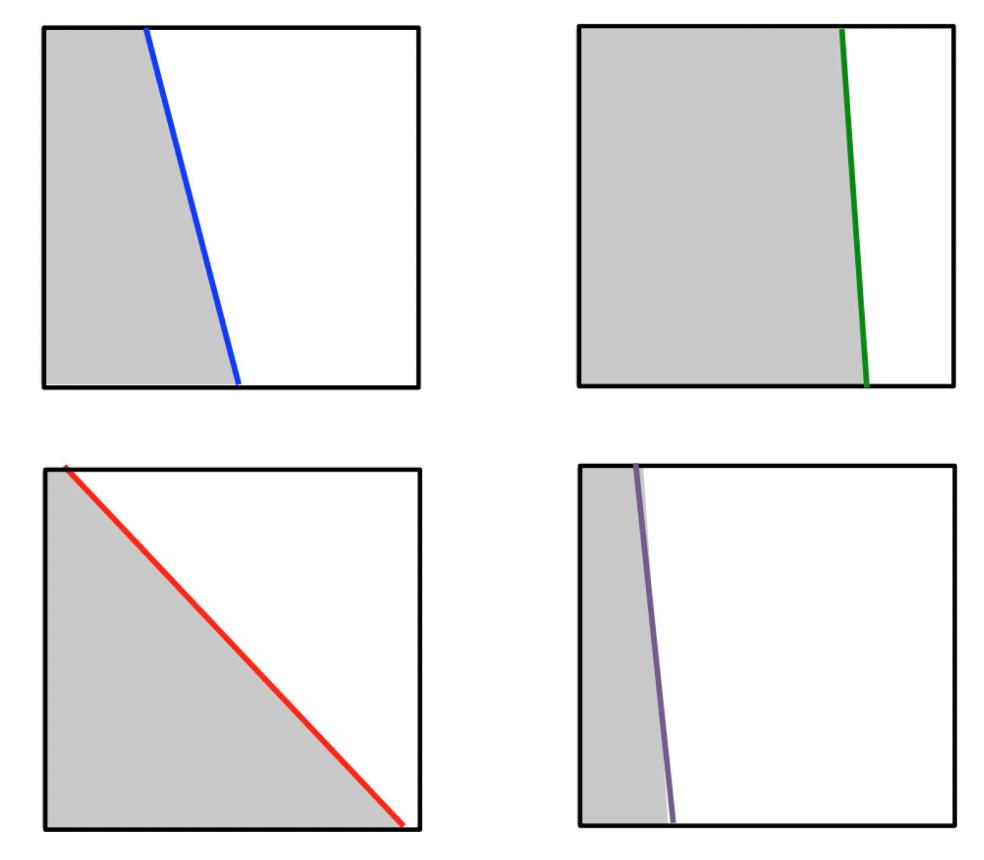
## Can we improve on | % | ?



• The event that  $|E_{\rm tr}(h_1)-E(h_1)|>\epsilon$  and  $|E_{\rm tr}(h_2)-E(h_2)|>\epsilon$  are largely overlapped

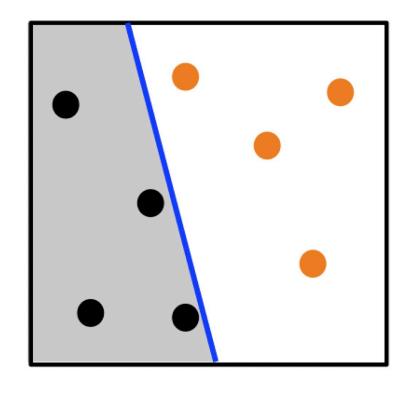
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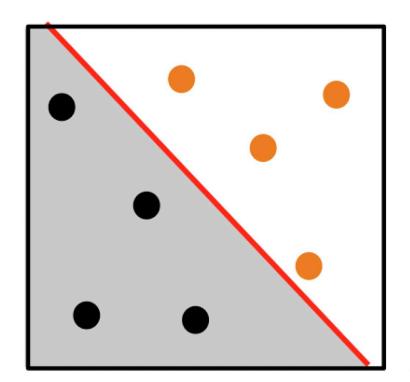
Instead of the whole input space

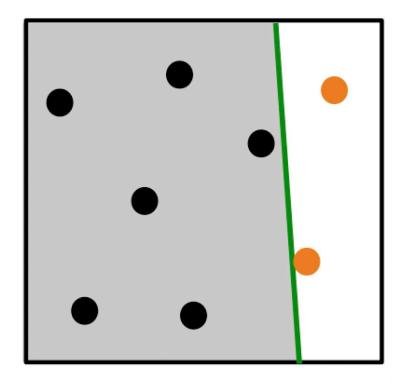


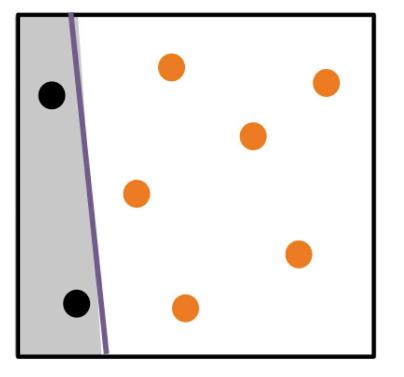
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- Instead of the whole input space
- Let's consider a finite set of input points



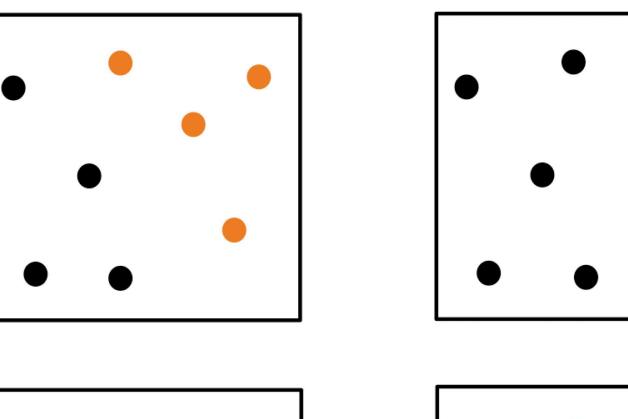


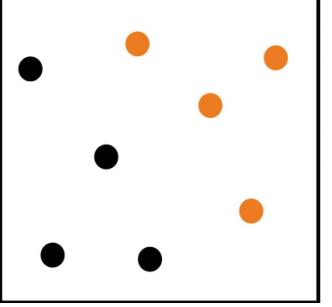


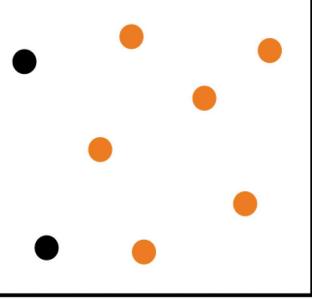


## What can we replace | \mathcal{H} | with?

- Instead of the whole input space
- Let's consider a finite set of input points
- How many patterns of colors can you get?







### Dichotomies: mini-hypotheses

- A hypothesis:  $h: \mathcal{X} \to \{-1, +1\}$
- A dichotomy:  $h: \{x_1, x_2, ..., x_N\} \rightarrow \{-1, +1\}$

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  - $\Rightarrow$  Candidate for replacing  $\mathscr{H}$
  - Why?

## Symmetrization lemma

• Imagine we have the ghost dataset S' with also size N:

$$P[SUP_{h\in\mathcal{H}}|E_{tr}(h) - E(h)| > \epsilon] \le 2P[SUP_{h\in\mathcal{H}}|E_{tr}(h) - E'_{tr}(h)| > \frac{\epsilon}{2}]$$

### **Growth function**

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- By union bound:

• 
$$P[SUP_{h \in \mathcal{H}_{S \cup S'}} | E_{tr}(h) - E'_{tr}(h) | > \frac{\epsilon}{2}] \le |\mathcal{H}_{S \cup S'}| P[|E_{tr}(h) - E'_{tr}(h)| > \frac{\epsilon}{2}]$$

### **Growth function**

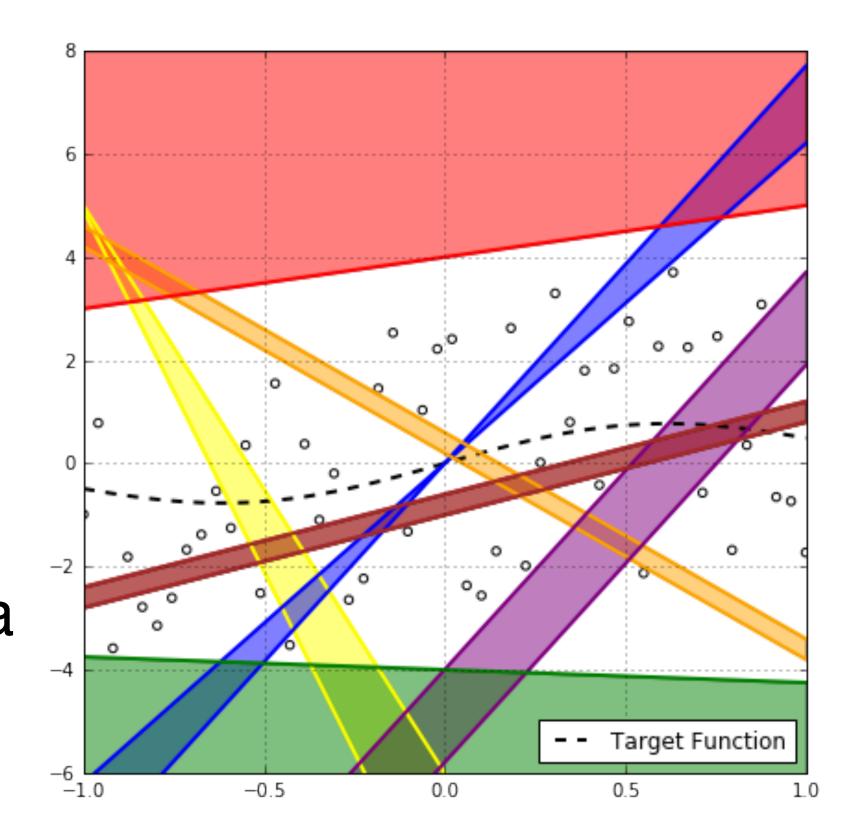
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• How to bound  $|\mathcal{H}_{S \cup S'}|$ 

### **Deduce the dimension**

- Why do we need to consider every possible hypothesis?
  - $P[SUP_{h\in\mathcal{H}}|E_{tr}(h)-E(h)|>\epsilon]$
  - If we omit one hypothesis, we might miss the biggest gap
- However, are the events of each hypothesis having a big generalization gap are likely to be independent?
  - No



## The growth function

The growth function counts the most dichotomies on any N points:

$$m_{\mathcal{H}}(N) = \max_{\substack{x_1, \dots, x_N \in \mathcal{X}}} |\mathcal{H}(x_1, \dots, x_N)|$$

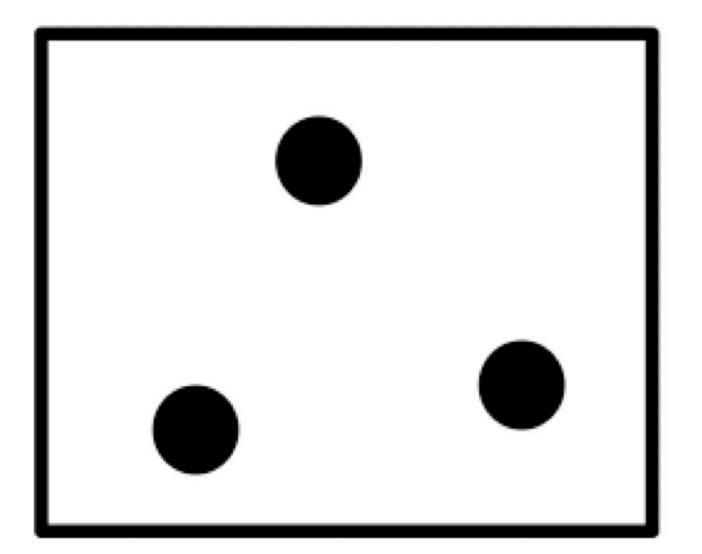
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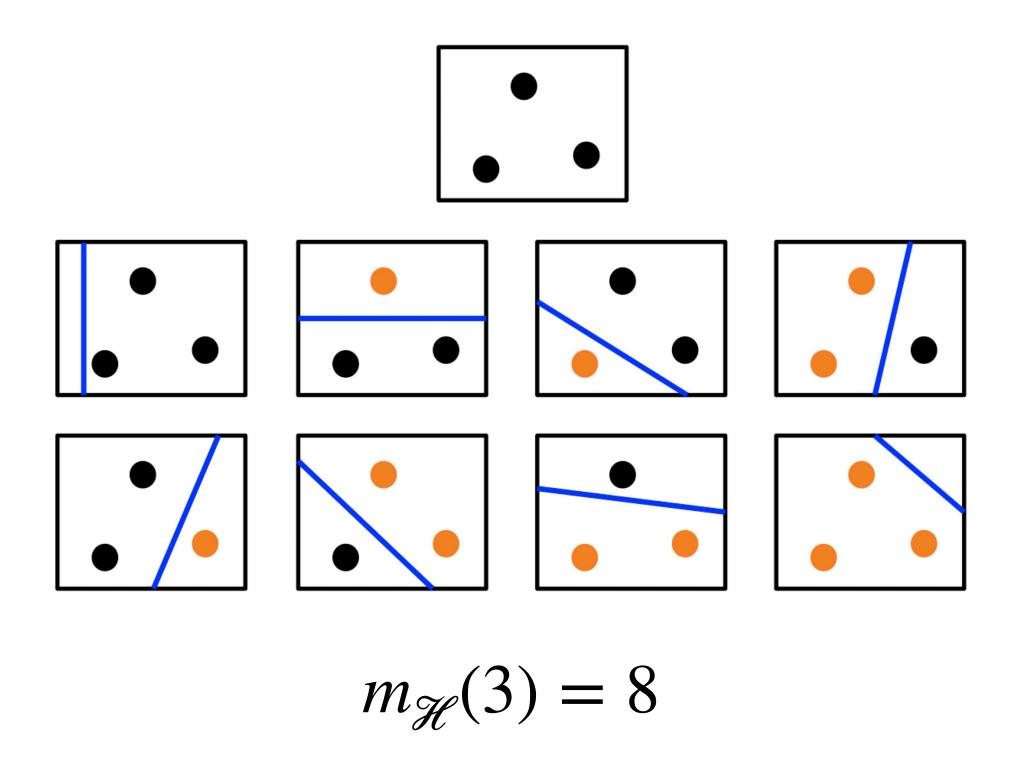
- The growth function satisfies:
  - $m_{\mathcal{H}}(N) \leq 2^N$

• Compute  $m_{\mathcal{H}}(3)$  in 2-D space

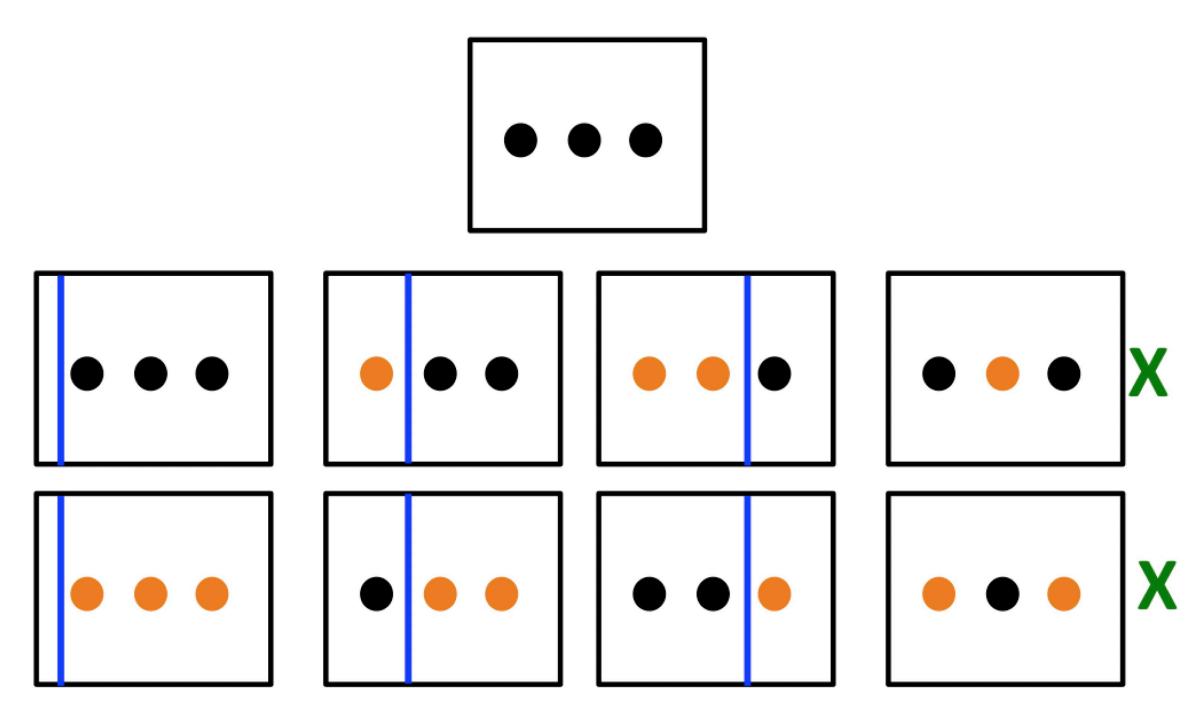


• What's  $|\mathcal{H}(x_1, x_2, x_3)|$ ?

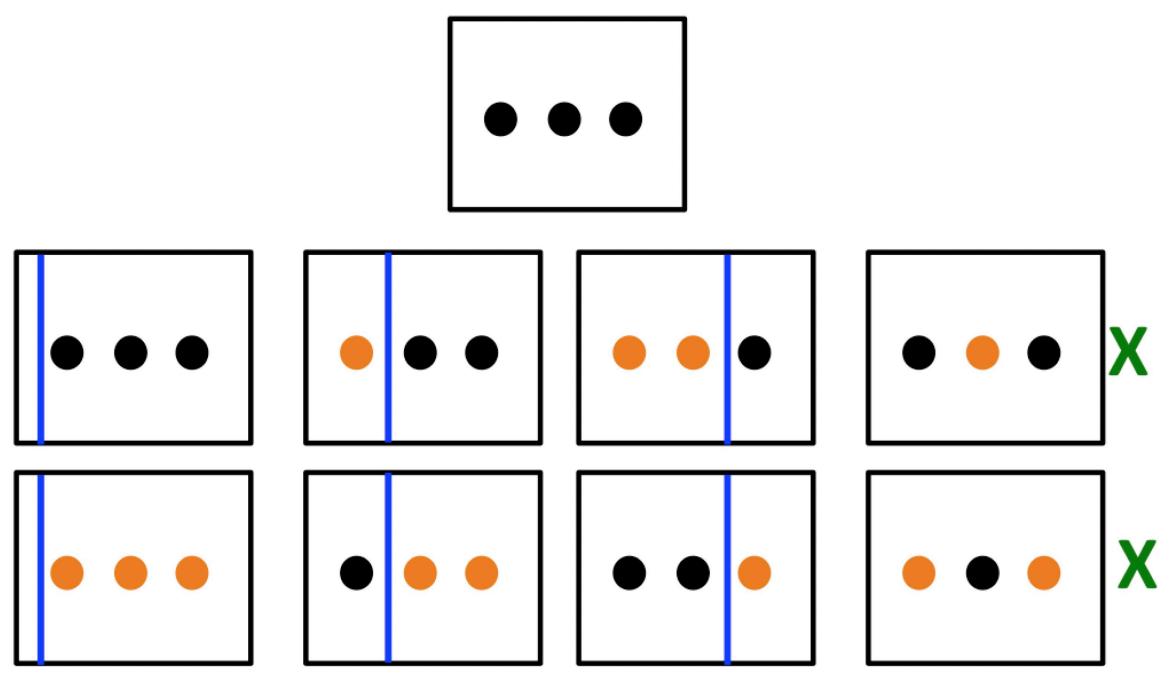
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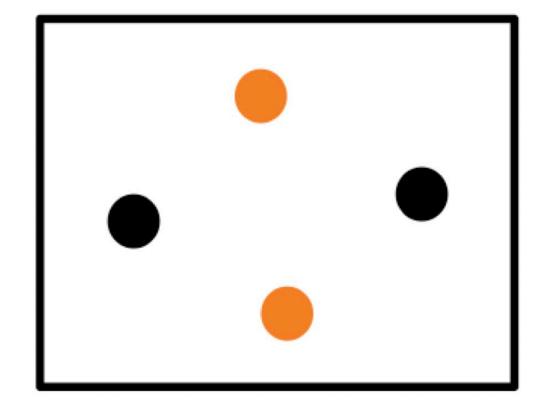
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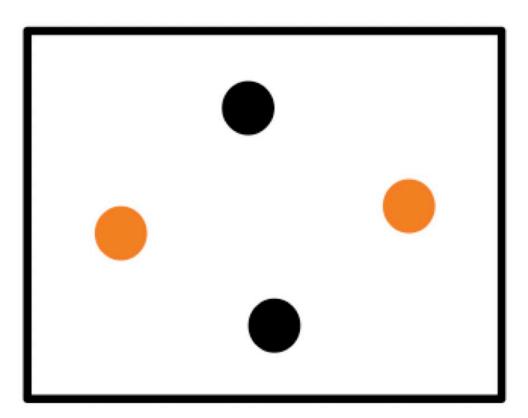


Doesn't matter because we only counts the most dichotomies

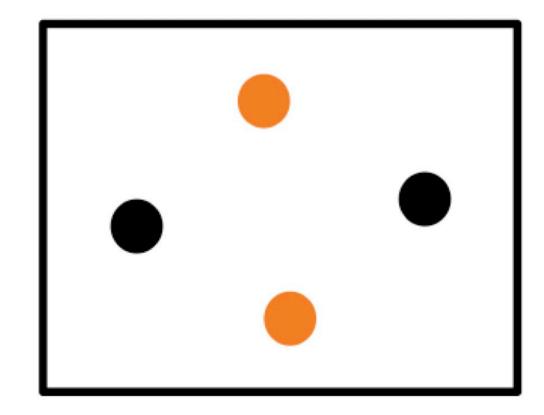
• What's  $m_{\mathcal{H}}(4)$ ?

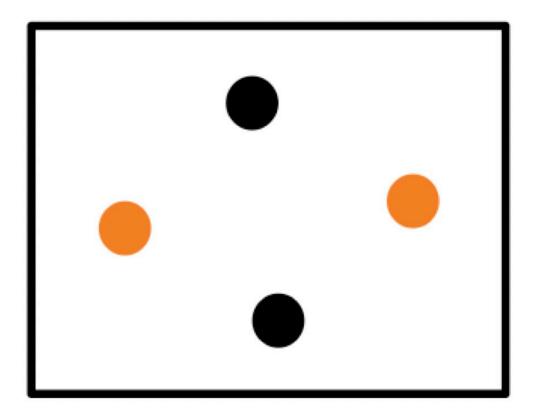
- What's  $m_{\mathcal{H}}(4)$ ?
- (At least) missing two dichotomies:





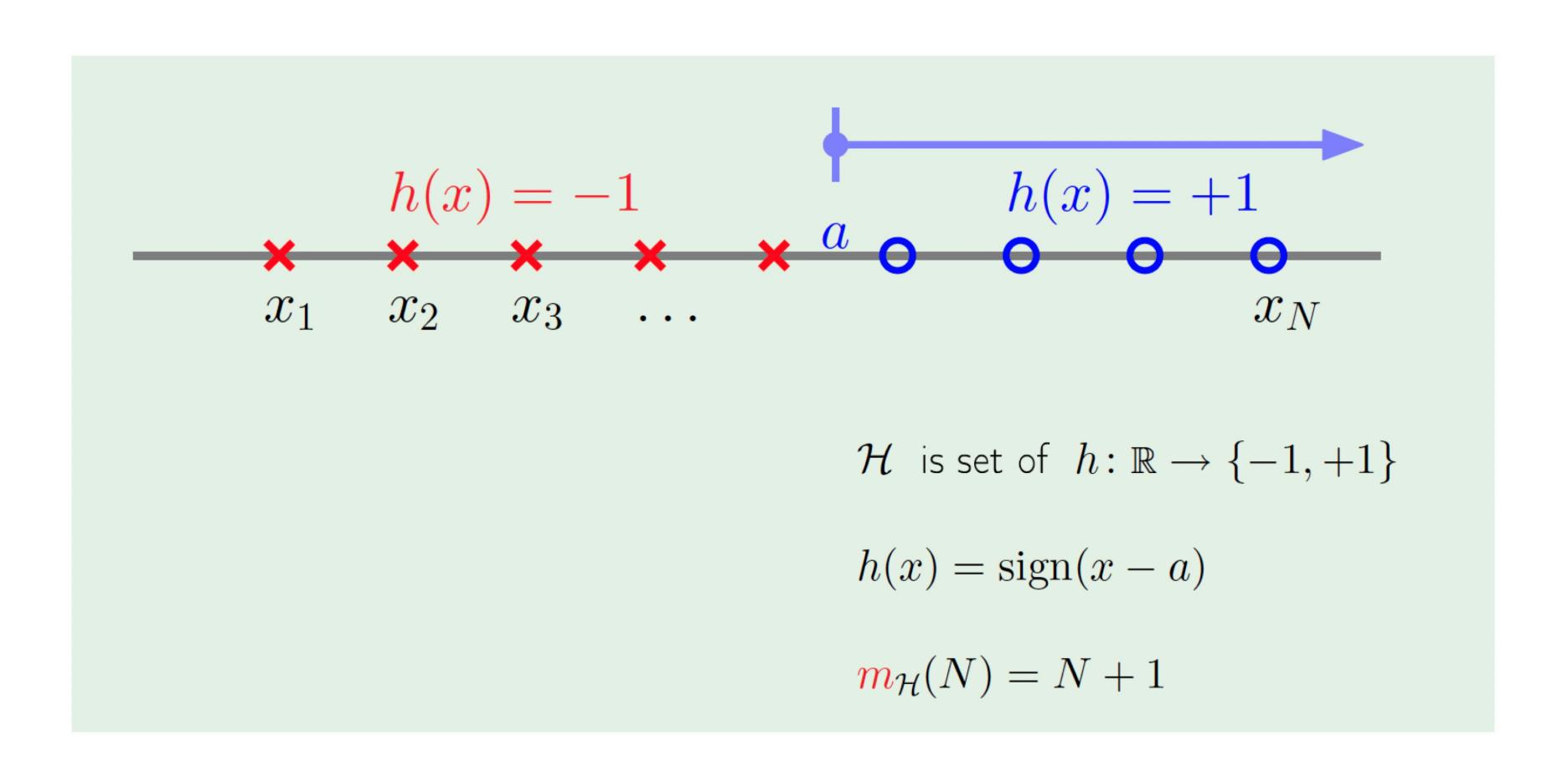
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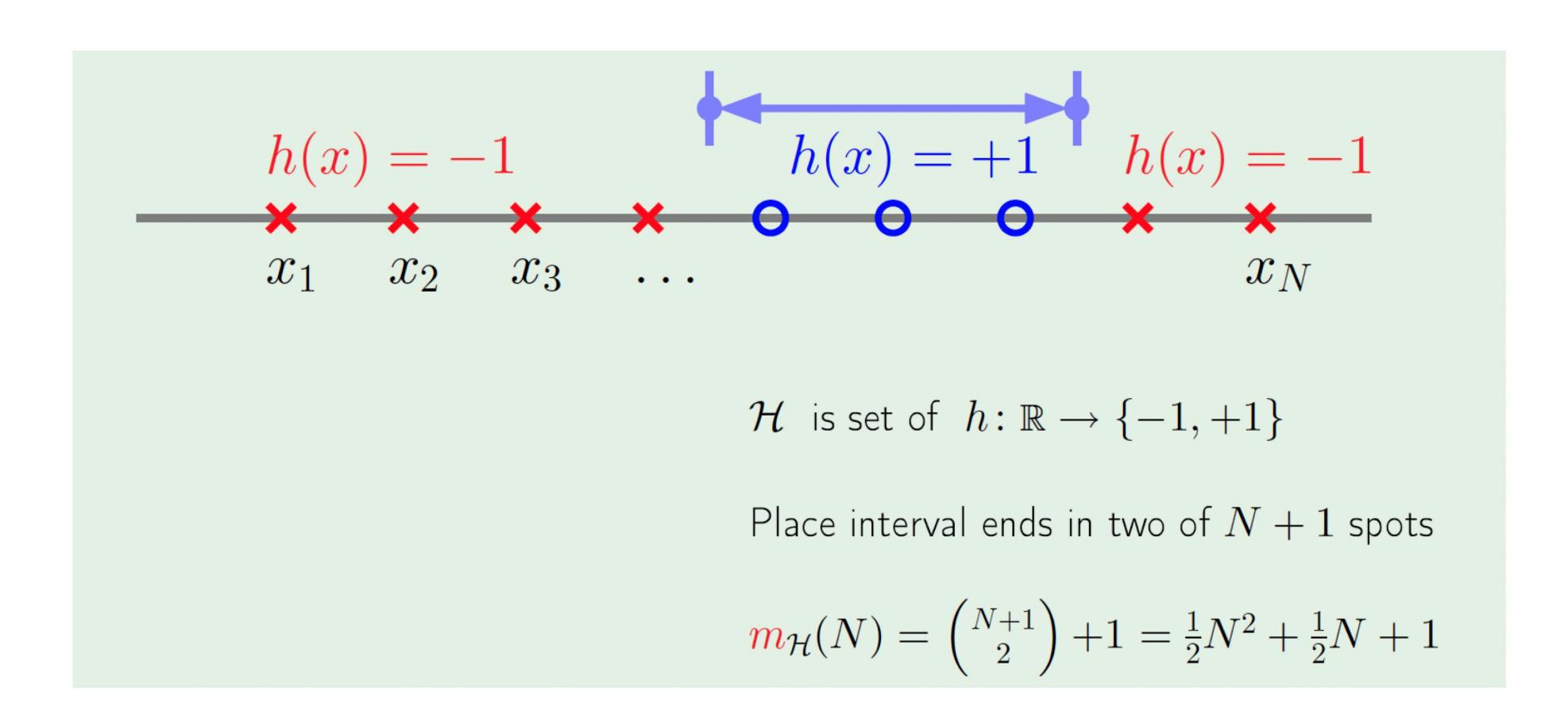


• 
$$m_{\mathcal{H}}(4) = 14 < 2^{2}$$

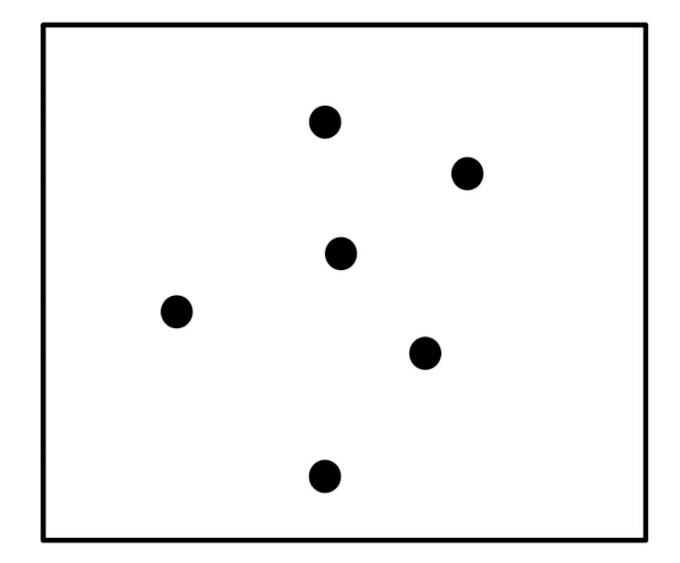
## **Example I: positive rays**



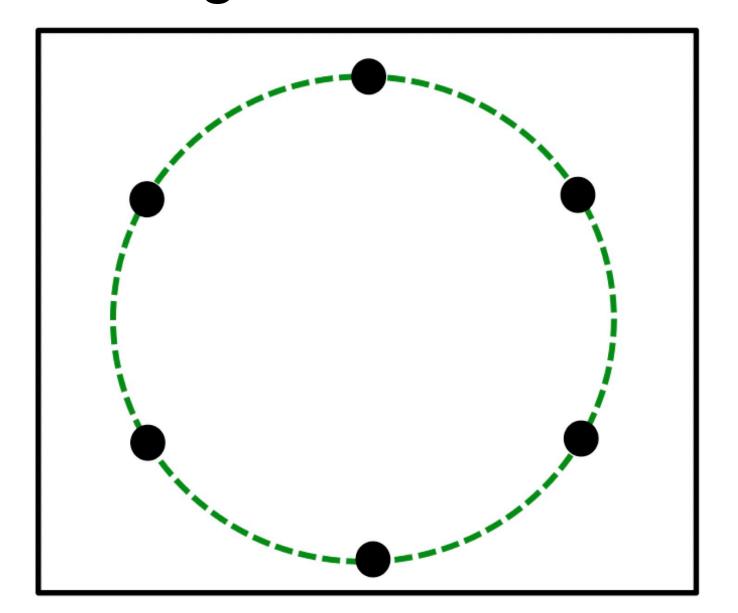
## **Example II: positive intervals**



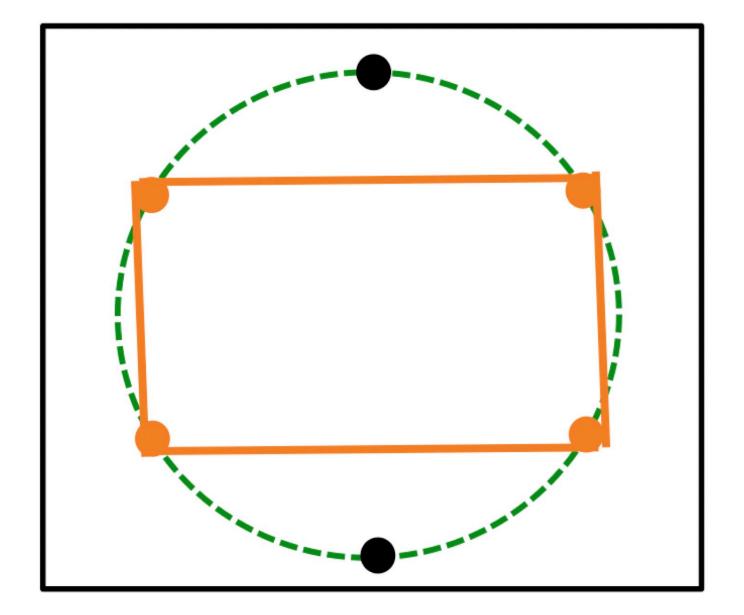
- $\mathcal{H}$  is set of  $h: \mathbb{R}^2 \to \{-1, +1\}$ 
  - h(x) = +1 is convex
- How many dichotomies can we generate?



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- $\mathcal{H}$  is set of  $h: \mathbb{R}^2 \to \{-1, +1\}$ 
  - h(x) = +1 is convex
- $m_{\mathscr{H}}(N) = 2^N$  for any  $N \Rightarrow We$  say the N points are "shattered" by h

### **Shattered**

• Given a set  $S = \{x^{(i)}, \dots, x^{(d)}\}$  (no relation to the training set) of points  $x^{(i)} \in \mathcal{X}$ , we say that  $\mathcal{H}$  shatters S if  $\mathcal{H}$  can realize any labeling on S. I.e, if for any set of labels  $\{y^{(i)}, \dots, y^{(d)}\}$ , there exist some  $h \in \mathcal{H}$  so that  $h(x^{(i)}) = y^{(i)}$  for all  $i = 1, \dots, d$ 

## The 3 growth functions

- *H* is positive rays:
  - $m_{\mathcal{H}}(N) = N + 1$
- $\mathcal{H}$  is positive intervals:

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

- $\mathcal{H}$  is convex sets:
  - $m_{\mathcal{H}}(N) = 2^N$

### What's next?

- Remember the inequality
  - $\mathbb{P}[|E_{\mathsf{in}} E_{\mathsf{out}}| > \epsilon] \le 2 |\mathcal{H}| e^{-2\epsilon^2 N}$
- What happens if we replace  $|\mathcal{H}|$  by  $m_{\mathcal{H}}(N)$ 
  - $m_{\mathcal{H}}(N)$  polynomial  $\Rightarrow$  Good!

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  - Why?
- How to show  $m_{\mathcal{H}}(N)$  is polynomial?