

1a) So for any  $x \dots$

$$f(x)g(x) \begin{cases} +1, & f(x) = g(x) \\ -1, & f(x) \neq g(x) \end{cases}$$

So we can write

$$1 \{ f(x) \neq g(x) \} = \frac{1 - f(x)g(x)}{2}$$

w.r.t.  $x$  is sampled according to  $D$

$$\mathbb{P}_{x \sim D}[f(x) \neq g(x)] = \mathbb{E}_{x \sim D}[1 \{ f(x) \neq g(x) \}] = \mathbb{E}_{x \sim D}\left[\frac{1 - f(x)g(x)}{2}\right] = \frac{1 - \mathbb{E}_{x \sim D}[f(x)g(x)]}{2}$$

or written as  $\mathbb{E}_{x \sim D}[f(x)g(x)] = 1 - 2\mathbb{P}_{x \sim D}[f(x) \neq g(x)]$

1b) Yes! The statement holds for any domain  $X$  and any  $D$  on  $X$  provided  $f, g: X \rightarrow \{-1, 1\}$ . Reason the identity which depends only on the range  $\{-1, 1\}$  not on the domain.

2) So to build one polynomial tree per leaf. Lets start with indicator for variable test.

▷ if branch checks  $x_i = 1$ , use

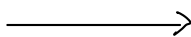
$$\frac{1 + x_i}{2} \quad \text{so that it equals } = 1 \text{ when } x_i = 1 \\ = 0 \text{ when } x_i = -1$$

▷ if branch checks  $x_i = -1$ , use

$$\frac{1 - x_i}{2} \quad \text{so that it equals } = 1 \text{ when } x_i = -1 \\ = 0 \text{ when } x_i = 1$$

So for indicator path we would get:

$$I_{\text{path}}(x) = \prod_{j=1}^k \frac{1 + a_j x_j}{2}$$



Lets attach leaf values  $b_\ell \in \{-1, 1\}$

$$P_\ell(x) = b_\ell \cdot I_{\text{path}_\ell}(x)$$

- $a_j$  is required value ( $\pm 1$ ) for  $x_j$ 
  - ↳ if all conditions are met every factor is 1, so the product is 1
  - ↳ if even one condition fails, one factor becomes 0, so the product is 0.
- On/off switch.

So this polynomial outputs the leaf's value  $\neq$  if on path  $\neq$  or else 0.

Finally, sum over all  $\ell$  leaves:

$$p(x) = \sum_{\ell=1}^L P_\ell(x).$$

At any input  $x$ , exactly one path indicator equals to 1.

↳ all others are 0. So therefore sum reduces to  $p(x) = f(x)$

3)

Totals:  $P_+ = 165$ ,  $N_+ = 85$ ,  $T_+ = 250$ Let's start with Root Node. For  $F \in \{X, Y, Z\}$ , group them by  $F=0$ ,  $F=1$ .Compute  $a = \frac{\#Pos}{\#Total}$  then  $C(F=v) = 2a(1-a)$ . Split score is weighted sum by group size.

$$C(P_r[Pos]) = 2 \cdot \frac{165}{250} \cdot \frac{85}{250} = 0.4488$$

## ① Root Split

→ Split on X

$$\bullet X=0: Pos=105, neg=45 \rightarrow a = \frac{105}{150}, C=0.42$$

$$\bullet X=1: Pos=60, neg=40 \rightarrow a = \frac{60}{100}, C=0.48 \quad \text{Weighted: } 0.444$$

→ Split on Y

$$\bullet Y=0: Pos=70, neg=50 \rightarrow a = \frac{70}{120}, C=0.467$$

$$\bullet Y=1: Pos=95, neg=35 \rightarrow a = \frac{95}{130}, C=0.391 \quad \text{Weighted: } 0.438$$

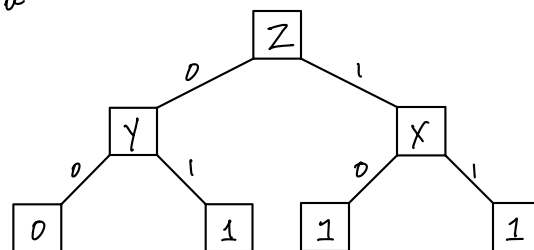
→ Split on Z

$$\bullet Z=0: Pos=60, neg=60 \rightarrow a = \frac{60}{120}, C=0.50$$

$$\bullet Z=1: Pos=105, neg=25 \rightarrow a = \frac{105}{130}, C=0.311 \quad \text{Weighted: } 0.402$$

Smallest score → root split on Z

Tree



## ② Depth two Split

Left child  $Z=0$  (Pos: 60, Neg: 60,  $N=120$ )

→ Split on X

$$\bullet X=0 (45, 35) \quad C=0.459$$

$$\bullet X=1 (15, 25) \quad C=0.469 \quad W=0.484$$

→ Split on Y

$$\bullet Y=0 (15, 35) \quad C=0.420$$

$$\bullet Y=1 (45, 25) \quad C=0.459 \quad W=0.443$$

Smallest Score → split on Y

→ Leaves:

$$Z=0, Y=0: (15, 35) \rightarrow P_{neg}$$

$$Z=0, Y=1: (45, 25) \rightarrow P_{pos}$$

Right child:  $Z=1$  (Pos: 105, Neg: 25,  $N=130$ )

→ Split on X

$$\bullet X=0 (60, 10) \rightarrow C=0.245$$

$$\bullet X=1 (45, 15) \rightarrow C=0.278 \quad W=0.305$$

→ Split on Y

$$\bullet Y=0 (55, 15) \rightarrow C=0.321$$

$$\bullet Y=1 (50, 10) \rightarrow C=0.278 \quad W=0.310$$

Smallest Score → Split on X

→ Leaves:

$$Z=1, X=0 (60, 10) \rightarrow P_{pos}$$

$$Z=1, X=1 (45, 15) \rightarrow P_{pos}$$

## ③ Final

• Root Split on Z

↳  $Z=0$ : Split on Y↳  $Y=0 \rightarrow Neg$  $Y=1 \rightarrow Pos$ ↳  $Z=1$ : Split on X↳  $X=0 \rightarrow Pos$  $X=1 \rightarrow Pos$ 

## ④ Training accuracy

$$\bullet Z=0, Y=0: 35$$

$$\bullet Z=0, Y=1: 45$$

$$\bullet Z=1, X=0: 60$$

$$\bullet Z=1, X=1: 45$$

$$T_c = 185$$

$$Acc = \frac{185}{250} = 0.74 \text{ or } \underline{\underline{74\%}}$$

#### 4) Pac Learning Algorithm

Training set  $S = \{x_1, y_1, \dots, x_m, y_m\}$  where  $x_i \in \mathbb{R}$   $y_i \in \{-1, 1\}$

① Input:  $m$  labeled point  $S = \{(x_i, y_i)\}$  with  $y_i \in \{-1, 1\}$

② set  $\hat{\theta} = \text{"no value"}$  \*  $\hat{\theta}$  - learned threshold.

③ Scan the sorted list from L or R:

if  $y_j = -1$ , set  $\hat{\theta} = x_j$  \* Keep largest  $x$  with label  $-1$

④ if  $\hat{\theta}$  has no value (there was no  $-1$  in  $S$ )

set  $\hat{\theta} = x_1 - 1$  \* makes all points predict  $1$

⑤ output classifier  $h_{\theta}$ :

$h_{\theta}(x) = -1$  if  $x \leq \theta$ , else  $+1$

Time: sort  $O(m \log m)$ , scan  $O(m)$

Assume there is some true threshold that labels the data correctly. Our algorithm chooses the learned threshold as the largest sample with label  $-1$ . Only mistakes can happen between learned and true threshold. if we get at least one training point close enough to true threshold, then total error is at most  $\epsilon$ . The chance that no point lands in that region is at most  $(1-\epsilon)^m$ , which is smaller than  $e^{-\epsilon m}$ . To make this failure probability at most  $\delta$  it is enough to take

$$m \geq \frac{1}{\epsilon} \ln \frac{1}{\delta}$$

so with probability at least  $1-\delta$ , the algorithm finds a classifier whose error is at most  $\epsilon$ .

5a) if  $\text{er}(h') > \epsilon$ , each fresh example is correct with prob  $< 1-\epsilon$ .

for a block of size  $k$ :

$$\begin{aligned} \Pr[h' \text{ makes 0 mistakes on the block}] \\ = (1-\epsilon)^k \leq e^{-\epsilon k} \end{aligned}$$

Pick  $k > \frac{1}{\epsilon} \ln \frac{1}{\delta}$ . Then

$$\Pr[\text{er}(h') > \epsilon \text{ and } h' \text{ passes its block}] \leq \delta$$

5b) Learner A makes at most  $t$  mistakes, so it visits at most  $t+1$  hypotheses  $h_1, \dots, h_{t+1}$ .

Use union bound with  $\delta' = \frac{\delta}{(t+1)}$

Choose

$$k \geq \frac{1}{\epsilon} \ln \frac{t+1}{\delta}$$

#### 5c) PAC Learner B (uses A as subroutine)

① Compute block size  $k$  as above.

② Draw  $m = (t+1) \cdot k$  training examples.

③ Split them into  $t+1$  blocks, each of size  $k$ .

④ Start with A's first hypothesis  $h_1$ .

⑤ for each block  $j=1, \dots, t+1$ :

- Test hypothesis  $h_j$  on the block  $j$ .
- if it makes no mistakes, output  $h_j$  and stop.
- if it makes a mistake, feed one mistake back to A so it updates to  $h_{j+1}$ .

⑥ if none pass, output the last hypothesis.

#### NOTE

- Each bad hypothesis passes its block with probability  $\leq \delta / (t+1)$
- with at most  $t+1$  hypotheses, union bound gives failure probability  $\leq \delta$
- So with probability  $\geq 1-\delta$ , the output has error  $\leq \epsilon$ .

Samples used:

$$m = (t+1) \cdot k = \frac{t+1}{\epsilon} \ln \frac{t+1}{\delta}$$