

Lecture 15.0

Hypothesis Testing

Some Problems: (Given samples of  $p$ )

Complexity (in terms of  $n = |D|$ )

is  $p = q$  (e.g.  $q = U_D$ )  
or  $\epsilon$ -far from  $q$

$\sqrt{n}$

is  $p$   $\epsilon$ -close to  $q$   
or  $\epsilon$ -far from  $q$

$\frac{n}{\log n}$

(Given samples of  $q$ ) is  $p = q$   
or  $p$   $\epsilon$ -far from  $q$

$n^{2/3}$

(Given samples of  $q$ ) is  $p$   $\epsilon$ -close to  $q$   
or  $\epsilon$ -far from  $q$

$\frac{n}{\log n}$

is  $p$  monotone  
or  $\epsilon$ -far from monotone

$\sqrt{n}$

is  $p$   $\epsilon$ -close to monotone  
or  $\epsilon$ -far from monotone

$n/\log n$

Other problems considered:

estimate entropy, support size

Independence?

represented well via k-histogram?

monotone hazard rate

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A useful tool:

Given: (1) collection of distributions (via complete description)  $\mathcal{H}$

(2) Samples of  $p$  such that  $\exists q \in \mathcal{H}$  for which  $\text{dist}(p, q)$  is small

$\mathcal{H}$  contains a good approx to  $p$

← Strong assumption

Goal: Output  $h \in \mathcal{H}$  s.t.  $\text{dist}(p, h)$  small

Question:

How many samples needed in terms of  $|\mathcal{H}|$  + domain size?

Is this the same as testing closeness, uniformity?

Do lower bounds apply?

**NO!**

$\left\{ \begin{array}{l} p \text{ is guaranteed to} \\ \text{be close to some } f \in \mathcal{H} \end{array} \right.$

What we want:

Given  $h_1, h_2$  explicit  
 $p$  via samples

procedure that outputs  $h_i$  that is closer to  $p$

What if both are roughly same distance?

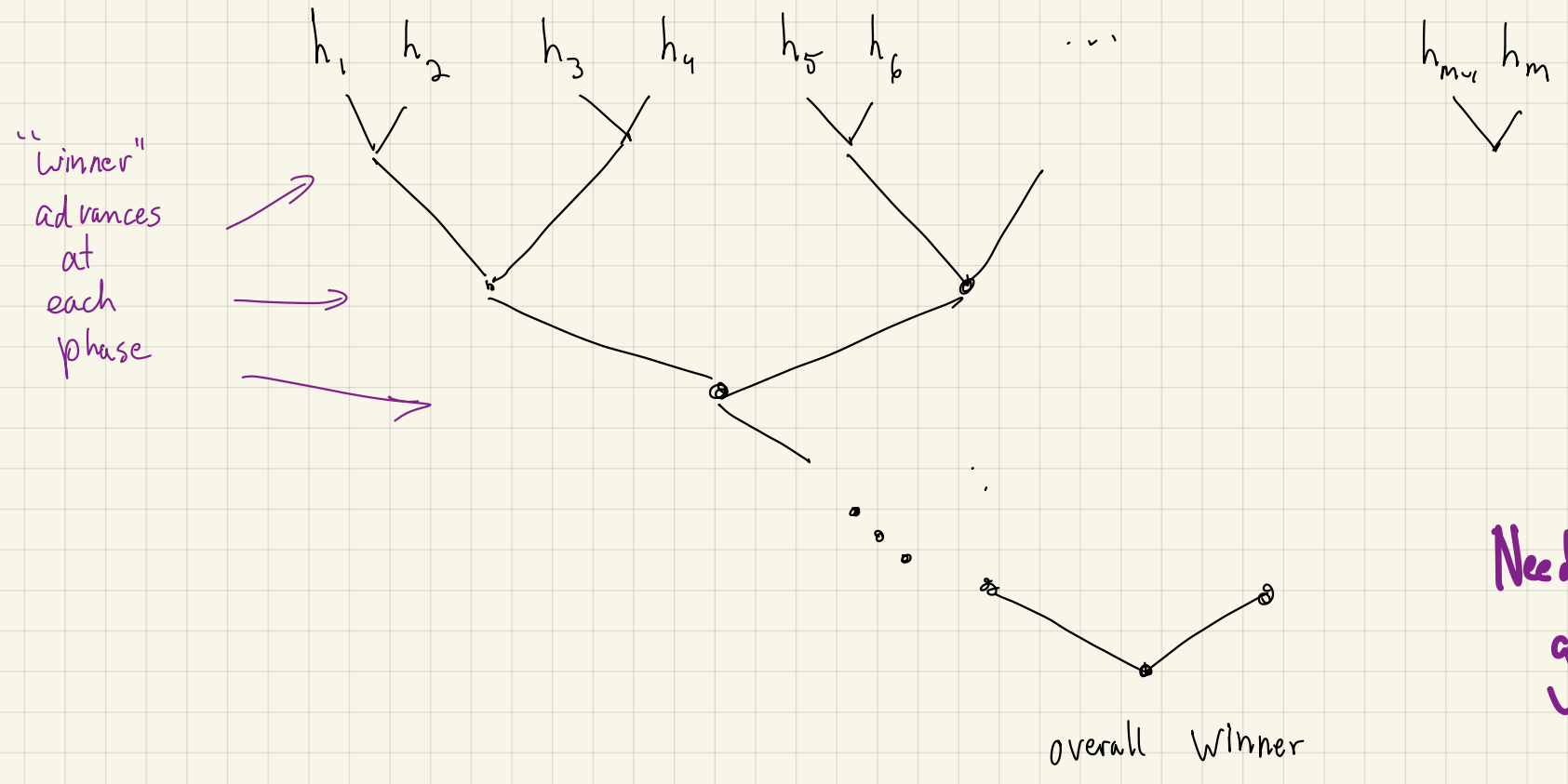
maybe either one is ok?

or maybe not...

More general Goal:

Given set of hypotheses  $\mathcal{H}$   
 $\dagger$   $p$  via samples  
find  $h \in \mathcal{H}$  closest to  $p$

Find best hypothesis via "tournament"?



maybe  $p = h_1$   
 $\|p - h_2\|_1 = \epsilon$  &  $h_2$  "wins"  
 then  $\|p - h_3\|_1 = 2\epsilon$  &  $h_3$  "wins"  
 then  $\|p - h_5\|_1 = 3\epsilon$  &  $h_5$  "wins"  
 ...

overall winner could be  $O(\log n \cdot \epsilon)$   
 far from best hypothesis?

- won't use simple tournament ← instead compare every pair
- will add notion of "tie"

Output hypothesis that wins or ties  
every match

(hopefully there is one, & it is the  
right one)

# A "subtool" for comparing two hypotheses:

- Thm given (1) sample access to  $p$   
(2)  $h_1, h_2$  hypothesis distributions (fully known to algorithm)  
(3) accuracy parameter  $\epsilon'$ , confidence parameter  $\delta'$

then Algorithm "choose" takes  $O(\log(\frac{1}{\delta'}) / (\epsilon')^2)$  samples + outputs  $h \in \{h_1, h_2\}$  satisfying:

if one of  $h_1, h_2$  has  $\|h_i - p\|_1 < \epsilon'$

then with prob  $\geq 1 - \delta'$ , output  $h_j$  has  $\|h_j - p\|_1 < 12\epsilon'$

ie. if both  $h_1, h_2$  far, no guarantees  
if one  $\epsilon'$ -close + one really far, will output  $\epsilon'$ -close hypothesis }  
if both  $\epsilon'$ -close then output  $12\epsilon'$ -close hypothesis } if at least one is close, will output pretty close hypothesis

e.g.  $\uparrow$  one is  $\epsilon'$ -close  
other is  $\leq 10\epsilon'$ -close

e.g.  $\rightarrow$   $\geq 12\epsilon'$

getting kind of complicated just to specify (??)



Actually a bit stronger:

Thm  $p$  given via samples  
 $h_1, h_2$  fully known &  $p$  is  $\epsilon'$ -close to at least one of  $h_1, h_2$   
 $\epsilon', \delta'$  given

Algorithm "choose" takes  $O((\log \frac{1}{\delta'}) (\frac{1}{\epsilon'})^2)$  samples & outputs  $h \in \{h_1, h_2\}$  such that:

(1) If  $h_i$  more than  $12\epsilon'$ -far from  $p$ , very bad unlikely to output  $h_i$  as winner or tie  
 $2e^{-m(\epsilon')^2/2}$

(2) If  $h_i$  more than  $10\epsilon'$ -far from  $p$ , not that bad unlikely to output  $h_i$  as winner  
 $\uparrow$  might tie but won't win

Can use  $\epsilon' \approx \frac{\epsilon}{10}$  ?

## Proof of subtool:

$h_1 + h_2$  are close  
can determine  $h_1 + h_2$  close w/o samples  
from  $p$

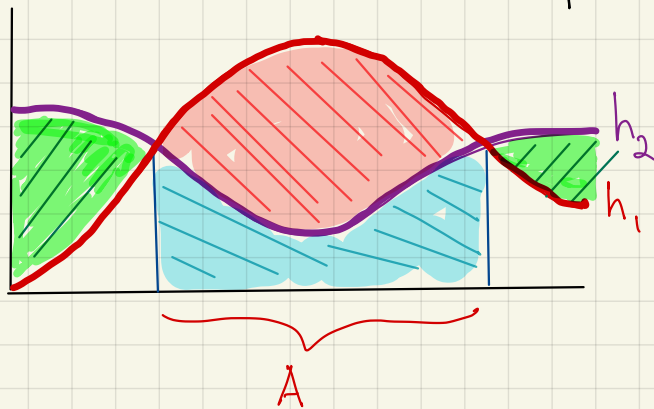
idea: wlog  $h_1$  is  $\epsilon'$ -close to  $p$

if  $h_2$  is  $10\epsilon'$ -close to  $p$ , then ok to output "tie" or either  $h_1, h_2$  as "winner"

else, if  $h_2$  is not  $10\epsilon'$ -close to  $p$  but is  $12\epsilon'$ -close, ok to "tie" or output  $h_1$  as "winner"

else  $h_2$  is  $12\epsilon'$  far from  $p$  +  $11\epsilon'$ -far from  $h_1$

so samples from  $p$  will fall in "difference" between  $h_1 + h_2$   
& will output  $h_1$



Since you know  $h_1 + h_2$ , you know  
where to look for this difference:

does  $p$  assign prob to  $A$  more like  $h_1$  or  $h_2$ ?  
(here you use samples)

Algorithm Choose: Input  $p, h_1, h_2$

$\swarrow$  samples  $\searrow$  explicit description

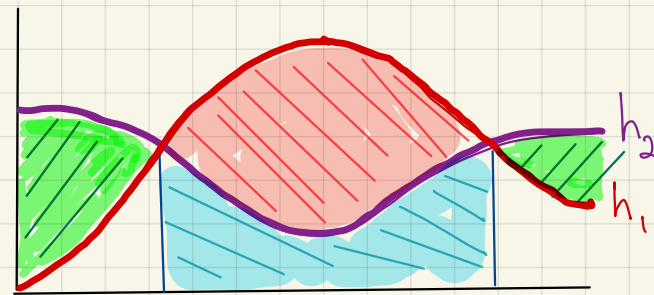
First some definitions:

$$A = \{x \mid h_1(x) > h_2(x)\}$$

$$a_1 = h_1(A) \quad \leftarrow \text{red + blue areas}$$

$$a_2 = h_2(A) \quad \leftarrow \text{blue area}$$

note  $\|h_1 - h_2\|_1 = 2(a_1 - a_2)$



$$\text{green area} = \text{red area} = a_1 - a_2$$

$$L_1 \text{ dist} = \text{green} + \text{red} = 2 \cdot \text{red}$$

will give factor of 2 in constants  $\rightarrow$

1. if  $a_1 - a_2 \leq 5\varepsilon'$  declare "tie" + return  $h_1$   
 $\underbrace{\quad}_{\frac{1}{2} L_1 \text{ distance}}$  (no samples needed)

2. draw  $m = 2 \log \frac{1}{\delta'} \frac{1}{(\varepsilon')^2}$  samples  $S_1, \dots, S_m$  from  $p$

3.  $\alpha \leftarrow \frac{1}{m} |\{i \mid S_i \in A\}|$   $\left. \begin{array}{l} \text{if } p = h_1, E[\alpha] = a_1 \\ \text{if } p = h_2, E[\alpha] = a_2 \end{array} \right\}$

4. if  $\alpha > a_1 - \frac{3}{2}\varepsilon'$  return  $h_1$   
 else if  $\alpha < a_2 + \frac{3}{2}\varepsilon'$  return  $h_2$   
 else declare "tie" + return  $h_1$

another  $\rightarrow$  additive error in constants

Why does it work?

- $h_1$  or  $h_2$  is  $\varepsilon'$ -close to  $A$  (given)
- If "tie" in step 1:

$h_1$  +  $h_2$  are  $10\varepsilon'$ -close (note  $L_1 \text{ dist} = 2(a_1 - a_2)$ )

$\Rightarrow$  both are  $\leq 11\varepsilon'$ -close to  $A$

So "tie" is ok

- Otherwise reach step 2:  $\|h_1 - h_2\|_1 > 10\varepsilon'$  ( $a_1 - a_2 > 5\varepsilon'$ )

Algorithm Choose:

$$A = \{x \mid h_1(x) > h_2(x)\}$$

$$a_1 = h_1(A)$$

$$a_2 = h_2(A)$$

$$\text{note } \|h_1 - h_2\|_1 = 2(a_1 - a_2)$$

1. if  $a_1 - a_2 \leq 5\varepsilon'$  declare "tie" + return  $h$   
(no samples needed)
2. draw  $m = 2 \frac{\log \frac{1}{\delta'}}{(\varepsilon')^2}$  samples  $S_1, \dots, S_m$  from  $p$
3.  $\alpha \leftarrow \frac{1}{m} |\{i \mid S_i \in A\}|$
4. if  $\alpha > a_1 - \frac{3}{2}\varepsilon'$  return  $h_1$   
else if  $\alpha < a_2 + \frac{3}{2}\varepsilon'$  return  $h_2$   
else declare "tie" + return  $h_1$

$\left\{ \begin{array}{l} \text{if } p = h_1, E[\alpha] = a_1 \\ \text{if } p = h_2, E[\alpha] = a_2 \end{array} \right.$



green area = red area =  $a_1 - a_2$   
 $L_1 \text{ dist} = \text{green} + \text{red}$   
 blue area =  $a_2$   
 blue + red area =  $a_1$

Why does it work?

- $h_1$  or  $h_2$  is  $\epsilon'$ -close to  $A$  (given)
- If "tie" in step 1, algorithm does right thing
- Otherwise reach step 2:  $\|h_1 - h_2\|_1 > 10\epsilon'$  ( $a_1 - a_2 > 5\epsilon'$ )

$$E[\alpha] = \Pr_{x \in p} [x \in A] = p(A)$$

assume (Chernoff) that with high prob  $|\alpha - E[\alpha]| \leq \frac{\epsilon'}{2}$

$h_1$  assigns  $a_1$  weight to  $A$   
 $h_2$  "  $a_2$  " "  $A$

if  $p$  is  $\epsilon'$ -close to  $h_1$ , assigns  $\geq a_1 - \epsilon'$  weight to  $A$

$$\alpha \geq a_1 - \epsilon' - \frac{\epsilon'}{2} = a_1 - \frac{3\epsilon'}{2} \quad \text{return } h_1 \text{ whp}$$

" " " " " "  $h_2$ , "  $\leq a_2 + \epsilon'$  weight to  $A$

$$\alpha \leq a_2 + \epsilon' + \frac{\epsilon'}{2} = a_2 + \frac{3\epsilon'}{2} \quad \text{return } h_2 \text{ whp}$$

Algorithm Choose:

$$A = \{x \mid h_1(x) > h_2(x)\}$$

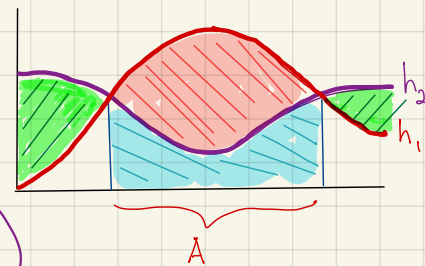
$$a_1 = h_1(A)$$

$$a_2 = h_2(A)$$

$$\text{note } \|h_1 - h_2\|_1 = 2(a_1 - a_2)$$

1. if  $a_1 - a_2 \leq 5\epsilon'$  declare "tie" + return  $h$  (no samples needed)
2. draw  $m = 2 \frac{\log \frac{1}{\delta'}}{\epsilon'^2}$  samples  $S_1, \dots, S_m$  from  $p$
3.  $\alpha \leftarrow \frac{1}{m} |\{i \mid S_i \in A\}|$
4. if  $\alpha > a_1 - \frac{3}{2}\epsilon'$  return  $h_1$   
 else if  $\alpha < a_2 + \frac{3}{2}\epsilon'$  return  $h_2$   
 else declare "tie" + return  $h_1$

if  $p = h_1$ ,  $E[\alpha] = a_1$   
 if  $p = h_2$ ,  $E[\alpha] = a_2$



green area = red area =  $a_1 - a_2$   
 $L_1$  dist = green + red  
 blue area =  $a_2$   
 blue + red area =  $a_1$

# The cover method - a method for learning distributions

def.  $\mathcal{C}$  is an " $\epsilon$ -cover" of  $\mathcal{D}$  if  $\forall p \in \mathcal{D}$   
 $\exists q \in \mathcal{C}$  st.  $\|p - q\|_1 \leq \epsilon$

$\uparrow$  smaller set of distributions

$\uparrow$  big set of distributions

Why useful?

hopefully  $\mathcal{C}$  is much smaller than  $\mathcal{D}$ , allows us to approximate  $\mathcal{D}$   
note  $\mathcal{C}$  not unique

Thm  $\exists$  algorithm, given  $p \in \mathcal{D}$ , which takes  $O(\frac{1}{\epsilon^2} \log |\mathcal{C}|)$  samples of  $p$  + outputs  $h \in \mathcal{C}$   
st.  $\|h - p\|_1 \leq 6\epsilon$  with prob  $\geq \frac{9}{10}$

big improvement:  $\Rightarrow$  union bnd over size of  $\mathcal{C}$  not  $\mathcal{D}$ !

Thm  $\exists$  algorithm, given  $p \in \mathcal{D}$ , which takes  
 $O\left(\frac{1}{\epsilon^2} \log |\mathcal{C}| \right)$  samples of  $p$  + outputs  $h \in \mathcal{C}$   
 s.t.  $\|h - p\|_1 \leq 6\epsilon$  with prob  $\geq \frac{9}{10}$

Pf.

Since  $p \in \mathcal{D}$ ,  $\exists q \in \mathcal{C}$  s.t.  $\|p - q\|_1 \leq \epsilon$   
 (could be more than one)

run "Choose" on  $p$  with every pair  $q_1, q_2 \in \mathcal{C}$   
 if best  $q_{\text{opt}}$  doesn't win all of its "matches" then it ties  
 with others that are not so bad

if  $q'$  is  $\geq 6\epsilon$ -far from  $p$ , then  $\geq \underbrace{6\epsilon - \epsilon}_{5\epsilon}$ -far from best  $q_{\text{opt}}$   
 $\Rightarrow$  loses to  $q_{\text{opt}}$

So all surviving  $q$  are  $\leq 5\epsilon$ -close to best  $q_{\text{opt}} \Rightarrow \leq 6\epsilon$ -close to  $p$ .  
 need all matches to give correct output — union bound on  $\binom{|\mathcal{C}|}{2}$  matches  $\square$

# Applications:

Example 1: learning distribution of a coin

domain =  $\{0, 1\}$

need to learn bias

Here  $\mathcal{D} = \mathbb{R}$

if use  $\mathcal{C} = \left\{0, \frac{1}{k}, \frac{2}{k}, \dots, \frac{k-1}{k}, 1\right\}$

then  $\forall$  bias  $p$ , let  $\frac{i}{k} \leq p \leq \frac{i+1}{k}$

then picking  $\tilde{p} = \frac{i}{k}$  gives  $\|p - \tilde{p}\|_1 \leq \frac{2}{k}$

biases of coin

So using  $k = \Theta\left(\frac{1}{\varepsilon}\right)$  gives  $\|p - \tilde{p}\|_1 \leq \varepsilon$

$|\mathcal{C}| = k+1$  # samples needed by cover method is  
 $= \Theta\left(\frac{1}{\varepsilon}\right)$   
 $O\left(\frac{1}{\varepsilon^2} \log \frac{1}{\varepsilon}\right)$



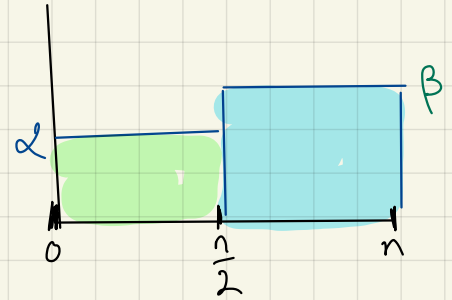
Example 2: 2-bucket distributions

now need to specify  $\alpha$  and  $\beta$

$$\text{so } \mathcal{C} = \left\{ \left( \frac{i}{k}, \frac{j}{k} \right) \mid i, j \in \{0, \dots, k\} \right\}$$

$$|\mathcal{C}| = \Theta\left(\left(\frac{1}{\varepsilon}\right)^2\right)$$

$$\# \text{ samples is } O\left(\frac{1}{\varepsilon^2} \log \frac{1}{\varepsilon}\right)$$



Example 3: monotone distributions

$$\text{Birge} \Rightarrow \mathcal{C} = \left\{ \left( \frac{i_1}{k}, \dots, \frac{i_{\lceil \log n / \varepsilon \rceil}}{k} \right) \mid i_1, i_2, \dots \in \{0, \dots, k\} \right\}$$

$$|\mathcal{C}| = \Theta\left(\frac{1}{\varepsilon^{\lceil \log n / \varepsilon \rceil}}\right) \Rightarrow \# \text{ samples is } O\left(\frac{1}{\varepsilon^3} \log n \log \frac{1}{\varepsilon}\right)$$