

Lecture 16:

Hypothesis Testing

Some Problems: (Given samples of p)

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

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

\sqrt{n}

is p ϵ -close to q
or ϵ -far from q

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

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

$n^{2/3}$

(Given samples of q) is p ϵ -close to q
or ϵ -far from q

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

is p monotone
or ϵ -far from monotone

\sqrt{n}

is p ϵ -close to monotone
or ϵ -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?

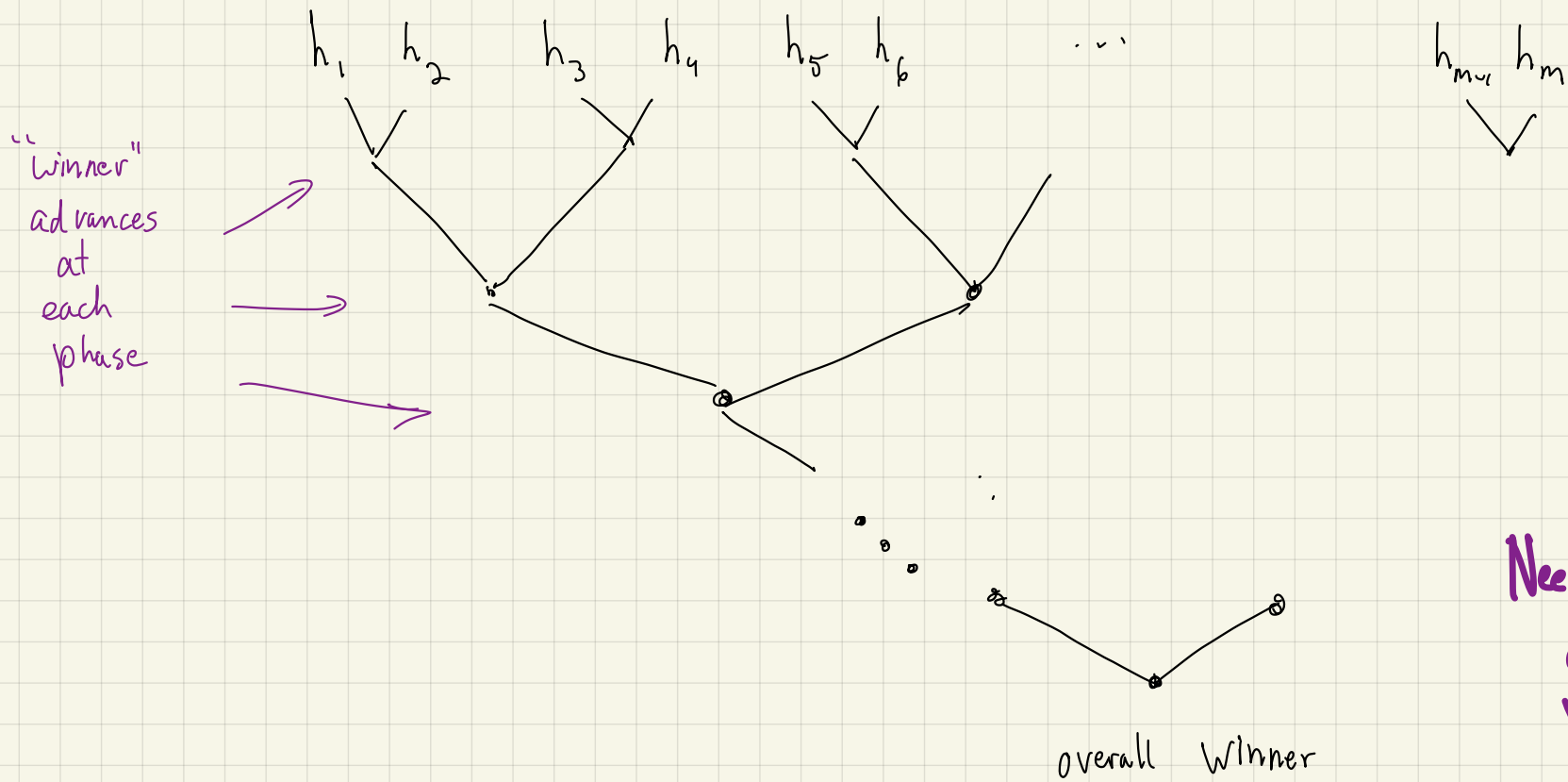
maybe either one is ok?

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"?



Need stronger guarantee!

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 ϵ' , confidence parameter δ'

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 ϵ' -close + one really far, will output ϵ' -close hypothesis }
if both ϵ' -close then output $12\epsilon'$ -close hypothesis } if at least one is close, will output pretty close hypothesis

e.g. \uparrow one is ϵ' -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 ϵ' -close to at least one of h_1, h_2
 ϵ', δ' 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
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

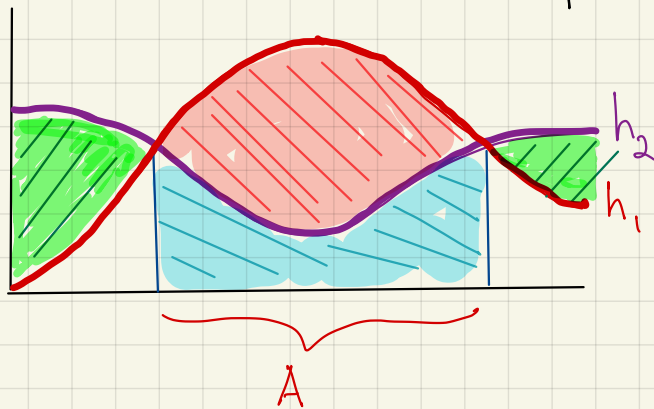
idea: wlog h_1 is ϵ' -close to p

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
 \nwarrow explicit description

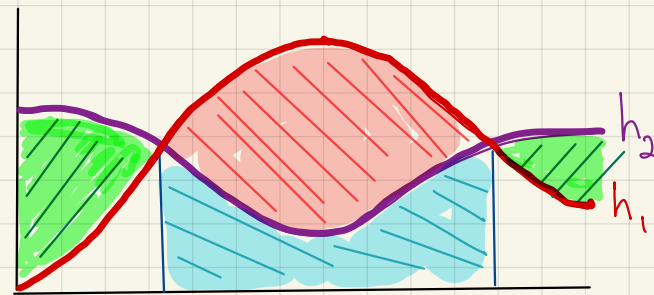
First some definitions:

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

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

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

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
 $\frac{1}{2} L_1$ 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 ε' -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 ϵ' -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 ϵ' -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:

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

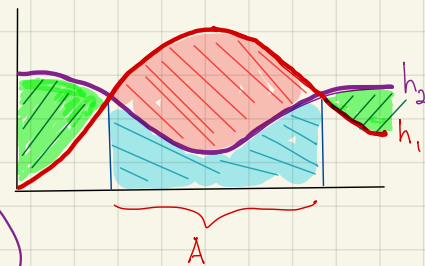
$$a_1 = h_1(A)$$

$$a_2 = h_2(A)$$

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

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 " ϵ -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_{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_{opt}
 \Rightarrow loses to q_{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 $\|p - \tilde{p}\|_1 \leq \varepsilon$

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

samples

needed

by cover method is

$$O\left(\frac{1}{\varepsilon^2} \log \frac{1}{\varepsilon}\right)$$

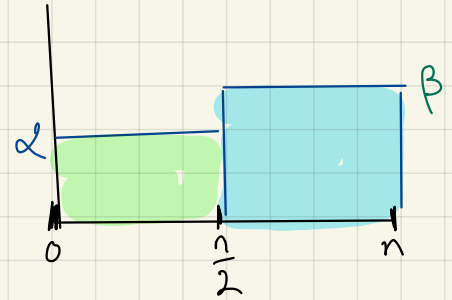
Example 2: 2-bucket distributions

now need to specify α and β

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

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

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



Example 3: monotone distributions

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

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