

Stat 202-2015S-W11 - Tuesday

## Review

Before the exam we studied Sampling Distributions. The scenario we studied was the following:

- We drew a sample of  $n$  individuals from a large population
- We measured a characteristic of each of the  $n$  individuals of the sample
- The characteristic could be height or weight (quantitative) or male/female } categorical  
· Prefer red over green  
· or prefer green over red }

For quantitative variables we sought to estimate the mean of the characteristic across the <sup>whole</sup> population,  $\mu$

For categorical variables we sought to estimate the proportion  $p$  of individuals with the characteristic across the whole population

In both cases we used only the sample to make the inference.

To estimate  $\mu$  we used the sample mean  $\bar{X}$   
 To estimate  $P$ , we used the sample proportion  $\hat{P}$

The sample mean  $\bar{X}$  and sample proportion  $\hat{P}$  were random variables. Every time we draw another sample we get different values for  $\bar{X}$  and  $\hat{P}$ .

What sample we draw is random  
 $\bar{X}$  and  $\hat{P}$  are numbers ~~that's why~~ ~~they~~ that depend upon the outcome of the random phenomenon, (sample)  
 That's why they are random variables,

Main results of chapter 5  
 was the distribution of  $\bar{X}$  and  $\hat{P}$ .

$$\begin{aligned}\mu_{\bar{X}} &= \mu \\ \sigma_{\bar{X}} &= \sigma / \sqrt{n}\end{aligned}$$

approx  $\left\{ \begin{array}{l} \mu_{\hat{P}} = p \\ \sigma_{\hat{P}} = \sqrt{\frac{p(1-p)}{n}} \end{array} \right\}$  } When the binomial distribution is a good approximation

Good approximation when population is at least 20x as large as sample

## Normal approximations

For large  $n$   $\bar{X} \xrightarrow{\text{approx}} N(\mu, \frac{\sigma^2}{n})$

IF  $np \geq 10$  and  $n(1-p) \geq 10$

then  $\hat{P} \xrightarrow{\text{approx}} N(P, \sqrt{\frac{P(1-P)}{n}})$

## New Chapter 6 Inference

For the rest of the semester  
we are doing Inference,

Purpose: draw conclusions from data

Formal inference emphasizes substantiating  
our conclusions via probability  
calculations,

Example: Are trees in a forest  
clustered or are they arranged  
randomly,

StatCrunch - longleaf

Are these trees clustered or  
random and how do we find out?  
531 trees old growth forest

Generate 584 random locations  
and see what it looks like;  
4 columns Seed 1

We already can see that most  
random forests look less clustered

But how do we be sure?

Key idea: Test Statistic

(Book cites another book which isn't  
in AC library and is ~~AGO~~  
I want to find it on interlibrary loan  
But I haven't done this yet)

So I am going to be vague

My name for test statistic

Clusterization Index.

Every forest has a clusterization  
index,

The higher the index the more  
clustered ~~all~~ the forest,

(PG5)

What forest has the highest  
clusterization index?

All trees in same spot.

What forest has the lowest  
clusterization index,

Trees tiled regularly like a tree  
farm:

Random forests have clusterization  
indexes somewhere in between

Because they are random each  
random forest has a different  
clusterization index.

In particular the distribution  
of the test statistic is important.

Where does our old growth forest  
fall on that distribution

If it is fall into the tails of the  
distribution we can conclude the  
forest is clustered more than random  
If not, we can't

## P-value

Who has heard of p-value?

The p-value is the probability that the test statistic would be ~~as extreme or~~ more extreme by random chance than what was observed in the data.

In other words the ~~p~~ p-value is the probability that the clustralization index would be greater in a random forest than what was observed in the old growth forest,

More extreme in this case is greater.

In some cases you may be asking if the clustralization index less than (more extreme would be less than) And in others more extreme would be greater than a threshold and less than a different threshold. For instance if you are asking if the forest random and you want to flag too clustered and not clustered enough, called a two sided alternative

(P97)

According to book the pvalue was 0.04

I don't know whether they used a one sided alternative or two.

$$0.04 < 0.05$$

at traditional cutoff for significance

You would say the test is significant at 0.05 level (the traditional level),

You would reject the null hypothesis

Null hypothesis being that the forest is random.

You would conclude that the forest is not random.

Pg 8

## Example #2

Researchers want to know if a new drug is more effective than placebo

20 patients receive new drug  
20 receive placebo

Twelve taking drug improve 60%  
8 taking placebo improve

State → proportion stats → two sample  
→ with summary

Sample 1

# successes 12  
# obs 20

Null hypothesis is

$H_0$  it is that both proportions are equal  $p_1 = p_2$

Sample 2

# successes 8  
# obs 20

Alternative hypothesis

do we care if its bigger than or both ≠

for two sided alternative  
p-value .20 large p-value

Pg 9

This means that if true proportions were the same,

In other words the probability of improving with the drug is the same as the probability of improving with the ~~the~~ placebo then the test statistic will be as extreme or more extreme 20% of the time

(I haven't told you what the test statistic is)

Because this is so high we cannot conclude that the drug has an effect.

That doesn't mean it doesn't have an effect. It could mean that it doesn't have an effect it could also mean the sample size is too small to see the effect because no effect is also not large enough.

PSG

# Review of terminology

test statistic

p-value

null hypothesis / alternative hypothesis

two sided alternative

versus one sided alternative

test statistic - a number you compute from data to be used to make an inference

probabilities and probability statement

null hypothesis - ~~stating~~ a statement that there is no effect - that the data observed ~~observed~~ are caused by random chance

alternative hypothesis - statement that there is an effect

p-value - the probability assuming the

null hypothesis is true that the test statistic will take on a value as extreme or more extreme as what is seen in data

one / two sided alternative - whether you consider ~~extreme~~ less than or greater than only one side or both (two sided)