STA201 Week 8: Statistical Inference, Statistics in the News, and Risk

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Robert Zimmerman (University of Toronto) STA201 Week 8: Inference/News/Risk

- Basic statistics (mean, standard deviation, quantiles, etc.)
- Histograms
- Normal distribution

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This Week

Statistical Inference

- Statistical Hypotheses
- NHST Framework
- Statistical Assumptions
- *p*-values
- Statistics in the News
 - The Replication Crisis
 - Poor Assumptions
- Financial Bisk
 Risk
 The 2008 Financial Crisis

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Why Statistics?

THIS IS WHY PEOPLE SHOULD LEARN STATISTICS:



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- We use statistics to extract information from our observations (raw data)
- We use mathematical techniques to ask what scientific claims we can infer from the raw data
- We can't "prove" or "disprove" what we can't fully observe (different from logical statements)
- We can only start with a hypothesis and then ask whether or not our observations support it

- Hypotheses must be testable they have to be expressible in statistical language
 e Usually, a mathematical statement
- Example: The average doily snauld amount in February in Tanto is 10 cm

· Non-example: It's snowing too much today (

Statistical Hypotheses II: H_0 and H_A

• We start with a *null hypothesis* and an *alternative hypothesis* which we label H_0 and H_A , respectively

"H-navaht

- H_0 represents the default scenario: nothing interesting is going on
- *H_A* represents the alternative situation: we suspect something interesting *is* going on
- H_A and H_0 can never both be true at the same time
- Often, H_A and H₀ are "opposites" either one is true or the other is true (but this isn't a requirement!)
- For example:

Ho: There tonds to be equal amounts of snowleds on Wadnewbays and Trasdays

Ha: There tonds to be more snowfell on Wednesdays than on Tuesdays

• Coin flip experiment:
$$M_0$$
: $p = \frac{1}{2}$ (ie, the coin is unbiased)
Let $P = \text{ probability of heads}$. M_a : $p = \frac{1}{2}$ (ie, the coin is biased)

• A clinical drug trial compares the effectiveness of a drug with that of a placebo to prevent heart attacks:

$$H_0$$
: this drug is no different from the placebo at preventing beat attacks H_A : this drug is better than the placebo at preventing beat attacks

PollEverywhere - Statistical Hypotheses

Which of the following is/are *not* a proper H_0 - H_A pair?

- H₀: The average Ontarian is more than 50% likely to vote for a Liberal candidate for Prime Minister
 H_A: The average Ontarian is at most 50% likely to vote for a Liberal candidate for Prime Minister
- H₀: If you buy Glop, you'll love its smooth texture H_A: If you buy Glop, you won't love its smooth texture
- H₀: There tend to be more red M&Ms in a pack than yellow M&Ms H_A: There tend to be equal amounts of red and yellow M&Ms in each pack

 \bullet H_0 : The average person in this room is at least 3 feet tall H_A : The average person in this room is at least 4 feet tall

NHST Framework I

The *Null Hypothesis Significance Testing (NHTS)* framework consists of the following steps:

1) We state H_0 and H_A , our statistical assumptions, and prespecify a significance level α between 0 and 1 (usually close to \mathcal{O})

"alpha"

2) We collect our data (observations)

- 3) We ask the key question: "If H₀ is true, what is the probability that we would observe data at least as extreme as this?"
- 4) We calculate this probability = probability at observing date at least as extreme, assuming the is true
- 5) We apply a rule: If the calculated probability is less than the significance level α , we reject H_0 i.e., the data we see is "too unlikely" to arise merely by chance (otherwise, we fail to reject H_0)

The order of these steps is *crucial* (why?)

NHST Framework II



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- In most studies, the observations are assumed to be independent (recall that two events are independent when the likelihood of one event occurring has no affect on the likelihood of the other event occurring)
- In terms of our data: none of the observations are affected by one another
- This is a critical assumption: without it, we would have to quantify *how* the observations depend on each other in order to perform statistical calculations

- Is it reasonable to assume the observations are independent?
- Sometimes: for example, estimating the bias of a (possibly biased) coin by flipping it – no flip affects the outcome of any other flip
- Sometimes not: for example, what is the probability of a random man on the street being color-blind? What is the probability of a color-blind man's brother being color-blind?
- Ensuring that an experiment is properly set up (in advance!) to satisfy this assumption is an important process called *experimental design*

Statistical Assumptions - Independence III



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Statistical Assumptions - Distributions I

- When we frame H_0 and H_A , we make the **statistical assumption** that the data follow some kind of underlying distribution with defining features (i.e., **parameters**) that we are trying to estimate
- For example: recall that the Normal distribution has two parameters: the mean μ and standard deviation $\sigma_{\underline{x_{signet}}}$
- A study of heights in adult males might conclude that the average (mean) height of the adult male population of Earth is 66 inches
- Assumption: Adult mole heights tend to follow a $N(\mu,\sigma)$ distribution

"Normel distribution with mean of and standard deviction or "

- Of course, we don't know for sure what distribution the data arises from there are *lots* more than just the Normal
- So how can we know which distribution to even *assume* the data arises from in the NHST framework?
- Sometimes it's straightforward, as in the case of coin flips (we have a distribution to describe exactly that situation)
- When we know very little about the data, it can be hard to even make a reasonable guess

Statistical Assumptions - Distributions III

• Fortunately, probability theory gives us very powerful tools to help

Definition

If X_1, X_2, \ldots, X_n represent *n* numerical observations, then we define the **sample mean** \bar{X}_n to be

$$\bar{X}_n = \frac{1}{n} (X_1 + X_2 + X_3 + \dots + X_n)$$

Eq: data:

$$X_{1}=9$$

 $X_{2}=10$ Then $\overline{X}_{4}=\frac{1}{4}(9+10+7+12)$
 $X_{3}=7$ $=\frac{1}{4}(38)$
 $X_{4}=12$ $= 9.5$
Eq: all use know about the data is that
 $X_{1}=-X_{2}$ and $X_{3}=-X_{4}$. Then $\overline{X}_{4}=0$. Why?
 $(u \ge \overline{X}_{4}=\frac{1}{4}(X_{4}+X_{2}+X_{3}+X_{4})$
 $=\frac{1}{4}(X_{4}+(-X_{1})+(-X_{2})+(-X_{3}))$
 $=\frac{1}{4}(0+0)$
 $= 0, \dots, 0$ as $x \ge -900$

Suppose $X_1 = 3$ and $X_3 = 3$ and the sample mean $\overline{X}_3 = 5$. What must X_2 be? χ_2 must be Q. Why?

$$\begin{array}{rcl} & & & & \\ & & & & \\ & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ &$$

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Suppose you know that each of X_1, \ldots, X_9 are negative numbers, and yet the sample mean \bar{X}_{10} is positive. What's the most you can say about X_{10} ?

- (\bullet) X_{10} is positive 44 %
 - X₁₀ is negative 23%
 - X_{10} is equal to $-(X_1 + \dots + X_9)$ 25%
 - Nothing 67.

$$\begin{array}{c} \frac{1}{10} \left(\chi_{1} + \cdots + \chi_{10} \right) = 0 \\ \chi_{1} + \cdots + \chi_{10} = 0 \\ \left[\chi_{1} + \cdots + \chi_{q} \right] + \chi_{10} = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{q} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} = - \left[\chi_{1} + \cdots + \chi_{1} \right] = 0 \\ \chi_{10} =$$

Suppose all observations X_i are equal to 5, except for $X_{43} = 543$ which is an **outlier**. What happens to the sample mean X_n as n gets larger?



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Statistical Assumptions - Distributions IV

- Two extremely important examples of these tools are the Law of Large Numbers and the Central Limit Theorem
- The Law of Large Numbers very roughly states that if the data are independent and arise from the same distribution (no matter which one) whose mean is μ, then the value of X
 _n will eventually approach the "true mean" μ as n grows higher and higher
- The Central Limit Theorem very roughly states that if the data are independent and arise from the same distribution (no matter which one) whose mean is μ and whose standard deviation is σ, then the distribution of the sample average *itself* eventually looks like a Normal(μ, σ/n) distribution n grows higher and higher

Mean & underlying data approaches O as n gots lager, so the average difference between In and u gets smaller

Statistical Assumptions - Distributions V



So how do these help?

- Suppose we want to compare the heights of the adult population of planet Stilt with those of adult Earthlings; we know that the average adult height on Earth is 165 cm
 - Stiltians appear to be quite tall...
- We set our significance level at $\alpha = 0.05$ and design our experiment, carefully stating our statistical hypotheses:
 - H_0 : There is no statistical difference between the average height of adults on the two planets
 - H_A : The average height of Stiltians is greater than the average height of Earthlings
- We collect 400 independent Stiltian height measurements and calculate the sample mean \bar{X}_{400} =180 cm

- To avoid technicalities: assume we know for a fact that $\sigma = 20$
- Under H_0 , the distribution of \bar{X}_n looks like a Normal distribution with mean 165 and standard deviation $\frac{20}{\sqrt{n}}$ for large *n* (in our case, n = 400 is definitely large enough): $\sum_{n=1}^{\infty} \frac{20}{2n} = \frac{20}{2n} = 1$



• Then under H_0 , $\bar{X}_{400} = 180$ cm is a random draw from the above distribution – that is, a Normal(165, 1) distribution

- Now we ask ourselves: what's the probability of drawing a number at least as high as 180 from the Normal(165, 1) distribution?
- Because the Normal distribution is so well understood, we can calculate this exactly: it's approximately 3.670966×10^{-51} ,

• Thus, we **reject** H_0 at the 5% significance level and conclude that Stiltians are, in fact, higher than Earthlings on average

NHST - Stilt Example IV



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- The probability we calculated on the previous slide is called a *p*-value
- Formally, the *p*-value is the probability that the underlying statistical model would generate data at least as extreme as our observations, given that the null hypothesis is true
- When our calculated *p*-value is below our pre-set confidence level, we conclude that the data is too unlikely to have occurred merely by chance under H_0
- Hence, we reject the null hypothesis:
- It is "very unlikely" that the sample of Stiltians whose height we measured just happened by chance to be atypically tall

- The study's conclusion might read "The Stiltish population is significantly taller than the Earthian population"
- **IMPORTANT:** We didn't **prove** that Stiltians actually *are* taller than Earthlings on average to do so would require measuring every Stiltian, computing the average, and comparing it to Earth's average
- If we could do that, we wouldn't need to use statistics!
- Instead, we concluded that it's (extremely) unlikely that Stiltians are not taller than Earthlings

- What significance level is low enough?
- Ronald Fisher, a pioneer of statistics and experimental design, suggested 5%
- Much of scientific literature/academia still uses this
- But it's completely arbitrary!
- "...statistical significance, an odd religion that researchers worship almost blindly"

Problems With *p*-values - Misinterpretation I

- *p*-values are *probabilities*, not *certainties*
- Suppose many published studies calculate p-values of 0.05
- Then on average, 1 out of 20 studies will incorrectly reject H_0
- How many studies are published every day? Kundudies

0.05 = 5% = 40

Problems With *p*-values - Misinterpretation II

Why would a Rats should be ashamed fly land on for falling in this trap something like this? **BEARS!!** This is ridiculous! p < .05

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Problems With *p*-values - Misinterpretation III

- *p*-values lead to publication bias
- The p < 0.05 is so entrenched that a study result with p = 0.06 is considered a "negative" study
- Journals with limited space want to publish new, interesting, "positive" findings – a study with p > 0.05 may contain important new findings, but is far less likely to be published

If we calculate
$$P=0.1$$
 (=10%), then there is a 10% choice that
our observations (or more extreme observations) arose due to chonce.
Not that high!

Correlation is not causation!

Problems With *p*-values - Misinterpretation IV



Problems With *p*-values - Misinterpretation V

- Extrapolating a population-based result to an individual can also be problematic
- For example: your well-conducted study compared the effectiveness of Drug A and Drug B to dangerously high serum rhubarb levels
- You find that Drug A was effective in 90% of patients who received it, while Drug B was effective in 30% of patients who received it
- You conclude that "Drug A works, but Drug B doesn't work"
- This is a disservice to the 30% of patients who responded to Drug B but might not respond to Drug A



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Problems With *p*-values - *p*-Hacking I

- Changing the prespecified threshold to declare results statistically significant
- Multiple comparisons
 - Multiple comparisons
- Increasing the size of the study population
 - *p*-value calculations are affected by the sample size
 - A very large medical study might produce a result that's statistically significant, but not *medically* significant
 - The time to achieve a normal temperature was 19.5 hours for Drug A and 19.8 hours for Drug B, a statistically significant difference
 - Drug A advertisement: "Expensive new Drug A reduced fever significantly faster than cheap old Drug B"

• Post-hoc analysis (testing analysis that were not prespecified)

- "If you torture the data long enough, it well tell you what you want to hear"
- Okay as a springboard to discover hypotheses for a new study
- Outright fraud
 - "Editing out" data points that sway the results away from the hoped-for conclusion
 - More difficult these days (still not impossible) now that many reputable journal require study authors to upload their raw data to a repository for scrutiny by other researchers

Suppose we have (properly) conducted an experiment which yielded a p-value of 0.002. Then H_0 must be false.

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• True 417.
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A hypothesis test is done in which H_A is that more than 10% of the population is left-handed. The *p*-value for the test is calculated to be 0.25. Which statement is correct?
We can conclude that more than 10% of the population is left-handed
We can conclude that more than 25% of the population is left-handed
We can conclude that exactly 25% of the population is left-handed
We cannot conclude that more than 10% of the population is left-handed

Statistical Significance



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Problems With *p*-values - What Can We Do

- It's easy to state why the framework is flawed but much harder to discover how to fix it!
- Suggestions:
 - Make the "standard" significance level (much) less than 0.05 (but many valuable results might be discarded/ignored if the *p*-value requirement is too stringent)
 - Report confidence intervals
 - Make studies more transparent transparent; i.e., *all* data must be submitted to the journal repository (but this won't stop data tampering before the data is submitted)
 - Combat publication bias all results, whether "negative" or "positive", must be uploaded to a central repository like Health Canada, FDA, APA, etc. (but this won't stop early-stage data tampering and might chill research efforts)

- An ongoing crisis, particularly in the field of social psychology
- Nobody questioned the results of many famous scientific studies until just recently, when scholars tried to replicate them without success
- According to a 2016 poll of 1,500 scientists reported in the journal Nature, 70% of them had failed to reproduce at least one other scientist's experiment. 50% had failed to reproduce one of *their own* experiments. (Wikipedia)
- Several well-respected researchers admitted to fabricating their results, and resigned in disgrace
- Optional (but interesting/disturbing) further reading

- Poll results are often stated in the form "correct to ±3%, 19 times out of 20"
- "19 times out of 20" is really a *p*-value of 0.05
- Even a perfectly conducted poll (free of cognitive biases!) will draw an atypical sample occasionally
- "19 times out of 20" implies that the pollster expects the sample to be atypical 1/20 times
- Framing effect!



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• Applying wrong assumptions to a statistical model can be disastrous: Sally Clark