## **Hypothesis Testing**

The scientific method in action

#### **Yordan Darakchiev**

Technical Trainer iordan93@gmail.com



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# Confidence Intervals Being confident is important

#### **Confidence Intervals**

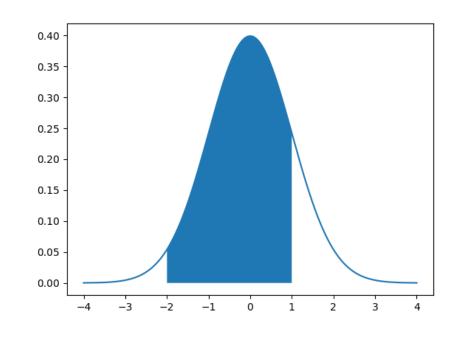
- In an experiment, we can't observe the variables' true values directly
  - We observe other values
  - We make assumptions as to how they are distributed
  - We can estimate the true value
    - Law of large numbers: when our sample is big enough, the sample parameters approach the population parameters
- With continuous values, it's useless to say that the mean is equal to a certain value (why?)
- Confidence interval a range of values that we're fairly sure contains the true value
  - How confident? A matter of choice
- Confidence level the probability that the value falls within the interval

## **Confidence Intervals - Interpretation**

- Similar to the probability interpretations
- To illustrate these, let's take a confidence interval [5; 7,3] and a 70% confidence level
- Frequency
  - If we perform the experiment many times, 70% of the values will fall in the interval [5; 7,3] and 30% outside it
- Certainty of next trial
  - Next time we perform the experiment, we are 70% certain that the value will fall within [5; 7,3]
  - Note that this is a statement about the interval, not about the value
- Typically used confidence intervals
  - **50%**; 90%; 95%; 99,7%

#### **Confidence Intervals and Z-Scores**

- Observe the Z-distribution (Gaussian,  $\mu = 0$ ,  $\sigma = 1$ )
- What's the probability that a value drawn from it  $x \in [-2; 1]$ ?
  - This corresponds to the shaded area in the graph
  - The cumulative function gives us the area to the left of some value
  - Shaded area = cdf(1) cdf(-2) = 0.819 = 81.9%
- Interpretations
  - If we draw many random numbers from the Z-distribution, we expect that 81,9% of them will be in [-2; 1]
  - If we draw one random number, there is 81,9% chance of it being in [-2; 1]
- Commonly used intervals
  - $1\sigma \rightarrow 68,27\%$ ;  $2\sigma \rightarrow 95,45\%$ ;  $3\sigma \rightarrow 99,73\%$
  - Also  $1,96\sigma \rightarrow 95\%$



## **Confidence Intervals Example**

- In the dataset heights.csv you're given the measured heights (in cm) of 351 elderly women (from an osteoporosis study)
  - Plot a histogram and / or boxplot to see what the distribution is
  - Print the mean  $\bar{x}$  and standard deviation s of the sample
  - Assume that the population follows a normal distribution
    - Real parameters unknown; our best guess:  $\mu = \bar{x}$ ,  $\sigma = s$
  - What are the confidence intervals of
    - **50%**, 90%, 95%
- To calculate the confidence intervals, we need to calculate the Z-scores
  - To do this, we'll use the percent point function, ppf
    - Inverse of the cdf
    - Returns the value at which the probability is less than or equal to the given probability
    - Example: Z-distribution
      - $ppf(0) = -\infty$ ;  $ppf(1) = \infty$ ; ppf(0.5) = 0; ppf(0.975) = 1.96

## **Confidence Intervals Example (2)**

- Note that once again we need to subtract the left white region
  - Area of shaded region: p (e.g. p = 0.95)
  - Area of both tails: 1 p
  - Percentage point of left tail:  $\frac{1-p}{2}$
  - Percentage point of right tail:  $\frac{1-p}{2} + p = \frac{1-p+2p}{2} = \frac{1+p}{2}$

```
import scipy.stats as st
def get_real_confidence_interval(probability, mean, std):
    lower_area = (1 - probability) / 2
    upper_area = (1 + probability) / 2
    return [
        st.norm.ppf(lower_area, mean, std),
        st.norm.ppf(upper_area, mean, std)]
```

95%

# **Testing Hypotheses**

The scientific method in action

## **Hypotheses**

- After performing an experiment and getting data, the scientific method requires that we form a hypothesis
  - Fact, law, theory and hypothesis are <u>different terms</u>
- In the simplest case, we have two hypotheses
  - Null hypothesis  $(H_0)$  the status quo is real, "nothing interesting happens"
  - Alternate hypothesis  $(H_1)$  what we're trying to demonstrate
- Types of hypotheses
  - Attributive something exists and can be measured
  - Associative there is a relationship between two behaviors
  - Causal differences in the amount / kind of one behavior cause differences in other behaviors

## **Hypotheses - Examples**

- Examples of hypotheses study of Disneyland visitors
  - Attributive
    - Most of the population has heard of Disneyland
    - Disneyland visitors are diverse in demographics
  - Associative
    - Income level is correlated with visiting Disneyland
    - People who live closer to Disneyland are more apt to visit Disneyland
  - Causal
    - Frequent exposure to Disneyland advertising results in increased attendance
    - Discounting tickets for local residents produces an increase in visitor numbers
- Note that attributive hypotheses involve one variable (univariate) while associative and causal hypotheses involve two variables (bivariate)

## **Testing a Hypothesis**

- In random experiments, we have error sources
  - Human error, systematic error, random errors, etc.
- We cannot prove (or reject) a hypothesis with complete certainty
- The errors we can make are two types
  - **Type I error** reject  $H_0$  while it's true (false positive)
  - **Type II error** accept  $H_0$  while  $H_1$  is true (false negative)
- The possible results can be summarized in the following truth table
  - Also called confusion matrix

#### **Action**

		Don't reject H <sub>0</sub>	Reject H <sub>o</sub>
Reality	H <sub>0</sub> true	<b>TN</b> true negative	<b>FP</b> (type l error) false positive
	H <sub>o</sub> false	<b>FN</b> (type II error) false negative	<b>TP</b> true positive

## Testing a Hypothesis (2)

- To measure the probability of producing a wrong hypothesis, we use a **test statistic** measure of deviations from  $H_0$ 
  - Different tests produce different measures (statistics)
  - We accept or reject the null hypothesis based on the value of the test statistic
- Let's denote the probability of getting a type I error with  $\alpha$ 
  - Each value of the selected test statistic has a corresponding alpha-value
  - We perform the experiment, get data and calculate the test statistic value
  - From that, we calculate the corresponding alpha-value
  - We reject the null hypothesis if  $\alpha < \alpha_c$ , where  $\alpha_c$  is a **critical confidence level**

#### **Z-test**

- A Z-test uses the Z-statistic
- $H_0$ : standard normal distribution
- Example: light bulb factory
  - A factory produces light bulbs with lifetime  $X \sim N(\mu = 500h, \ \sigma = 50h)$
  - A sample of 25 bulbs has a mean lifetime  $\bar{x} = 480h$
  - Is there something wrong with the production line?
- Forming hypotheses
  - $H_0$ : The production line works normally, the observed deviation of the sample mean from the population mean is due to chance
  - $H_1$ : The production line is broken

## Z-test (2)

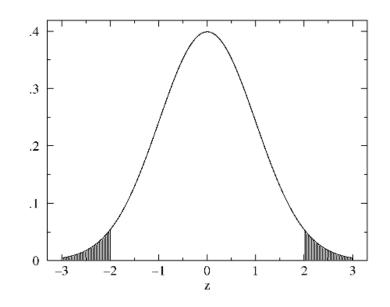
- Suppose we take a lot of samples from the entire population
  - Each sample mean will be different
  - The distribution of sample means will be more or less Gaussian
    - Parameters (our best estimate):  $\mu_{\bar{\chi}} = \mu$ ,  $\sigma_{\bar{\chi}} = \sigma/\sqrt{n}$
    - Here's why the parameters are chosen as such
- If  $H_0$  is correct, we assume that  $\bar{x} \sim N(\mu, \sigma/\sqrt{n})$
- Z-statistic

$$Z = \frac{\bar{x} - \mu}{\sigma_{\overline{x}}} = \frac{480 - 500}{50/\sqrt{25}} = -2$$

- We can see that we are 2 std's below the mean
- How extreme is that?
  - What's the probability that we get results as extreme or more extreme than we observed, assuming the null hypothesis is true?
    - Less than 5%

#### **Two-tailed Z-test**

- We can get the confidence interval from the Z-statistic
- We are looking for more extreme values
  - Values outside the confidence interval
  - What's the probability  $P(|Z| \ge 2)$ ?
  - We're looking for a value different than the mean
    - We can't assume whether it's smaller or larger
    - Therefore, we have to look at both "tails" of the distribution



- If we assume a critical value (also called a p-value) of 5%,
   the results are significant
  - P(|Z| > 2) ≈ 0.0455 = 4.55%
- We can reject H<sub>0</sub> at the 5% level
  - Even at lower levels, up to 4,55%

#### **One-tailed Z-test**

- The same logic applies, but now we're looking at one tail only
- Question: Is the lifespan **significantly lower** than it should be? Cutoff point:  $\alpha_c = 5\%$ , Z = -2
  - $P(Z \le -2) = \frac{0,00455}{2} 0,02275 = 2,275\% < \alpha_c$
  - Answer: Yes, at the given significance level
- Question: Is the lifespan significantly higher than it should be?
  - $P(Z \ge -2) = 97,725\% \gg \alpha_c$
  - Answer: No, at the given significance level

#### t-test

- The Z-test requires that we know the standard deviation of the population
  - Usually not available
- We can use another test statistic, called t
- Advantages over the Z-test
  - lacktriangle We don't need to know the population  $\sigma$
  - It's better when we have very small sample sizes (e.g., n < 30)
  - It can be used for testing the mean of a sample against a standard, but also for comparing two means
    - We can see whether two sets of data are significantly different from each other
- Null hypothesis: The test statistic follows Student's t-distribution
  - Similar to Gaussian distribution, with "fatter" tails

### One-Sample t-test

- The details of the calculation are fairly complex but we can do this in code
  - Using scipy.stats
- First, we generate 100 random numbers with  $\mu = 5$ ,  $\sigma = 10$
- Then we ask whether the sample mean is equal to the true mean (and other values, just for testing)
- We get the p-value probability of the null hypothesis being true
  - I.e. probability that the mean is equal to the given mean

```
sample_data = st.norm.rvs(5, 10, 100)

print(st.ttest_1samp(sample_data, 5).pvalue) # 0.9301
print(st.ttest_1samp(sample_data, 4).pvalue) # 0.3352
print(st.ttest_1samp(sample_data, 0).pvalue) # 1.104e-6
```

## **Independent Two-Sample t-test**

- We compare two independent distributions
  - We want to see whether they have the same mean
  - We assume equal variances (scipy can also do tests with unequal variances – important when sample sizes differ)
- Example: Grain size
  - We are given data (in grain\_data.csv) of grain sizes from two different farms
  - Do they differ significantly (at the 95% level)?
  - \* We can also plot histograms to see what the distributions look like

```
grain_data = ...
st.ttest_ind(grain_data.GreatNorthern, grain_data.BigFour)
# Ttest_indResult(statistic=1.312336706487564,
# pvalue=0.20792200785311768)
```

## Paired Two-Sample t-test

- We compare two distributions
  - Observations in samples can be paired
  - Examples before / after observations; comparison between two different treatments applied to the same subjects
- Example: Drinking water
  - We are given data (in water\_data.csv) of Zn concentration in surface and bottom water at 10 different locations
  - Does the true average concentration in bottom water exceed that of top water?
  - We use a paired t-test because the samples are from the same locations
  - It reduces experimental error (and provides stronger evidence)

```
water_data = ...
# We use a one-tailed t-test
st.ttest_rel(water_data.surface, water_data.bottom).pvalue / 2
# 0.00044555772891127738
```

#### **Generalizations to More Variables**

- Sometimes it's not enough to compare two distributions
  - We may want to compare multiple distributions against the same null hypothesis
  - E.g. how is the percentage of smokers distributed by income and age?
- Other times, we create a model and want to evaluate it
  - E.g. a linear regression
  - We can explain some of the variance in the sample
- There are other tests to perform these "checks"
  - ANOVA (Analysis of Variance) useful for grouped data
    - Observe the variance inside groups and between groups
  - Chi-square(d) test can be applied to categorical data
    - Two common types
      - How good a model is (goodness of fit)
      - Whether two variables are independent

## **Analysis of Variance (ANOVA)**

- We want to compare several groups
- $H_0$ : The means of the groups are the same
- Method (scipy.stats.f oneway())
  - For each group ⇒ group mean
    - In-group variance: distances from an individual point to the group mean
    - Between-group variance: distances between the means of two groups
  - For the entire data ⇒ total mean (mean of all data)
    - Also equal to the mean of all group means
    - Total variance: in-group + between-group
- F-statistic (Fisher)
  - $F = \frac{\text{variance between groups}}{\text{variance within groups}}$ 
    - F large  $\Rightarrow$  the variance between groups dominates
    - For each value of F, there's a corresponding p-value
      - If  $p \le p_c$ , we can reject  $H_0$

## Chi-Squared ( $\chi^2$ ) Test

- Compares expected (predicted) and observed frequencies
  - Is there a significant difference between these?
  - This is a goodness-of-fit measure
    - How well were we able to predict
- Statistic:  $\chi^2 = \frac{(f_{\text{observed}} f_{\text{estimated}})^2}{f_{\text{estimated}}}$
- $H_0$ : No significant difference between observed and estimated
  - The test returns the value of the statistic and the p-value corresponding to it
  - Works the same as any other test
  - Python: scipy.stats.chisquare()

# Common Misconceptions

Because everyone can be wrong

## Some p-value Misconceptions

- Goodman, S. (2011), <u>source</u>
- "If p = 0.05,  $H_0$  has 5% chance of being true"
  - The data alone can't tell us how likely we are to be wrong
  - p is calculated under  $H_0$ , so it can't be the probability of  $H_0$  being false
- "p = 0.05 means that if we reject  $H_0$ , the probability of type I error (false positive) is only 5%"
  - I.e. seeing a difference where there isn't any
  - $\blacksquare$  => 5% chance of false rejection = 5% chance  $H_0$  is true
    - Wrong, see first bullet
- "If p = 0.05, we have observed data that will occur **only** 5% of the time assuming  $H_0$ "
  - The p-value is the probability of observing data as extreme or more extreme under  $H_0$

## Some p-value Misconceptions (2)

- "A nonsignificant difference means the groups are the same"
  - It only means we don't have enough data to reject  $H_0$
- "A scientific conclusion or treatment policy must be based on whether or not the *p*-value is significant"
  - The results have to be checked against prior data
- Failing to reject  $H_0$  means that  $H_1$  is true
  - It means that we don't have enough evidence to reject it
  - We can't accept (or reject) any other hypothesis
  - "Absence of evidence is not evidence of absence"
- https://xkcd.com/882/
- https://www.xkcd.com/1478/
- "Still. Not. Significant" article

## Summary

- Confidence intervals
  - Confidence level
- Hypothesis tests
  - Z-test
  - t-test (one-sample, two-sample)
- Hypothesis tests of many variables
  - ANOVA
  - Chi-squared
- p-value misconceptions

# Questions?