



Research method - 10004

Mind, Brain And Behaviour 2 (University of Melbourne)

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1. Inference

- A. Introduction to quantitative psychological research
- B. Inference as the goal of psychological research
- C. Populations and samples
- D. Examples
- E. Summary

A . intro to quantitative psychological research

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Quantitative psychological research addresses a broad range of topics. Even though there are many types of questions asked, they tend to fall into one of three categories:

1. Difference - is one group of people different to another in some way?

MBB2 online modules focus on this.

2. Association - is one construct related to another? A MBB2 tutorial class will address this

3. Prediction - does one construct influence another? You will learn about this in future psychology subjects

B. The Goal of Psychological Research

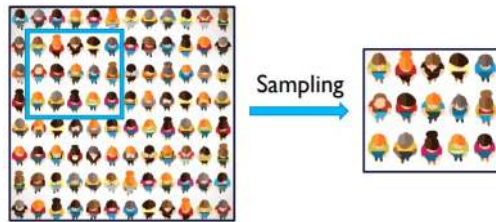
- When we conduct a psychological research project, our aim is to **make inferences** (in other words, suggestions or claims) about a **population**. Put simply, we want to say something about a **population**
- A population is **everyone of interest to a research question**. In other words, it is the research question that defines the population.

c. samples & populations

Taking Samples from Populations

It is usually not possible to recruit all people in a population to participate in a study.

Instead use a **sample**: a group of people taken from the population to participate in a study.



Making Inferences Based on Samples

We can then make **inferences** about the population based on what happens with **measurement** of our sample.

We aim to infer that what is typical for our sample should also be typical for the population.

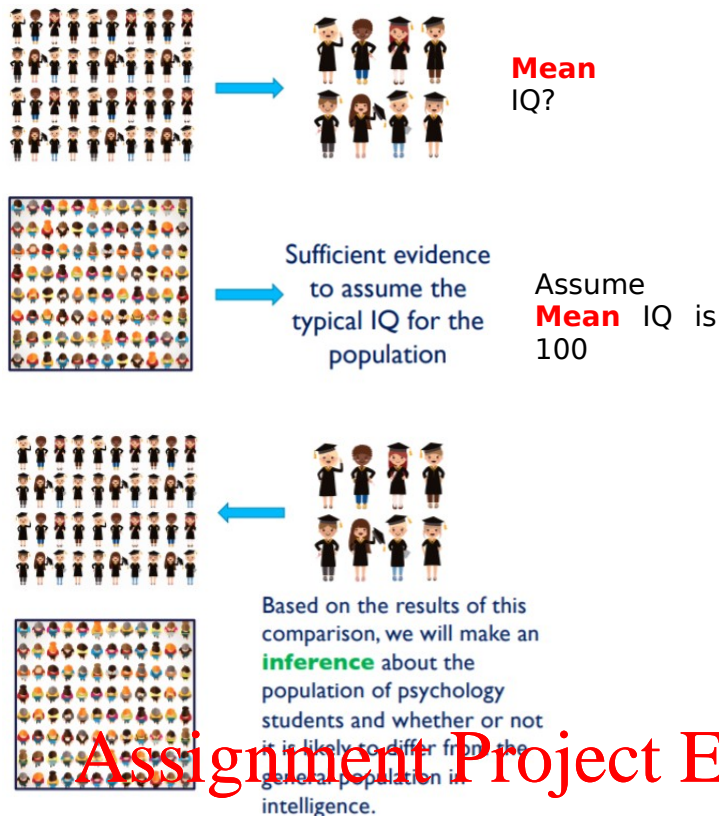


D. Example

Example 1: Are Psychology Students Smarter?

- **Research Question:**
Are psychology students smarter than the general population?
- Answering this question involves knowing the **typical IQ for**:
 1. Psychology students
 2. The general population.

Once we know **what is typical** for each of these groups, then we can compare them and assess the evidence for a difference.



In quantitative terms, **mean scores** will be our indicator of what is typical.

We know the population mean in this example. We could compare our sample mean to this value to assess evidence for a **difference**.

Based on the results of this comparison, we will make an **inference** about the population of psychology students and whether or not it is likely to differ from the general population in intelligence.

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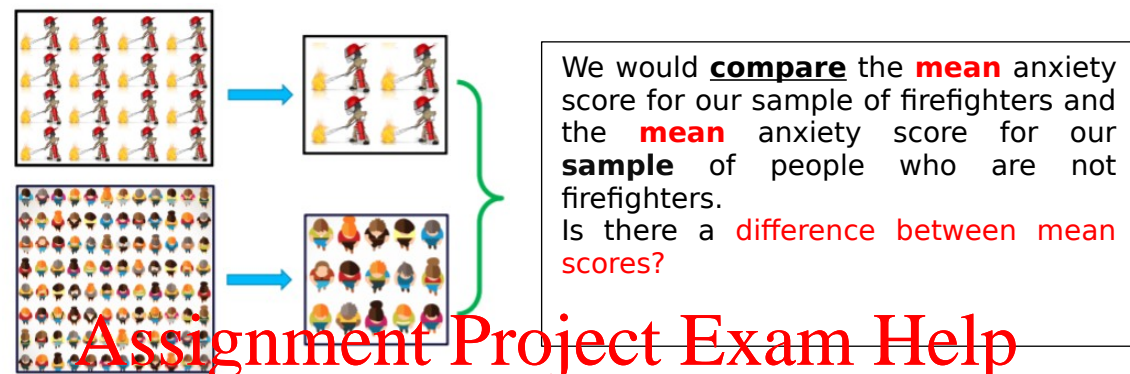
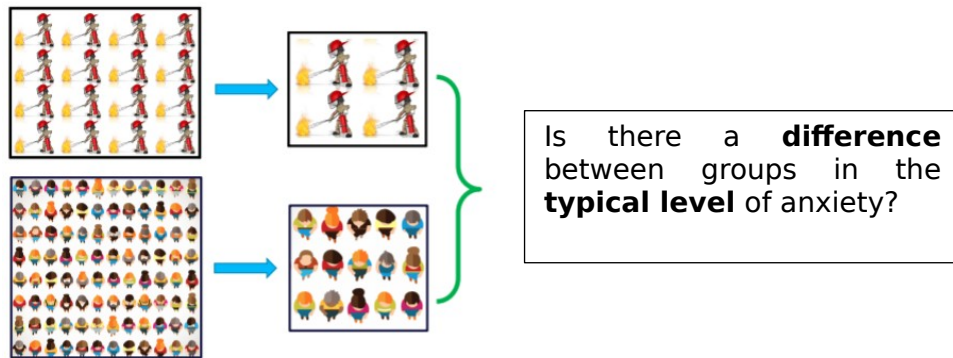
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EXAMPLE 2: Anxiety in Firefighters

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- Research Question:
Do Firefighters differ from the general population in their experience of anxiety?
- Answering this question involves knowing the **typical level of anxiety** for:
 1. Firefighters
 2. People who are not Firefighters

Once we know what is **typical** for each of these groups, then we can **compare** them and **assess the evidence for a difference**.



E, Summary

- Quantitative psychological research aims to generate knowledge about **populations**.
- A population is **everyone** of interest to a research question.
- Because we usually can't measure a population in its entirety, **a sample is drawn from the population and measured**.
- We can **make inferences about the population based on the evidence observed in the sample**.

- In these MBB2 modules, we will focus on inference about **differences in means scores.**

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2, Typical and Extreme Scores

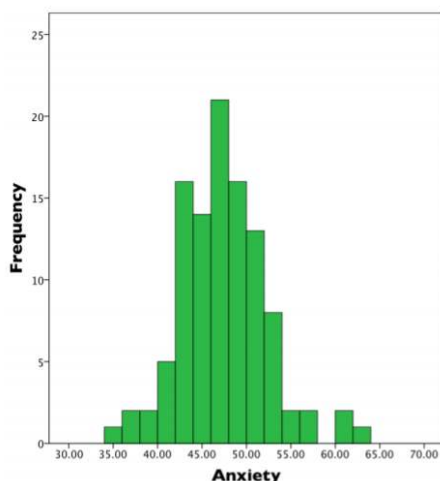
- A. Recap on distributions of individuals' data
- B. The Normal Distribution
- C. Typical and extreme scores
- D. The 2s rule of thumb
- E. Summary

A. Recap on distributions of individuals' data

Distribution of Data:

- To understand a psychological construct, we need to know how it is **distributed across a population**.
- When measured, constructs takes on **different values** for **different people** in a sample.
- Collectively, those different values form a **distribution** of data, which can be described in terms of **central tendency (mean)** and **variability (the standard deviation)**.
 - ➔ SD = A common measure of average spread (or variability) around the centre score (i.e., the mean)

Typically we estimate σ using s , the standard deviation of a **sample**

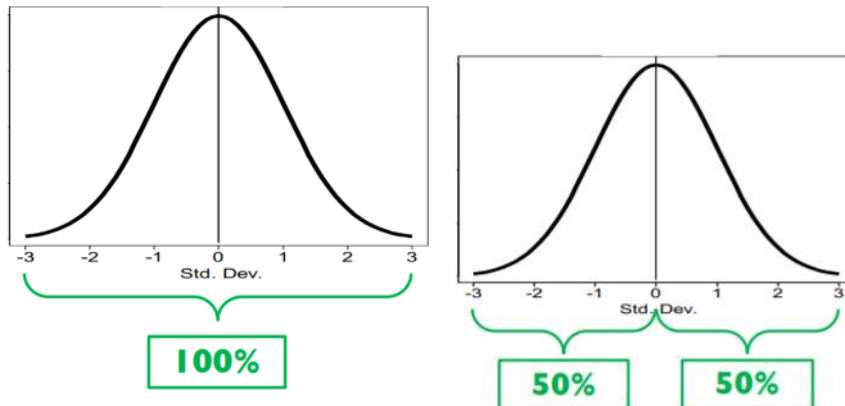


A **histogram** is one way to represent a **distribution**

Distributions of data can be described according to their **central tendency (m)** and **variability (s)**
 $m = 46.87$ $s = 4.84$

- ➔ a **normal** shape.
- **Most of the people are in the middle.** The peak of the graph.
- **Relatively fewer people are on the outsides.** The tails of the graph.

B. Normal distribution



Majority of observations are in the **middle**.

Observations **reduce in frequency towards the tails**.

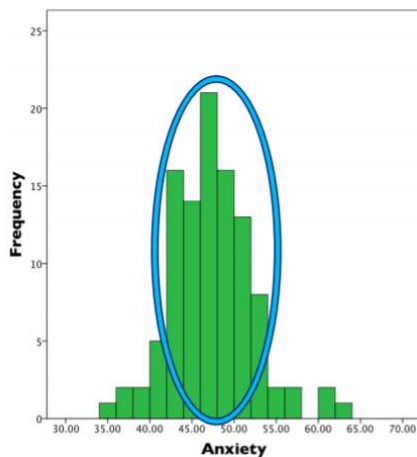
The distribution is **symmetrical**.

C. Typical and Extreme Scores

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Typical scores

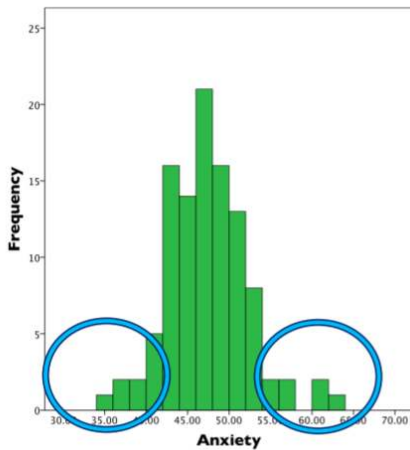
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In this distribution, it is **expected** or **typical** to find scores around 40 - 55 in this distribution. Why? These scores **occur frequently**

Typical scores are **expected**, occur with **high frequency/probability**

Extreme scores



it is relatively **unusual** to find extreme scores – ones that are **very low or very high**. Why? They occur **infrequently** in this distribution.

They indicate a **difference** to typical scores in terms of whatever is being measured.

Typical VS Extreme

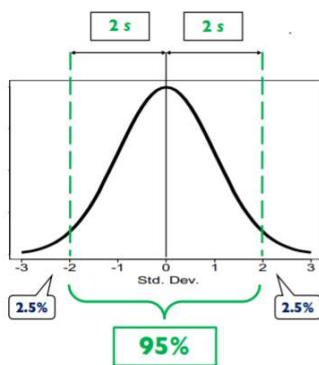
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How can we more reliably decide what counts as **typical vs extreme**?

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D. The 2s Rule of Thumb

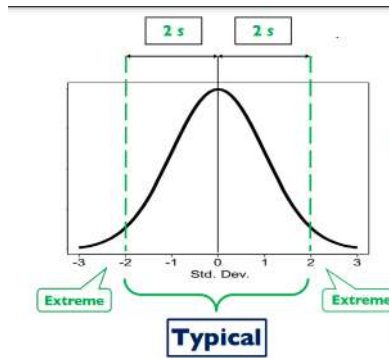
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In a distribution with a normal shape, **95%** of scores fall **within** approximately **2 standard deviations (s)** of the mean.

those scores **outside 2 standard deviations (s)** of the mean as being **typical**. They are **expected** as they occur **frequently** in this distribution

those scores outside 2 standard deviations (s) of the mean as being **extreme**. They are not expected as they occur **infrequently** in this distribution



Applying the 2s Rule of Thumb

We can apply our **2s rule of thumb** to decide what might be **typical** and **extreme** in this real distribution of data.

$$m = 46.87, s = 4.84$$

$$s = 4.84 \times 2 = 9.68$$

Lower limit

$$46.87 - 9.68 = 37.19$$

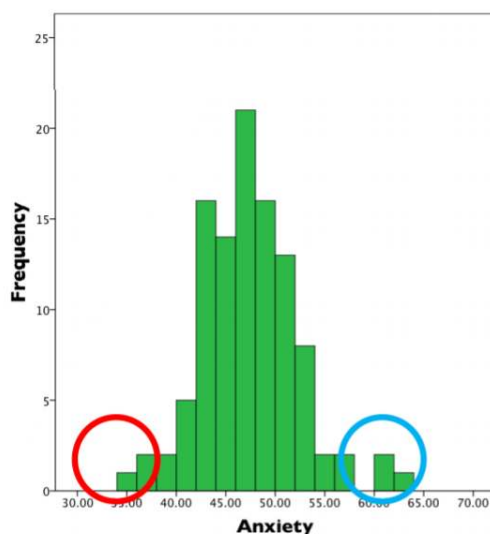
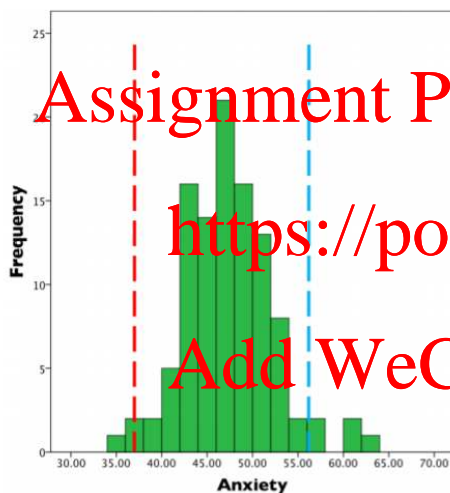
Upper limit

$$46.87 + 9.68 = 56.55$$

→ More extreme than lower limit: **2.02%**(m-lower l)

More extreme than upper limit: **4.04%**

→ Within 2s of distribution mean: **93.94%**



The extreme cases are noteworthy. They are unusual in terms of reported level of anxiety and indicate a difference from what is typical.

E. Summary

- A distribution of data can be fundamentally described according to its **central tendency (m)** and **variability (s)**.
- In a normal distribution, most observations are close to **m**, and they reduce in frequency towards the distribution's tails.
- We can use the **2s Rule of Thumb** to define a **critical limit**, beyond which, cases are considered **extreme**.
- We are most interested in extreme cases. These cases are unusual and indicate a difference in terms of what is being measured

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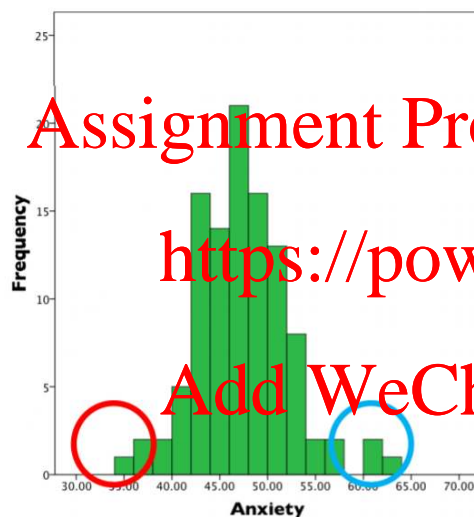
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3, Distribution of Sample Means

- A. Recap: Distributions of Individuals' Data
- B. The Distribution of Sample Means
- C. Central Limit Theorem
- D. Summary

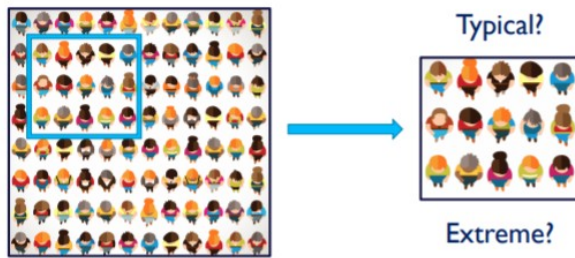
A. Recap: Distributions of Individuals' Data



We previously determined if **an** individual's score was typical or if it was extreme according to the characteristics of a **distribution of OTHER individuals' scores**.

B. What about an Entire Sample - the distribution of Sample Means

What if we wanted to know if not just an individual, but a **sample**, was typical or if it was extreme?



One population, many sample

- When we conduct research, we usually **recruit one sample** from each population of interest.
- However, there are **many samples** that could possibly be recruited from any population.
- How can we tell if our one sample is typical of a certain population, or if it is extreme and indicates a difference?

→ We previously examined **individual scores** in a **distribution of other individual's scores**. We can do much the same thing to determine if a sample is typical or if it is extreme.

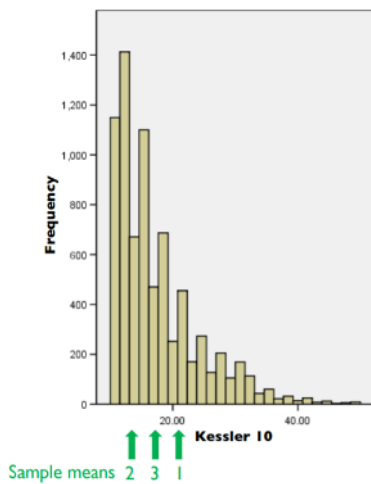
→ ① we would need a score for our one sample. This would be **the sample mean**. ② we would need a **distribution** made up of **sample means**, within which, we could examine our **one sample mean**.

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THE DISTRIBUTION OF SAMPLE MEANS

Collection of all the possible random sample of a particular size (n) that can be obtained from a population

Examples:



Suppose this is a **population**

Let's take some samples of 5 people at random.

Scores – sample 1

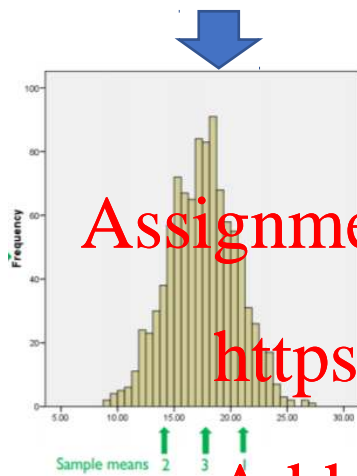
13, 16, 18, 28, 29 Mean = 20.8

Scores – sample 2

12, 13, 13, 14, 18 Mean = 14.0

Scores – sample 3

15, 15, 15, 20, 21 Mean = 17.2

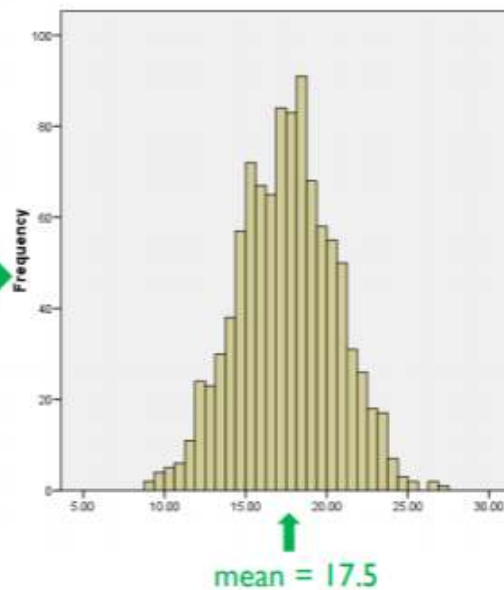
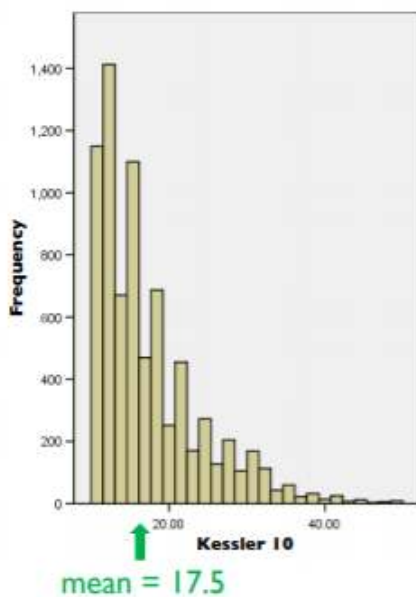


A **Distribution of Sample Means** with 1000 random samples, each with $n = 5$.

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C. Central Limit Theorem

- In real research, it is not pragmatically possible to collect all possible random samples from a population. However, we don't need to.
- The **Distribution of Sample Means** is a **theoretical distribution** governed by a **mathematical theorem** the **Central Limit Theorem**.
- **The Central Limit Theorem** tells us the precise characteristics of the distribution of any distribution of sample means.

Central Limit Theorem Tells Us

- The precise characteristics of a distribution of sample means for samples of any size (n).
- The distribution of sample means is the **SAME as the population mean**.
- For **large sample sizes (≥ 30)**, the distribution of sample means will have a **NORMAL** shape, regardless of shape, mean, SD of population
- so we can work out what is 2SDs from the mean
- the standard deviation of the sampling distribution means is called the **standard error** of the mean

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formula of standard error

The standard error formula: $\sigma_M = \frac{\sigma}{\sqrt{n}}$

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As sample size \uparrow , standard error \downarrow .

In turn, estimation of the population mean becomes **more precise**. When a sample is large enough, its mean provides a reliable estimate of the population mean.

What does this all mean?

- We know if our sample is large enough, the distribution of sample means will be **normal**.
- We know the mean of the distribution of sample means, and we can calculate its **standard error**.
- Therefore, we can use our **2s rule of thumb** to test if our sample mean is typical or if it is extreme

D. Summary

- We can decide if individuals' scores are typical or extreme by **comparing**

- one score with a distribution of other individuals' scores.
- We can apply a similar process to determine if a sample is typical or extreme, with the use of our **sample mean** and a **distribution of sample means**.
 - **Central Limit Theorem** tells us:
 - 1) **the distribution of sample means = population mean**;
 - 2) the details of standard error; how the standard error is related to the population standard deviation
 - 3) as sample size increases, standard error decreases.infer the population mean from the sample mean

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4: Null Hypothesis Significance

Testing

- A) Hypotheses
- B) The Null Hypothesis
- C) Null Hypothesis Significance Testing
- D) Determining the probability of a sample mean
- E) Summary

A) Hypotheses

- A hypothesis is a predictive statement. When we conduct psychological research, we pose and test an **experimental hypothesis**.
- An experimental hypothesis is a statement that **predicts an EFFECT**. This effect might be one of: DIFFERENCE / ASSOCIATION
- **Experimental** hypotheses = **alternative** hypotheses. But, alternative to what?

B) Null Hypotheses

- A **null hypothesis** is also a prediction. The null always predicts the one basic concept regardless of what is being investigated – that **nothing is happening**.
- States that in the general population there is no change, no difference, or no relationship
- In other words, the null hypothesis is a hypothesis of **NO effect**
➔ NO DIFFERENCE / NO ASSOCIATION
- The **experimental** hypothesis is the **alternative** to the **null** hypothesis. Only one of these hypotheses can be supported by the research data

Statistical Notation:

Null Hypotheses: H_0

Alternative (Experimental): H_1

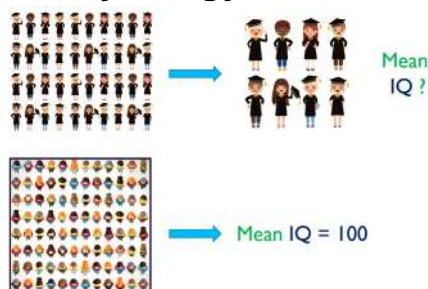
C) Null Hypothesis Significance Testing

1. Propose a null hypothesis that a population parameter (mean) has a particular value.
2. Proceed assuming the null hypothesis is **true**.
3. Determine the **probability** of the sample mean occurring if the null hypothesis is true.
4. If the probability of the sample mean occurring is **small**, **reject** the null hypothesis.
If the probability is **large**, do **not reject** the null hypothesis.

D) Example: Null Hypothesis Significance Testing

Testing

Are Psychology Students Smarter?



Step 1:

Pose a null hypothesis that a **population parameter** has a certain value.

$H_0 = 100$

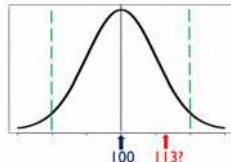
Step 2:

Proceed assuming $H_0 = 100$ is **true**. Collect the data. Observe the sample mean.

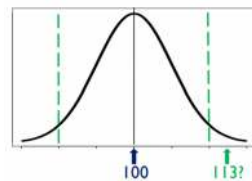
$H_0 = 100$, Observed $m = 113$

Step 3:

- Determine the **probability** of the sample mean occurring if the null hypothesis is true. Given $H_0 = 100$, is our sample mean of 113 **typical**, or is it **extreme**?
- Involves a **statistical test** based on a distribution of sample means with a mean of 100 (the same as the null hypothesized population mean). We know the shape will be **normal**. We can also calculate **standard error**.
 - If our sample mean is **typical**, then it **does not** provide evidence for a difference.



- If our sample mean is **extreme**, then it **does** provide evidence for a difference.



Step 4: Either reject or do not reject the null hypothesis.

Extreme sample mean? Evidence for a **difference**. **Reject** null hypothesis.

- Probability is **low**

Typical sample mean? **No** evidence for a difference. **Do not reject** null hypothesis.

- Probability is **HIGH**

if the null hypothesis is **true** we expect our sample mean to be **within 2 SDs** of the null hypothesis mean

We can calculate the probability using **z-scores**

E) Summary

When we conduct research, we pose an **experimental hypothesis** – one of **effect**.

The opposite of this is a **null hypothesis** – one of **no effect**.

Null Hypothesis Significance Testing involves

- 1) **assigning a value** to the null hypothesis.
- 2) then conduct **a statistical test** to determine the **probability of our sample mean occurring if the null hypothesis is true**.
- 3) If the probability is **low**, we **reject the null** hypothesis
if the probability is **high**, we do **not reject** it

5 The Single Sample z-Test

A. Null Hypothesis Significance Testing Recap

1. Propose a null hypothesis that a population parameter (mean) has a particular value.
2. Proceed assuming the null hypothesis is **true**.
3. Determine the **probability** of the sample mean occurring if the null hypothesis is true.
 - If the probability of the sample mean occurring is **small**, **reject** the null hypothesis. If the probability is **large**, do **not reject** the null hypothesis.

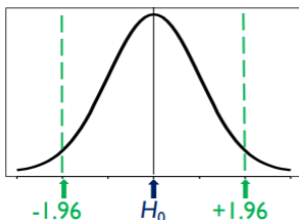
Probability: What Counts as Small?

5%: Alpha Level / Level of Significance

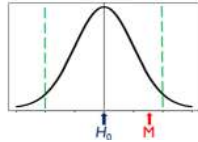
B. 5% Alpha level / Level of significance

The **alpha level** defines which sample means in a distribution of sample means are expected or typical, and which are unlikely or extreme, if the null hypothesis is true

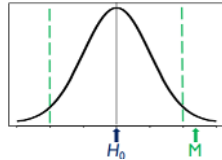
When the comparison distribution is **perfectly normal**, the critical limits set by the 5% Alpha Level are **precisely ± 1.96 standard errors** from the mean of the distribution.



- If our sample mean is **inside these limits**, the **probability is greater than 5%**, and therefore, **high**. Do **not reject** the null hypothesis.



→ If our sample mean is outside these limits, the probability is lower than 5%, and therefore, low. Reject the null hypothesis



C.The Single Sample z-Test

Now that an **Alpha Level of 5%** has been set, we must determine the probability of our sample mean occurring.

We can do this by performing a **single sample z-test**.

In other words, we will **calculate a z-score for our sample mean**.

In this context, a **z-score** will express **how many standard errors our sample mean is away from H0**.

For example, a z-score of **$z = 1.5$** would indicate that our sample mean is **1.5 standard errors above** the mean of the distribution of sample means.

a z-score of **$z = -1.5$** would indicate that our sample mean is **1.5 standard errors below** the mean of the distribution of sample means.

A single sample z-test is calculated using the following formula

$$z = \frac{M - \mu}{\sigma_M}$$

- $\sigma_M = \sigma / \sqrt{n}$

z = the z-score for our sample mean
 M = our sample mean
 μ = population mean
 σ_M = standard error of the mean.
 Note that **population standard deviation is known**.

It is a single sample test: compares the sample mean with a given number and asks is there a difference

D. Summary

When determining if our sample mean provides evidence to support the null hypothesis or not, we...

Set an alpha level. Here we have used 5%. This defines which sample means are expected and which are unlikely if the null hypothesis is true.

In a normal distribution of sample means, the 5% alpha level corresponds to ± 1.96 standard errors from the null hypothesised mean value.

Calculate a z-score for our sample mean. Standard error is based on population standard deviation in this instance.

If the z-score is less than 1.96, do not reject the null hypothesis.

If the z-score is greater than 1.96, reject the null hypothesis.

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6 The Single Sample t-test

A. z-test recap

$$z = \frac{M - \mu}{\sigma_M}$$

z-score: how many standard errors our sample mean is away from H_0 .

z = the z-score for our sample mean

M = our sample mean

μ = population mean

σ_M = standard error of the mean. $\sigma_M = \sigma / \sqrt{n}$

Note that population standard deviation is known.

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B. Handling the case of unknown population

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standard deviation

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However, population standard deviation is rarely known.

In instances when we don't know the population standard deviation, we can't use a **z-test** and the normal distribution to assess our sample mean.

That's okay! Instead, we can use a **t-test** and the '**t-distribution**'.

We use **SAMPLE standard deviation** as an **ESTIMATE of POPULATION standard deviation**. Single sample t-tests and z-tests are very similar otherwise.

Z-Test VS T-Test

Z-Test standard error (σ)	T-Test (estimate σ with s)
------------------------------------	-----------------------------------

$\sigma_M = \frac{\sigma}{\sqrt{n}}$	$s_M = \frac{s}{\sqrt{n}}$
Z-Test Formula	t-test formula
$z = \frac{M - \mu}{\sigma_M}$ <div> M = sample mean μ = population </div>	$t = \frac{M - \mu}{s_M}$

Almost all aspects of the process are the **same** when conducting either a single sample t-test or z-test.

We still use **Null Hypothesis Significance Testing**. We still assign a value that indicates no effect to the null hypothesis, and proceed assuming the null is true.

We still apply an alpha level of 5% and determine the probability of our mean occurring. The result determines whether the null hypothesis is rejected or not.

One Difference of t-distribution

- In the t-distribution, the critical limit corresponding to our alpha level of 5% will not be fixed at +/- 1.96 as it was with the z-test and normal distribution.
- Instead, the t-distribution requires that we consider sample size and **degrees of freedom (df)**, when determining the probability of the sample mean.
 - ➔ Degrees of Freedom are one less than our sample size for a single sample t-test (**n-1**). n = sample size
- The critical limit varies along with **df**

Why do we use **n-1** when calculating sample standard deviation?

- We use n-1 as the process of inferring our sample is a representative of the population is an **estimation** – imprecise process
 - This process is **biased** because samples are often less variable (less spread out) than the population they come from
 - **The underestimation bias is corrected by using n-1 degrees of freedom**
 - Dividing by n-1 gives us **a larger result**

- Both tests have very similar calculations to get a probability for a sample mean
- The t-score gives the probability for the sample mean, just like the z-score
 - ➔ We **reject** the null hypothesis if the probability is **too small**, just as before
- However:
 - ➔ T distribution requires **degrees of freedom, df**, to calculate the **correct probability**
df = n – 1, where n= sample size

Example Does Head Injury Affect IQ?

- Imagine that we have a sample of 10 people with an acquired head injury. Has IQ been affected?
- The **null hypothesis** would be that **head injury has not affected IQ**.
H₀: μ = 100
- Sample data: **M** = 94.20, **s** = 6.16, **n** = 10

- T-score $t = \frac{M - \mu}{s_M}$

Step 1: Calculate s_M (standard error)

$$t = \frac{M - \mu}{s_M} \rightarrow s_M = \frac{s}{\sqrt{n}}$$

$$s_M = \frac{6.16}{\sqrt{10}}$$

$$s_M = \frac{6.16}{3.16}$$

$$s_M = 1.95$$

Step 2: Calculate the t-score for our sample mean

$$t = \frac{94.2 - 100}{1.95}$$

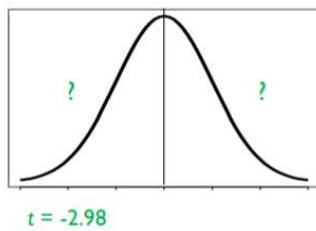
$$t = \frac{-5.8}{1.95}$$

$$t = -2.98$$

$$t = \frac{M - \mu}{s_M}$$

Is the t-score of **-2.98** for our sample mean more extreme than the critical limit, taking **df** into account?

- Yes! A t-score of **-2.98** is extreme for df = n-1=10-1= 9.



This means that the probability of our sample mean occurring, assuming the null hypothesis is true, is less than the alpha level of 5%. In fact, it is just 1.6%

Summary **Assignment Project Exam Help**

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