

Probability and Statistics

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Topics

- 1 Test for the Difference Between Two Normal Means $\mu_1 - \mu_2$ when σ_1 and σ_2 are Known
- 2 Large Sample Test for the Difference Between Two General Population Means $\mu_1 - \mu_2$
- 3 Test for the Difference Between Two Normal Means $\mu_1 - \mu_2$ When σ_1 and σ_2 are Unknown

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Objectives

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- Carry out:
 - Two-sample z test for the difference between two normal means $\mu_1 - \mu_2$ when σ_1 and σ_2 are known.
 - Two-sample z test for the difference between two general population means $\mu_1 - \mu_2$ when m and n are large.
 - Two-sample t test for the difference between two normal means $\mu_1 - \mu_2$ when σ_1 and σ_2 are unknown.

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Test for the Difference Between Two Normal Means $\mu_1 - \mu_2$ when σ_1 and σ_2 are Known (9.1)

- Suppose
 1. X_1, X_2, \dots, X_m are a random sample from a $\mathbf{N}(\mu_1, \sigma_1)$ population.
 2. Y_1, Y_2, \dots, Y_n are a random sample from a $\mathbf{N}(\mu_2, \sigma_2)$ population.
 3. The population means μ_1 and μ_2 are **unknown** but the standard deviations σ_1 and σ_2 are **known**.
 4. The X and Y samples are **independent** of each other.
- We'll see how to test whether μ_1 and μ_2 are different from each other.

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- When σ_1 and σ_2 are known, the appropriate test is called the **two-sample z test**
- The difference $\bar{X} - \bar{Y}$ between the two sample means is an **estimator** of the (unknown) difference between the population means $\mu_1 - \mu_2$.
- $\bar{X} - \bar{Y}$ is a difference between two **normal** random variables, so it too follows a **normal distribution**.

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Normality of $\bar{X} - \bar{Y}$: If X_1, X_2, \dots, X_m is a random sample from a $N(\mu_1, \sigma_1)$ distribution, and Y_1, Y_2, \dots, Y_n is a random sample from a $N(\mu_2, \sigma_2)$ distribution, and the two samples are **independent** of each other, then

$$\bar{X} - \bar{Y} \sim N\left(\mu_1 - \mu_2, \sqrt{\frac{\sigma_1^2}{m} + \frac{\sigma_2^2}{n}}\right).$$

It follows that

$$\frac{\bar{X} - \bar{Y} - (\mu_1 - \mu_2)}{\sqrt{\sigma_1^2/m + \sigma_2^2/n}} \sim N(0, 1).$$

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Proof: From Slides 14,

$$\bar{X} \sim N\left(\mu_1, \frac{\sigma_1}{\sqrt{m}}\right) \quad \text{and} \quad \bar{Y} \sim N\left(\mu_2, \frac{\sigma_2}{\sqrt{n}}\right).$$

Furthermore, $\bar{X} - \bar{Y}$ is a linear combination of \bar{X} and \bar{Y} (which are independent because the two samples are), so (also from Slides 14)

$$\bar{X} - \bar{Y} \sim N\left(\mu_1 - \mu_2, \sqrt{\frac{\sigma_1^2}{m} + \frac{\sigma_2^2}{n}}\right).$$

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- We're seeking to "disprove" the claim that μ_1 is equal to μ_2 , so the **null hypothesis** is that they *are* equal.

Null Hypothesis:

$$H_0 : \mu_1 - \mu_2 = 0$$

(H_0 could also be written as $H_0 : \mu_1 = \mu_2$.)

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- The **alternative hypothesis** will depend on what we're trying to "prove":

Alternative Hypothesis: The alternative hypothesis will be one of

1. $H_a : \mu_1 - \mu_2 > 0$ (one-sided, upper-tailed)
2. $H_a : \mu_1 - \mu_2 < 0$ (one-sided, lower-tailed)
3. $H_a : \mu_1 - \mu_2 \neq 0$ (two-sided, two-tailed)

depending on what we're trying to verify using the data.

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- The **test statistic** for the **two-sample z test** for $\mu_1 - \mu_2$ is

Two-Sample Z Test Statistic:

$$Z = \frac{\bar{X} - \bar{Y} - 0}{\sqrt{\sigma_1^2/m + \sigma_2^2/n}}$$

When H_0 is true, $Z \sim N(0, 1)$.

- Z measures how many standard errors $\bar{X} - \bar{Y}$ is away from **zero**.

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- $\bar{X} - \bar{Y}$ is an estimator of the unknown difference between population means $\mu_1 - \mu_2$, so ...
 1. Z will be approximately **zero** if $\mu_1 - \mu_2 = 0$.
 2. It will be **positive** if $\mu_1 - \mu_2 > 0$.
 3. It will be **negative** if $\mu_1 - \mu_2 < 0$.

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1. **Large positive** values of Z provide **evidence against H_0 in favor of**
 $H_a : \mu_1 - \mu_2 > 0$.
2. **Large negative** values of Z provide **evidence against H_0 in favor of**
 $H_a : \mu_1 - \mu_2 < 0$.
3. **Large positive and large negative** values of Z provide **evidence against H_0 in favor of**
 $H_a : \mu_1 - \mu_2 \neq 0$.

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- Recall that

$$\frac{\bar{X} - \bar{Y} - (\mu_1 - \mu_2)(\mu_1 - \mu_2)}{\sqrt{\sigma_1^2/m + \sigma_2^2/n}} \sim N(0, 1).$$

- It follows that if H_0 is true (so $\mu_1 - \mu_2 = 0$),

$$\frac{\bar{X} - \bar{Y} - 00}{\sqrt{\sigma_1^2/m + \sigma_2^2/n}} \sim N(0, 1).$$

Sampling Distribution of the Test Statistic Under H_0 :

If Z is the two-sample Z test statistic, then when

$$H_0: \mu_1 - \mu_2 = 0$$

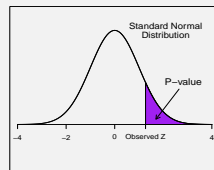
is true,

$$Z \sim N(0, 1).$$

- The **p-value** is the probability that just by chance (under H_0) we'd get a test statistic value as far from zero, in the direction predicted by H_a , as the observed value.

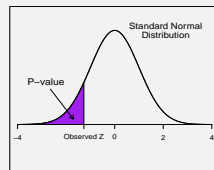
- P-value** = Area to the **right** of the observed z if the alternative hypothesis is $H_a: \mu_1 - \mu_2 > 0$.

P-Value for Upper-Tailed Z Test

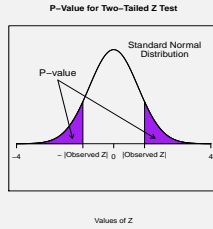


- P-value** = Area to the **left** of the observed z if the alternative hypothesis is $H_a: \mu_1 - \mu_2 < 0$.

P-Value for Lower-Tailed Z Test



1. **P-value** = Area to the left of $-|z|$ and right of $|z|$ if the alternative hypothesis is $H_a : \mu_1 - \mu_2 \neq 0$.



Two-Sample Z Test for $\mu_1 - \mu_2$ when σ_1 and σ_2 are Known

Assumptions: The data x_1, x_2, \dots, x_m are a random sample from a $N(\mu_1, \sigma_1)$ distribution and y_1, y_2, \dots, y_n are a random sample from a $N(\mu_2, \sigma_2)$ distribution, where σ_1 and σ_2 are known. Also, the two samples are independent.

Null hypothesis: $H_0 : \mu_1 - \mu_2 = 0$.

Test statistic value: $z = \frac{\bar{x} - \bar{y} - 0}{\sqrt{\sigma_1^2/m + \sigma_2^2/n}}$

Decision rule: Reject H_0 if p-value $< \alpha$.

| Alternative hypothesis | P-value = area under $N(0, 1)$ distribution: |
|------------------------------|--|
| $H_a : \mu_1 - \mu_2 > 0$ | to the right of z |
| $H_a : \mu_1 - \mu_2 < 0$ | to the left of z |
| $H_a : \mu_1 - \mu_2 \neq 0$ | to the left of $- z $ and right of $ z $ |

Large Sample Test for the Difference Between Two General Population Means $\mu_1 - \mu_2$ (9.1)

- When the **sample sizes m and n are both large**, two things are true:

- Regardless of the shape of the populations, by the Central Limit Theorem,

$$Z = \frac{\bar{X} - \bar{Y} - (\mu_1 - \mu_2)}{\sqrt{\sigma_1^2/m + \sigma_2^2/n}} \sim N(0, 1) \quad (\text{approximately}).$$

- The **sample standard deviations S_1 and S_2** will remain fairly **constant** from one sample to the next, and **approximately equal to σ_1 and σ_2** .

- As a consequence, **when n is large**,

$$Z = \frac{\bar{X} - \bar{Y} - (\mu_1 - \mu_2)}{\sqrt{S_1^2/m + S_2^2/n}} \sim N(0, 1) \quad (\text{approximately}).$$

even if the samples are from **non-normal** populations.
 (Note that σ_1 and σ_2 were replaced by S_1 and S_2 above.)

- It follows that **if H_0 is true** (so $\mu_1 - \mu_2 = 0$),

$$\frac{\bar{X} - \bar{Y} - 0}{\sqrt{S_1^2/m + S_2^2/n}} \sim N(0, 1) \quad (\text{approximately}).$$

- Thus we can use this as our **test statistic** in a **two-sample z test for $\mu_1 - \mu_2$** .

Two-Sample Z Test Statistic:

$$Z = \frac{\bar{X} - \bar{Y} - 0}{\sqrt{S_1^2/m + S_2^2/n}}$$

When H_0 is true, $Z \sim N(0, 1)$.

Sampling Distribution of the Test Statistic Under H_0 :

If Z is the two-sample Z test statistic (from the previous slide), then when

$$H_0: \mu_1 - \mu_2 = 0$$

is true,

$$Z \sim N(0, 1).$$

- The **p-value** is the appropriate tail area under the $N(0, 1)$ curve.
- In practice, m and n are **large enough** if $m > 40$ and $n > 40$.

Two-Sample Z Test for $\mu_1 - \mu_2$ when m and n are Large

Assumptions: The data x_1, x_2, \dots, x_m are a random sample from any distribution whose mean and standard deviation are μ_1 and σ_1 and y_1, y_2, \dots, y_n are a random sample from any distribution whose mean and standard deviation are μ_2 and σ_2 . Also, the two samples are independent of each other.

Null hypothesis: $H_0: \mu_1 - \mu_2 = 0$.

Test statistic value: $z = \frac{\bar{x} - \bar{y} - 0}{\sqrt{s_1^2/m + s_2^2/n}}$.

Decision rule: Reject H_0 if p-value $< \alpha$.

Alternative hypothesis

- $H_a: \mu_1 - \mu_2 > 0$
- $H_a: \mu_1 - \mu_2 < 0$

P-value = area under $N(0, 1)$ distribution:

- to the right of z
- to the left of z

Test for the Difference Between Two Normal Means $\mu_1 - \mu_2$ When σ_1 and σ_2 are Unknown (9.2)

- It can be shown that if X_1, X_2, \dots, X_n and Y_1, Y_2, \dots, Y_n are independent random samples from two **normal** populations, the random variable

$$\frac{\bar{X} - \bar{Y} - (\mu_1 - \mu_2)}{\sqrt{S_1^2/m + S_2^2/n}} \sim t(\nu),$$

a **t distribution** with ν **degrees of freedom**, where

$$\nu = \frac{\left(\frac{s_1^2}{m} + \frac{s_2^2}{n}\right)^2}{\frac{(s_1^2/m)^2}{m-1} + \frac{(s_2^2/n)^2}{n-1}},$$

which should be truncated **down** to the nearest integer.

- It follows that if H_0 is true (so $\mu_1 - \mu_2 = 0$),

$$\frac{\bar{X} - \bar{Y} - 0}{\sqrt{S_1^2/m + S_2^2/n}} \sim t(\nu),$$

- Thus we can use this as our **test statistic** in a **two-sample t test for $\mu_1 - \mu_2$** .

- The **test statistic** for the **two-sample t test for $\mu_1 - \mu_2$** is

Two-Sample t Test Statistic:

$$T = \frac{\bar{X} - \bar{Y} - 0}{\sqrt{S_1^2/m + S_2^2/n}}.$$

When H_0 is true, $T \sim t(\nu)$.

Sampling Distribution of the Test Statistic Under H_0 :

If T is the two-sample t test statistic, then when

$$H_0 : \mu_1 - \mu_2 = 0$$

is true,

$$T \sim t(\nu).$$

- The **p -value** is the appropriate tail area under the $t(\nu)$ curve.

Two-Sample t Test for $\mu_1 - \mu_2$

Assumptions: The data x_1, x_2, \dots, x_m are a random sample from a $N(\mu_1, \sigma_1)$ distribution and y_1, y_2, \dots, y_n are a random sample from a $N(\mu_2, \sigma_2)$ distribution. Also, the two samples are independent of each other.

Null hypothesis: $H_0 : \mu_1 - \mu_2 = 0$.

Test statistic value: $t = \frac{\bar{x} - \bar{y} - 0}{\sqrt{s_1^2/m + s_2^2/n}}$.

Decision rule: Reject H_0 if p -value $< \alpha$.

Alternative hypothesis

$H_a : \mu_1 - \mu_2 > 0$
 $H_a : \mu_1 - \mu_2 < 0$
 $H_a : \mu_1 - \mu_2 \neq 0$

P-value = area under $t(\nu)$ distribution*:

to the right of t
 to the left of t
 to the left of $-|t|$ and right of $|t|$

* $t(\nu)$ is the t distribution with d.f. ν given a few slides back.

Example

An engineer in a garment factory must compare two different work sequences for measuring the strength of polyester fibers **to decide if one sequence is, on average, faster than the other.**

Twelve workers are randomly assigned to two groups of **six workers each.**

The first group measures the strength of the fabric using **Work Sequence 1** and the second measures it using **Work Sequence 2.**

The following data are the **completion times (in seconds)** for each group:

| Work Sequence 1 | Work Sequence 2 |
|-----------------|-----------------|
| 220 | 247 |
| 235 | 223 |
| 214 | 215 |
| 197 | 219 |
| 206 | 207 |
| 214 | 236 |

The **summary statistics** for the two groups are:

| Work Sequence 1 | Work Sequence 2 |
|-------------------|-------------------|
| $m = 6$ | $n = 6$ |
| $\bar{x} = 214.3$ | $\bar{y} = 224.5$ |
| $s_1 = 12.9$ | $s_2 = 14.6$ |

We'll carry out a **two-sample t test** to decide **which work sequence, if any, is faster.**

The **hypotheses** are

$$H_0 : \mu_1 - \mu_2 = 0$$
$$H_a : \mu_1 - \mu_2 \neq 0$$

where μ_1 and μ_2 are the true (unknown) population mean completion times.

The observed **test statistic** is

$$\begin{aligned}
 t &= \frac{\bar{x} - \bar{y} - 0}{\sqrt{s_1^2/m + s_2^2/n}} \\
 &= \frac{214.3 - 224.5 - 0}{\sqrt{12.9^2/6 + 14.6^2/6}} \\
 &= -1.28.
 \end{aligned}$$

Thus the observed difference between **sample mean** completion times, $\bar{x} - \bar{y} = -10.2$, is about **1.28 standard errors below zero**.

The **p-value** is the **probability** that we'd get a t value this far away from zero (in either direction) by chance if there was **no difference** in the **population means** μ_1 and μ_2 .

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Under H_0 , the test statistic would follow a $t(\nu)$ **distribution** with **degrees of freedom**

$$\nu = \frac{\left(\frac{s_1^2}{m} + \frac{s_2^2}{n}\right)^2}{\frac{(s_1^2/m)^2}{m-1} + \frac{(s_2^2/n)^2}{n-1}} = \frac{\left(\frac{12.9^2}{6} + \frac{14.6^2}{6}\right)^2}{\frac{(12.9^2/6)^2}{6-1} + \frac{(14.6^2/6)^2}{6-1}} = 9.8,$$

which we round **down to 9**.

From the **two tail** areas of the $t(9)$ **distribution**, to the **left of -1.28** and **right of 1.28**,

$$\text{p-value} = 2(0.116) = 0.232.$$

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Thus we'd get a result like the one we got **23.2%** of the time **even if** the **population mean** completion times μ_1 and μ_2 were equal.

Using a **level of significance** $\alpha = 0.05$, the **decision rule** is

- Reject H_0 if p-value < 0.05 .
- Fail to reject H_0 if p-value ≥ 0.05 .

Because $0.232 \geq 0.05$, we **fail to reject H_0** .

There's **no statistically significant evidence** for any difference in the mean completion times for the two work sequences.

The observed difference can be explained by chance variation (sampling error).

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