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## Difference between Two Means (Independent Groups)

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### Prerequisites

Sampling Distribution of Difference between Means, Confidence Intervals, Confidence Interval on the Difference between Means, Logic of Hypothesis Testing, Testing a Single Mean

#### Learning Objectives

- 1. State the assumptions for testing the difference between two means
- 2. Estimate the population variance assuming homogeneity of variance
- 3. Compute the standard error of the difference between means
- 4. Compute t and p for the difference between means
- 5. Format data for computer analysis

It is much more common for a researcher to be interested in the difference between means than in the specific values of the means themselves. This covers how to test for differences between means from two separate ground subjects. A <u>later section</u> describes how to test for differences between the of two conditions in designs where only one group of subjects is used and subject is tested in each condition.

We take as an example the data from the "Animal Research" case stu this experiment, students rated (on a 7-point scale) whether they though research is wrong. The sample sizes, means, and variances are shown sel for males and females in Table 1.

Table 1. Means and Variances in Animal Research study.

III Alliniai Research study.				
Group	n	Mean	Variance	
Females	17	5.353	2.743	
Males	17	3.882	2.985	

As you can see, the females rated animal research as more wrong than dimales. This sample difference between the female mean of 5.35 and the imean of 3.88 is 1.47. However, the gender difference in this particular sample very important. What is important is whether there is a difference in the population means.

In order to test whether there is a difference between population mea are going to make three assumptions:

- 1. The two populations have the same variance. This assumption is the assumption of *homogeneity of variance*.
- 2. The populations are *normally distributed*.
- 3. Each value is sampled <u>independently</u> from each other value. Thi assumption requires that each subject provide only one value. I subject provides two scores, then the scores are not independently analysis of data with two scores per subject is shown in the second the <u>correlated t test</u> later in this chapter.

The consequences of violating the first two assumptions are investigated <u>simulation in the next section</u>. For now, suffice it to say that small-to-mod violations of assumptions 1 and 2 do not make much difference. It is import to violate assumption 3.

We saw the following general formula for significance testing in the se  $\underline{\text{testing a single mean}}$ :

 $t = \frac{\text{statistic-hypothesized value}}{\text{estimated standard error of the statistic}}$ 

In this case, our statistic is the difference between sample means and hypothesized value is 0. The hypothesized value is the null hypothesis that difference between population means is 0.

We continue to use the data from the "Animal Research" case study a

compute a significance test on the difference between the mean score of females and the mean score of the males. For this calculation, we will ma three assumptions specified above.

The first step is to compute the statistic, which is simply the differenc between means.

$$M_1 - M_2 = 5.3529 - 3.8824 = 1.4705$$

Since the hypothesized value is 0, we do not need to subtract it from the The next step is to compute the estimate of the standard error of the In this case, the statistic is the difference between means, so the estimat standard error of the statistic is  $(S_{M_1-M_2})$ . Recall from the <u>relevant section</u> chapter on sampling distributions that the formula for the standard error difference between means is:

$$\sigma_{M_1-M_2} = \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}} = \sqrt{\frac{\sigma^2}{n} + \frac{\sigma^2}{n}} = \sqrt{\frac{2\sigma^2}{n}}$$

In order to estimate this quantity, we estimate  $\sigma^2$  and use that estimate i of  $\sigma^2$ . Since we are assuming the two population variances are the same, estimate this variance by averaging our two sample variances. Thus, our of variance is computed using the following formula:

$$MSE = \frac{s_1^2 + s_2^2}{2}$$

where MSE is our estimate of  $\sigma^2$ . In this example,

$$MSE = (2.743 + 2.985)/2 = 2.864.$$

Since n (the number of scores in each group) is 17,

$$S_{M_1-M_2} = \sqrt{\frac{2MSE}{n}} = \sqrt{\frac{(2)(2.864)}{17}} = 0.5805.$$

The next step is to compute t by plugging these values into the formula:

$$t = 1.4705/.5805 = 2.533.$$

Finally, we compute the probability of getting a t as large or larger than 2 as small or smaller than -2.533. To do this, we need to know the <u>degrees freedom</u>. The degrees of freedom is the number of independent estimates variance on which MSE is based. This is equal to  $(n_1 - 1) + (n_2 - 1)$ , wher the sample size of the first group and  $n_2$  is the sample size of the second For this example,  $n_1 = n_2 = 17$ . When  $n_1 = n_2$ , it is conventional to use "refer to the sample size of each group. Therefore, the degrees of freedom  $n_1 = n_2 = 17$ .

Once we have the degrees of freedom, we can use the  $\underline{t}$  distribution  $\underline{c}$  to find the probability. Figure 1 shows that the probability value for a two-test is 0.0164. The two-tailed test is used when the null hypothesis can be rejected regardless of the direction of the effect. As shown in Figure 1, it probability of a t < -2.533 or a t > 2.533.

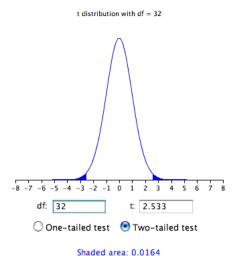


Figure 1. The two-tailed probability.

The results of a one-tailed test are shown in Figure 2. As you can see, the probability value of 0.0082 is half the value for the two-tailed test.

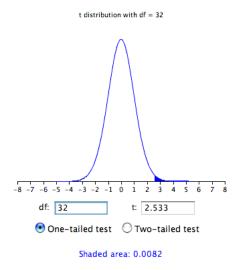


Figure 2. The one-tailed probability.

# Online Calculator: t distribution

FORMATTING DATA FOR COMPUTER ANALYSIS

Most computer programs that compute t tests require your data to be in  $\epsilon$  form. Consider the data in Table 2.

Table 2. Example Data.

Group 1	Group 2
3	2
4	6
5	8

Here there are two groups, each with three observations. To format these a computer program, you normally have to use two variables: the first sp the group the subject is in and the second is the score itself. The reformary version of the data in Table 2 is shown in Table 3.

Table 3. Reformatted

Data. **G**1

1	4
1	5
2	2
2	6
2	8

To use Analysis Lab to do the calculations, you would copy the data and tl

- Click the "Enter/Edit Data" button. (You may be warned that for secureasons you must use the keyboard shortcut for pasting data.)
- 2. Paste your data.
- 3. Click "Accept Data."
- 4. Set the Dependent Variable to Y.
- 5. Set the Grouping Variable to G.
- 6. Click the "t-test/confidence interval" button.

The t value is -0.718, the df = 4, and p = 0.512.

COMPUTATIONS FOR UNEQUAL SAMPLE SIZES (OPTIONAL)

The calculations are somewhat more complicated when the sample sizes equal. One consideration is that MSE, the estimate of variance, counts the group with the larger sample size more than the group with the smaller size. Computationally, this is done by computing the sum of squares errors follows:

$$SSE = \sum (X - M_1)^2 + \sum (X - M_2)^2$$

where  $M_1$  is the mean for group 1 and  $M_2$  is the mean for group 2. Consi following small example:

Table 4. Unequal n.

Group 1	Group 2
3	2
4	4
5	

$$M_1 = 4$$
 and  $M_2 = 3$ .

SSE = 
$$(3-4)^2 + (4-4)^2 + (5-4)^2 + (2-3)^2 + (4-3)^2 = 4$$

Then, MSE is computed by: MSE = SSE/df

where the degrees of freedom (df) is computed as before:  $df = (n_1 - 1) + (n_2 - 1) = (3 - 1) + (2 - 1) = 3$ . MSE = SSE/df = 4/3 = 1.333.

The formula

$$S_{M_1 - M_2} = \sqrt{\frac{2MSE}{n}}$$

is replaced by

$$S_{M_1-M_2} = \sqrt{\frac{2MSE}{n_h}}$$

where  $n_{\mbox{\scriptsize h}}$  is the harmonic mean of the sample sizes and is computed as 1

$$n_h = \frac{2}{1/n_1 + 1/n_2} = \frac{2}{1/3 + 1/2} = 2.4.$$

and

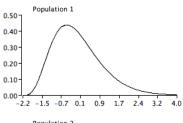
$$S_{M_1-M_2} = \sqrt{\frac{(2)(1.333)}{2.4}} = 1.054.$$

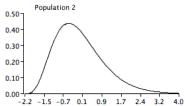
Therefore,

$$t = (4-3)/1.054 = 0.949$$

and the two-tailed p = 0.413.

**Question 1 out of 9.**The graphs show a violation of the assumption of (check all that apply)





- normality
- homogeneity of variance

Check Answer | Previous Question | Next Question

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