power twomeans — Power analysis for a two-sample means test

Description

power twomeans computes sample size, power, or the experimental-group mean for a two-sample means test. By default, it computes sample size for the given power and the values of the control-group and experimental-group means. Alternatively, it can compute power for given sample size and values of the control-group and experimental-group means or the experimental-group mean for given sample size, power, and the control-group mean. For power and sample-size analysis in a cluster randomized design, see [PSS-2] power twomeans, cluster. Also see [PSS-2] power for a general introduction to the power command using hypothesis tests.

For precision and sample-size analysis for a CI for the difference between two means from independent samples, see [PSS-3] ciwidth twomeans.

Quick start

Sample size for a test of \( H_0: \mu_1 = \mu_2 \) versus \( H_a: \mu_1 \neq \mu_2 \) given alternative control-group mean \( m_1 = 8 \) and alternative experimental-group mean \( m_2 = 12 \) with shared standard deviation of 9 using default power of 0.8 and significance level \( \alpha = 0.05 \)

\[ \text{power twomeans 8 12, sd(9)} \]

As above, but for \( m_2 \) equal to 10, 11, 12, 13, and 14

\[ \text{power twomeans 8 (10(1)14), sd(9)} \]

As above, but display results in a graph of sample size versus \( m_2 \)

\[ \text{power twomeans 8 (10(1)14), sd(9) graph} \]

As above, but specify different standard deviations \( s_1 = 7 \) and \( s_2 = 10 \)

\[ \text{power twomeans 8 (10(1)14), sd1(7) sd2(10) graph} \]

Sample size for one-sided test with power of 0.9

\[ \text{power twomeans 8 12, sd(9) power(.9) onesided} \]

Same as above, specified as \( \mu_1 \) and difference between means \( m_2 - m_1 = 4 \)

\[ \text{power twomeans 8, sd(9) power(.9) onesided diff(4)} \]

Power for a total sample size of 74 with balanced group sizes

\[ \text{power twomeans 8 12, sd(9) n(74)} \]

As above, but for sample sizes of 45 and 30 in groups 1 and 2, respectively

\[ \text{power twomeans 8 12, sd(9) n1(45) n2(30)} \]

Effect size and target mean difference for a sample size of 200 with power of 0.8

\[ \text{power twomeans 8, sd(9) power(.8) n(200)} \]
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Menu
Statistics > Power, precision, and sample size

Syntax

Compute sample size

```
power twomeans m1 m2 [ , power(numlist) options ]
```

Compute power

```
power twomeans m1 m2 , n(numlist) [ options ]
```

Compute effect size and experimental-group mean

```
power twomeans m1 , n(numlist) power(numlist) [ options ]
```

where \( m_1 \) is the mean in the control (reference) group and \( m_2 \) is the mean in the experimental (comparison) group. \( m_1 \) and \( m_2 \) may each be specified either as one number or as a list of values in parentheses (see \[U\] 11.1.8 numlist).
## power twomeans — Power analysis for a two-sample means test

<table>
<thead>
<tr>
<th>options</th>
<th>Description</th>
</tr>
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<tbody>
<tr>
<td><strong>Main</strong></td>
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</tr>
<tr>
<td><em>alpha</em>(numlist)</td>
<td>significance level; default is alpha(0.05)</td>
</tr>
<tr>
<td><em>power</em>(numlist)</td>
<td>power; default is power(0.8)</td>
</tr>
<tr>
<td><em>beta</em>(numlist)</td>
<td>probability of type II error; default is beta(0.2)</td>
</tr>
<tr>
<td><em>n</em>(numlist)</td>
<td>total sample size; required to compute power or effect size</td>
</tr>
<tr>
<td><em>n1</em>(numlist)</td>
<td>sample size of the control group</td>
</tr>
<tr>
<td><em>n2</em>(numlist)</td>
<td>sample size of the experimental group</td>
</tr>
<tr>
<td><em>nratio</em>(numlist)</td>
<td>ratio of sample sizes, N2/N1; default is nratio(1), meaning equal group sizes</td>
</tr>
<tr>
<td>compute(N1</td>
<td>N2)</td>
</tr>
<tr>
<td>nfractional</td>
<td>allow fractional sample sizes</td>
</tr>
<tr>
<td><em>diff</em>(numlist)</td>
<td>difference between the experimental-group mean and the control-group mean, $m_2 - m_1$; specify instead of the experimental-group mean $m_2$</td>
</tr>
<tr>
<td><em>sd</em>(numlist)</td>
<td>common standard deviation of the control and the experimental groups assuming equal standard deviations in both groups; default is sd(1)</td>
</tr>
<tr>
<td><em>sd1</em>(numlist)</td>
<td>standard deviation of the control group; requires sd2()</td>
</tr>
<tr>
<td><em>sd2</em>(numlist)</td>
<td>standard deviation of the experimental group; requires sd1()</td>
</tr>
<tr>
<td>knownsds</td>
<td>request computation assuming known standard deviations for both groups; default is to assume unknown standard deviations</td>
</tr>
<tr>
<td>direction(upper</td>
<td>lower)</td>
</tr>
<tr>
<td>onesided</td>
<td>one-sided test; default is two sided</td>
</tr>
<tr>
<td>parallel</td>
<td>treat number lists in starred options or in command arguments as parallel when multiple values per option or argument are specified (do not enumerate all possible combinations of values)</td>
</tr>
<tr>
<td>Table</td>
<td></td>
</tr>
<tr>
<td>[no]table[(tablespec)]</td>
<td>suppress table or display results as a table; see [PSS-2] power, table</td>
</tr>
<tr>
<td>saving(filename [, replace])</td>
<td>save the table data to filename; use replace to overwrite existing filename</td>
</tr>
<tr>
<td>Graph</td>
<td></td>
</tr>
<tr>
<td>graph[(graphopts)]</td>
<td>graph results; see [PSS-2] power, graph</td>
</tr>
</tbody>
</table>
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Iteration

init(#) initial value for sample sizes or experimental-group mean
iterate(#) maximum number of iterations; default is iterate(500)
tolerance(#) parameter tolerance; default is tolerance(1e-12)
ftolerance(#) function tolerance; default is ftolerance(1e-12)
log suppress or display iteration log
dots suppress or display iterations as dots

cluster perform computations for a CRD; see [PSS-2] power twomeans, cluster

notitle suppress the title

* Specifying a list of values in at least two starred options, or at least two command arguments, or at least one starred option and one argument results in computations for all possible combinations of the values; see [U] 11.1.8 numlist. Also see the parallel option.

collect is allowed; see [U] 11.1.10 Prefix commands.

cluster and notitle do not appear in the dialog box.

where tablespec is

    column[ :label] [ column[ :label] [...] ] [ , tableopts ]

column is one of the columns defined below, and label is a column label (may contain quotes and compound quotes).

<table>
<thead>
<tr>
<th>column</th>
<th>Description</th>
<th>Symbol</th>
</tr>
</thead>
<tbody>
<tr>
<td>alpha</td>
<td>significance level</td>
<td>$\alpha$</td>
</tr>
<tr>
<td>power</td>
<td>power</td>
<td>$1 - \beta$</td>
</tr>
<tr>
<td>beta</td>
<td>type II error probability</td>
<td>$\beta$</td>
</tr>
<tr>
<td>N</td>
<td>total number of subjects</td>
<td>$N$</td>
</tr>
<tr>
<td>N1</td>
<td>number of subjects in the control group</td>
<td>$N_1$</td>
</tr>
<tr>
<td>N2</td>
<td>number of subjects in the experimental group</td>
<td>$N_2$</td>
</tr>
<tr>
<td>nratio</td>
<td>ratio of sample sizes, experimental to control</td>
<td>$N_2/N_1$</td>
</tr>
<tr>
<td>delta</td>
<td>effect size</td>
<td>$\delta$</td>
</tr>
<tr>
<td>m1</td>
<td>control-group mean</td>
<td>$\mu_1$</td>
</tr>
<tr>
<td>m2</td>
<td>experimental-group mean</td>
<td>$\mu_2$</td>
</tr>
<tr>
<td>diff</td>
<td>difference between the experimental-group mean and the control-group mean</td>
<td>$\mu_2 - \mu_1$</td>
</tr>
<tr>
<td>sd</td>
<td>common standard deviation</td>
<td>$\sigma$</td>
</tr>
<tr>
<td>sd1</td>
<td>control-group standard deviation</td>
<td>$\sigma_1$</td>
</tr>
<tr>
<td>sd2</td>
<td>experimental-group standard deviation</td>
<td>$\sigma_2$</td>
</tr>
<tr>
<td>target</td>
<td>target parameter; synonym for m2</td>
<td></td>
</tr>
<tr>
<td>_all</td>
<td>display all supported columns</td>
<td></td>
</tr>
</tbody>
</table>

Column beta is shown in the default table in place of column power if specified.
Columns nratio, diff, sd, sd1, and sd2 are shown in the default table if the corresponding options are specified.
Options

Main

alpha(), power(), beta(), n(), n1(), n2(), nratio(), compute(), nfractional; see [PSS-2] power.

diff(numlist) specifies the difference between the experimental-group mean and the control-group mean, \( m_2 - m_1 \). You can specify either the experimental-group mean \( m_2 \) as a command argument or the difference between the two means in diff(). If you specify diff(#), the experimental-group mean is computed as \( m_2 = m_1 + # \). This option is not allowed with the effect-size determination.

sd(numlist) specifies the common standard deviation of the control and the experimental groups assuming equal standard deviations in both groups. The default is sd(1).

sd1(numlist) specifies the standard deviation of the control group. If you specify sd1(), you must also specify sd2().

sd2(numlist) specifies the standard deviation of the experimental group. If you specify sd2(), you must also specify sd1().

knownsds requests that standard deviations of each group be treated as known in the computations. By default, standard deviations are treated as unknown, and the computations are based on a two-sample \( t \) test, which uses a Student’s \( t \) distribution as a sampling distribution of the test statistic. If knownsds is specified, the computation is based on a two-sample \( z \) test, which uses a normal distribution as the sampling distribution of the test statistic.

direction(), onesided, parallel; see [PSS-2] power.

Table

table, table(); notable; see [PSS-2] power, table.

saving(); see [PSS-2] power.

Graph

graph, graph(); see [PSS-2] power, graph. Also see the column table for a list of symbols used by the graphs.

Iteration

init(#) specifies the initial value for the estimated parameter. For sample-size determination, the estimated parameter is either the control-group size \( n_1 \) or, if compute(N2) is specified, the experimental-group size \( n_2 \). For the effect-size determination, the estimated parameter is the experimental-group mean \( m_2 \). The default initial values for a two-sided test are obtained as a closed-form solution for the corresponding one-sided test with the significance level \( \alpha/2 \). The default initial values for the \( t \) test computations are based on the corresponding large-sample normal approximation.

iterate(), tolerance(), ftolerance(), log, nolog, dots, nodots; see [PSS-2] power.

The following options are available with power twomeans but are not shown in the dialog box:
cluster; see [PSS-2] power twomeans, cluster.

notitle; see [PSS-2] power.
Remarks and examples

Remarks are presented under the following headings:

- Introduction
- Using power twomeans
- Computing sample size
- Computing power
- Computing effect size and experimental-group mean
- Testing a hypothesis about two independent means

This entry describes the `power twomeans` command and the methodology for power and sample-size analysis for a two-sample means test. See [PSS-2] Intro (power) for a general introduction to power and sample-size analysis and [PSS-2] power for a general introduction to the `power` command using hypothesis tests. Also see [PSS-2] power twomeans, cluster for power and sample-size analysis in a cluster randomized design.

Introduction

The analysis of means is one of the most commonly used approaches in a wide variety of statistical studies. Many applications lead to the study of two independent means, such as studies comparing the average mileage of foreign and domestic cars, the average SAT scores obtained from two different coaching classes, the average yields of a crop due to a certain fertilizer, and so on. The two populations of interest are assumed to be independent.

This entry describes power and sample-size analysis for the inference about two population means performed using hypothesis testing. Specifically, we consider the null hypothesis $H_0: \mu_2 = \mu_1$ versus the two-sided alternative hypothesis $H_a: \mu_2 \neq \mu_1$, the upper one-sided alternative $H_a: \mu_2 > \mu_1$, or the lower one-sided alternative $H_a: \mu_2 < \mu_1$.

The considered two-sample tests rely on the assumption that the two random samples are normally distributed or that the sample size is large. Suppose that the two samples are normally distributed. If variances of the considered populations are known a priori, the test statistic has a standard normal distribution under the null hypothesis, and the corresponding test is referred to as a two-sample $z$ test. If variances of the two populations are not known, then the null sampling distribution of the test statistic depends on whether the two variances are assumed to be equal. If the two variances are assumed to be equal, the test statistic has an exact Student’s $t$ distribution under the null hypothesis. The corresponding test is referred to as a two-sample $t$ test. If the two variances are not equal, then the distribution can only be approximated by a Student’s $t$ distribution; the degrees of freedom is approximated using Satterthwaite’s method. We refer to this test as Satterthwaite’s $t$ test. For a large sample, the distribution of the test statistic is approximately normal, and the corresponding test is a large-sample $z$ test.

The `power twomeans` command provides power and sample-size analysis for the above tests.

Using power twomeans

`power twomeans` computes sample size, power, or experimental-group mean for a two-sample means test. All computations are performed for a two-sided hypothesis test where, by default, the significance level is set to 0.05. You may change the significance level by specifying the `alpha()` option. You can specify the `onesided` option to request a one-sided test. By default, all computations assume a balanced- or equal-allocation design; see [PSS-4] Unbalanced designs for a description of how to specify an unbalanced design.
By default, all computations are for a two-sample $t$ test, which assumes equal and unknown standard deviations. By default, the common standard deviation is set to one but may be changed by specifying the `sd()` option. To specify different standard deviations, use the respective `sd1()` and `sd2()` options. These options must be specified together and may not be used in combination with `sd()`. When `sd1()` and `sd2()` are specified, the computations are based on Satterthwaite’s $t$ test, which assumes unequal and unknown standard deviations. If standard deviations are known, use the `knownsds` option to request that computations be based on a two-sample $z$ test.

To compute the total sample size, you must specify the control-group mean $m_1$, the experimental-group mean $m_2$, and, optionally, the power of the test in the `power()` option. The default power is set to 0.8.

Instead of the total sample size, you can compute one of the group sizes given the other one. To compute the control-group sample size, you must specify the `compute(N1)` option and the sample size of the experimental group in the `n2()` option. Likewise, to compute the experimental-group sample size, you must specify the `compute(N2)` option and the sample size of the control group in the `n1()` option.

To compute power, you must specify the total sample size in the `n()` option, the control-group mean $m_1$, and the experimental-group mean $m_2$.

Instead of the experimental-group mean $m_2$, you may specify the difference $m_2 - m_1$ between the experimental-group mean and the control-group mean in the `diff()` option when computing sample size or power.

To compute effect size, the difference between the experimental-group mean and the null mean, and the experimental-group mean, you must specify the total sample size in the `n()` option, the power in the `power()` option, the control-group mean $m_1$, and, optionally, the direction of the effect. The direction is upper by default, `direction(upper)`, which means that the experimental-group mean is assumed to be larger than the specified control-group value. You can change the direction to be lower, which means that the experimental-group mean is assumed to be smaller than the specified control-group value, by specifying the `direction(lower)` option.

Instead of the total sample size `n()`, you can specify individual group sizes in `n1()` and `n2()`, or specify one of the group sizes and `nratio()` when computing power or effect size. Also see Two samples in [PSS-4] Unbalanced designs for more details.

In the following sections, we describe the use of `power twomeans` accompanied by examples for computing sample size, power, and experimental-group mean.

### Computing sample size

To compute sample size, you must specify the control-group mean $m_1$, the experimental-group mean $m_2$, and, optionally, the power of the test in the `power()` option. A default power of 0.8 is assumed if `power()` is not specified.

> **Example 1: Sample size for a two-sample means test**

Consider a study investigating the effects of smoking on lung function of males. The response variable is forced expiratory volume (FEV), measured in liters (L), where better lung function implies higher values of FEV. We wish to test the null hypothesis $H_0$: $\mu_1 = \mu_2$ versus a two-sided alternative hypothesis $H_a$: $\mu_1 \neq \mu_2$, where $\mu_1$ and $\mu_2$ are the mean FEV for nonsmokers and smokers, respectively.

Suppose that the mean FEV from previous studies was reported to be 3 L for nonsmokers and 2.7 L for smokers. We are designing a new study and wish to find out how many subjects we need
to enroll so that the power of a 5%-level two-sided test to detect the specified difference between means is at least 80%. We assume equal numbers of subjects in each group and a common standard deviation of 1.

```
. power twomeans 3 2.7
Performing iteration ... 
Estimated sample sizes for a two-sample means test 
t test assuming sd1 = sd2 = sd
H0: m2 = m1 versus Ha: m2 != m1
Study parameters:
    alpha =    0.0500
    power =    0.8000
    delta =    -0.3000
    m1 =     3.0000
    m2 =     2.7000
    sd =     1.0000
Estimated sample sizes:
    N =      352
    N per group =    176
```

We need a total sample of 352 subjects, 176 per group, to detect the specified mean difference between the smoking and nonsmoking groups with 80% power using a two-sided 5%-level test.

The default computation is for the case of equal and unknown standard deviations, as indicated by the output. You can specify the `knownsd` option to request the computation assuming known standard deviations.

Example 2: Sample size assuming unequal standard deviations

Instead of assuming equal standard deviations as in example 1, we use the estimates of the standard deviations from previous studies as our hypothetical values. The standard deviation of FEV for the nonsmoking group was reported to be 0.8 L and that for the smoking group was reported to be 0.7 L. We specify standard deviations in the `sd1()` and `sd2()` options.

```
. power twomeans 3 2.7, sd1(0.8) sd2(0.7)
Performing iteration ... 
Estimated sample sizes for a two-sample means test 
Satterthwaite’s t test assuming unequal variances
H0: m2 = m1 versus Ha: m2 != m1
Study parameters:
    alpha =    0.0500
    power =    0.8000
    delta =    -0.3000
    m1 =     3.0000
    m2 =     2.7000
    sd1 =     0.8000
    sd2 =     0.7000
Estimated sample sizes:
    N =      200
    N per group =    100
```

The specified standard deviations are smaller than one, so we obtain a smaller required total sample size of 200 compared with example 1.
Example 3: Specifying difference between means

Instead of the mean FEV of 2.7 for the smoking group as in example 2, we can specify the difference between the two means of $2.7 - 3 = -0.3$ in the `diff()` option.

```
. power twomeans 3, sd1(0.8) sd2(0.7) diff(-0.3)
```

Performing iteration ... 
Estimated sample sizes for a two-sample means test
Satterthwaite’s $t$ test assuming unequal variances
$H_0$: $m_2 = m_1$ versus $H_a$: $m_2 \neq m_1$

Study parameters:
- $\alpha = 0.0500$
- $\beta = 0.8000$
- $\delta = -0.3000$
- $m_1 = 3.0000$
- $m_2 = 2.7000$
- $\text{diff} = -0.3000$
- $sd_1 = 0.8000$
- $sd_2 = 0.7000$

Estimated sample sizes:
- $N = 200$
- $N \text{ per group} = 100$

We obtain the same results as in example 2. The difference between means is now also reported in the output following the individual means.

Example 4: Computing one of the group sizes

Suppose we anticipate a sample of 120 nonsmoking subjects. We wish to compute the required number of subjects in the smoking group, keeping all other study parameters as in example 2. We specify the number of subjects in the nonsmoking group in the `n1()` option and specify the `compute(N2)` option.

```
. power twomeans 3 2.7, sd1(0.8) sd2(0.7) n1(120) compute(N2)
```

Performing iteration ...
Estimated sample sizes for a two-sample means test
Satterthwaite’s $t$ test assuming unequal variances
$H_0$: $m_2 = m_1$ versus $H_a$: $m_2 \neq m_1$

Study parameters:
- $\alpha = 0.0500$
- $\beta = 0.8000$
- $\delta = -0.3000$
- $m_1 = 3.0000$
- $m_2 = 2.7000$
- $sd_1 = 0.8000$
- $sd_2 = 0.7000$
- $N_1 = 120$

Estimated sample sizes:
- $N = 202$
- $N_2 = 82$

We need a sample of 82 smoking subjects given a sample of 120 nonsmoking subjects.
Example 5: Unbalanced design

By default, *power twomeans* computes sample size for a balanced- or equal-allocation design. If we know the allocation ratio of subjects between the groups, we can compute the required sample size for an unbalanced design by specifying the `nratio()` option.

Continuing with example 2, we will suppose that we anticipate to recruit twice as many smokers than nonsmokers; that is, \(n_2/n_1 = 2\). We specify the `nratio(2)` option to compute the required sample size for the specified unbalanced design.

```
. power twomeans 3 2.7, sd1(0.8) sd2(0.7) nratio(2)
Performing iteration ... 
Estimated sample sizes for a two-sample means test
Satterthwaite’s t test assuming unequal variances
H0: m2 = m1 versus Ha: m2 != m1
Study parameters:
  alpha =   0.0500
  power =   0.8000
  delta =  -0.3000
  m1 =     3.0000
  m2 =     2.7000
  sd1 =    0.8000
  sd2 =    0.7000
N2/N1 =   2.0000
Estimated sample sizes:
  N =    237
  N1 =     79
  N2 =   158
```

We need a total sample size of 237 subjects, which is larger than the required total sample size for the corresponding balanced design from example 2.

Also see Two samples in [PSS-4] Unbalanced designs for more examples of unbalanced designs for two-sample tests.

Computing power

To compute power, you must specify the total sample size in the `n()` option, the control-group mean \(m_1\), and the experimental-group mean \(m_2\).

Example 6: Power of a two-sample means test

Continuing with example 1, we will suppose that we have resources to enroll a total of only 250 subjects, assuming equal-sized groups. To compute the power corresponding to this sample size given the study parameters from example 1, we specify the total sample size in `n()`:
power twomeans 3 2.7, n(250)

Estimated power for a two-sample means test
t test assuming sd1 = sd2 = sd
H0: m2 = m1 versus Ha: m2 != m1

Study parameters:
- alpha = 0.0500
- N = 250
- N per group = 125
- delta = -0.3000
- m1 = 3.0000
- m2 = 2.7000
- sd = 1.0000

Estimated power:
- power = 0.6564

With a total sample of 250 subjects, we obtain a power of only 65.64%.

Example 7: Multiple values of study parameters

In this example, we assess the effect of varying the common standard deviation (assuming equal standard deviations in both groups) of FEV on the power of our study.

Continuing with example 6, we compute powers for a range of common standard deviations between 0.5 and 1.5 with the step size of 0.1. We specify the corresponding numlist in the sd() option.

```
. power twomeans 3 2.7, sd(0.5(0.1)1.5) n(250)
```

<table>
<thead>
<tr>
<th>alpha</th>
<th>power</th>
<th>N</th>
<th>N1</th>
<th>N2</th>
<th>delta</th>
<th>m1</th>
<th>m2</th>
<th>sd</th>
</tr>
</thead>
<tbody>
<tr>
<td>.05</td>
<td>.9972</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>.5</td>
</tr>
<tr>
<td>.05</td>
<td>.976</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>.6</td>
</tr>
<tr>
<td>.05</td>
<td>.9215</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>.7</td>
</tr>
<tr>
<td>.05</td>
<td>.8397</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>.8</td>
</tr>
<tr>
<td>.05</td>
<td>.747</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>.9</td>
</tr>
<tr>
<td>.05</td>
<td>.6564</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>1</td>
</tr>
<tr>
<td>.05</td>
<td>.5745</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>1.1</td>
</tr>
<tr>
<td>.05</td>
<td>.5036</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>1.2</td>
</tr>
<tr>
<td>.05</td>
<td>.4434</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>1.3</td>
</tr>
<tr>
<td>.05</td>
<td>.3928</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>1.4</td>
</tr>
<tr>
<td>.05</td>
<td>.3503</td>
<td>250</td>
<td>125</td>
<td>125</td>
<td>-.3</td>
<td>3</td>
<td>2.7</td>
<td>1.5</td>
</tr>
</tbody>
</table>

The power decreases from 99.7% to 35.0% as the common standard deviation increases from 0.5 to 1.5 L.

For multiple values of parameters, the results are automatically displayed in a table, as we see above. For more examples of tables, see [PSS-2] power, table. If you wish to produce a power plot, see [PSS-2] power, graph.
Computing effect size and experimental-group mean

Effect size $\delta$ for a two-sample means test is defined as the difference between the experimental-group mean and the control-group mean $\delta = \mu_2 - \mu_1$.

Sometimes, we may be interested in determining the smallest effect and the corresponding experimental-group mean that yield a statistically significant result for prespecified sample size and power. In this case, power, sample size, and control-group mean must be specified. In addition, you must also decide on the direction of the effect: upper, meaning $m_2 > m_1$, or lower, meaning $m_2 < m_1$. The direction may be specified in the `direction()` option; `direction(upper)` is the default.

Example 8: Minimum detectable change in the experimental-group mean

Continuing with example 6, we compute the smallest change in the mean of the smoking group that can be detected given a total sample of 250 subjects and 80% power, assuming equal-group allocation. To solve for the mean FEV of the smoking group, after the command name, we specify the nonsmoking-group mean of 3, total sample size `n(250)`, and power `power(0.8)`.

Because our initial study was based on the hypothesis that FEV for the smoking group is lower than that of the nonsmoking group, we specify the `direction(lower)` option to compute the smoking-group mean that is lower than the specified nonsmoking-group mean.

```plaintext
.power twomeans 3, n(250) power(0.8) direction(lower)
Performing iteration ...
Estimated experimental-group mean for a two-sample means test
  t test assuming sd1 = sd2 = sd
  H0: m2 = m1 versus Ha: m2 != m1; m2 < m1
Study parameters:
  alpha =  0.0500
  power =  0.8000
  N =  250
  N per group = 125
  m1 =  3.0000
  sd =  1.0000
Estimated effect size and experimental-group mean:
  delta = -0.3558
  m2 =  2.6442
```

We find that the minimum detectable value of the effect size is $-0.36$, which corresponds to the mean FEV of 2.64 for the smoking group.

Testing a hypothesis about two independent means

After data are collected, we can use the `ttest` command to test the equality of two independent means using a t test; see [R] `ttest` for details. In this section, we demonstrate the use of `ttesti`, the immediate form of the `test` command, which can be used to test a hypothesis using summary statistics instead of the actual data values.
Example 9: Two-sample \( t \) test

Consider an example from van Belle et al. (2004, 129), where newborn infants were divided into two groups: a treatment group, where infants received daily “walking stimulus” for eight weeks, and a control group, where no stimulus was provided. The goal of this study was to test whether receiving the walking stimulus during stages of infancy induces the walking ability to develop sooner.

The average number of months before the infants started walking was recorded for both groups. The authors provide estimates of the average of 10.125 months for the treatment group with estimated standard deviation of 1.447 months and 12.35 months for the control group with estimated standard deviation of 0.9618 months. The sample sizes for treatment and control groups were 6 and 5, respectively. We supply these estimates to the \texttt{ttesti} command and use the \texttt{unequal} option to perform a \( t \) test assuming unequal variances.

\[
\texttt{ttesti 6 10.125 1.447 5 12.35 0.9618, unequal}
\]

Two-sample \( t \) test with unequal variances

<table>
<thead>
<tr>
<th></th>
<th>Obs</th>
<th>Mean</th>
<th>Std. err.</th>
<th>Std. dev.</th>
<th>[95% conf. interval]</th>
</tr>
</thead>
<tbody>
<tr>
<td>( x )</td>
<td>6</td>
<td>10.125</td>
<td>.5907353</td>
<td>1.447</td>
<td>8.606467 – 11.64353</td>
</tr>
<tr>
<td>( y )</td>
<td>5</td>
<td>12.35</td>
<td>.43013</td>
<td>.9618</td>
<td>11.1577 – 13.54423</td>
</tr>
</tbody>
</table>

Combined: 11 | 11.13636 | .501552 | 1.66346 | 10.01884 – 12.25389 |

diff: -2.225 | .7307394 | -3.887894 – -.562106 |

diff = mean(\( x \)) - mean(\( y \))

\( H_0: \text{diff} = 0 \)

\( Satterthwaite’s \) degrees of freedom = 8.66326

\( H_a: \text{diff} < 0 \quad H_a: \text{diff} \neq 0 \quad H_a: \text{diff} > 0 \)

\( \Pr(T < t) = 0.0073 \quad \Pr(|T| > |t|) = 0.0145 \quad \Pr(T > t) = 0.9927 \)

We reject the null hypothesis of \( H_0: \mu_C = \mu_T \) against the two-sided alternative \( H_a: \mu_C \neq \mu_T \) at the 5% significance level; the \( p \)-value = 0.0145.

We use the estimates of this study to perform a sample-size analysis we would have conducted before a new study. In our analysis, we assume equal-group allocation.

\[
\texttt{power twomeans 10.125 12.35, power(0.8) sd1(1.447) sd2(0.9618)}
\]

Performing iteration ...

Estimated sample sizes for a two-sample means test

Satterthwaite’s \( t \) test assuming unequal variances

\( H_0: m_2 = m_1 \quad \text{versus} \quad H_a: m_2 \neq m_1 \)

Study parameters:

\[
\begin{align*}
\text{alpha} & = 0.0500 \\
n & = 14 \\
\text{power} & = 0.8000 \\
delta & = 2.2250 \\
m_1 & = 10.1250 \\
m_2 & = 12.3500 \\
ds_1 & = 1.4470 \\
ds_2 & = 0.9618
\end{align*}
\]

Estimated sample sizes:

\[
\begin{align*}
N & = 14 \\
N \text{ per group} & = 7
\end{align*}
\]

We find that the sample size required to detect a difference of 2.225 \((12.35 - 10.125 = 2.225)\) given the control-group standard deviation of 1.447 and the experimental-group standard deviation of 0.9618 using a 5%-level two-sided test is 7 in each group.
power twomeans stores the following in r():

Scalars
- r(alpha) : significance level
- r(power) : power
- r(beta) : probability of a type II error
- r(delta) : effect size
- r(N) : total sample size
- r(N_a) : actual sample size
- r(N1) : sample size of the control group
- r(N2) : sample size of the experimental group
- r(nratio) : ratio of sample sizes, N2/N1
- r(nratio_a) : actual ratio of sample sizes
- r(nfractional) : 1 if nfractional is specified, 0 otherwise
- r(onesided) : 1 for a one-sided test, 0 otherwise
- r(m1) : control-group mean
- r(m2) : experimental-group mean
- r(diff) : difference between the experimental- and control-group means
- r(sd) : common standard deviation of the control and experimental groups
- r(sdi) : standard deviation of the control group
- r(sd2) : standard deviation of the experimental group
- r(knownsds) : 1 if option knownsds is specified, 0 otherwise
- r(separator) : number of lines between separator lines in the table
- r(divider) : 1 if divider is requested in the table, 0 otherwise
- r(init) : initial value for sample sizes or experimental-group mean
- r(maxiter) : maximum number of iterations
- r(iter) : number of iterations performed
- r(tolerance) : requested parameter tolerance
- r(deltax) : final parameter tolerance achieved
- r(ftolerance) : requested distance of the objective function from zero
- r(function) : final distance of the objective function from zero
- r(converged) : 1 if iteration algorithm converged, 0 otherwise

Macros
- r(type) : test
- r(method) : twomeans
- r(direction) : upper or lower
- r(columns) : displayed table columns
- r(labels) : table column labels
- r(widths) : table column widths
- r(formats) : table column formats

Matrices
- r(pss_table) : table of results

Methods and formulas

Consider two independent samples with $n_1$ subjects in the control group and $n_2$ subjects in the experimental group. Let $x_{11}, \ldots, x_{1n_1}$ be a random sample of size $n_1$ from a normal population with mean $\mu_1$ and variance $\sigma_1^2$. Let $x_{21}, \ldots, x_{2n_2}$ be a random sample of size $n_2$ from a normal population with mean $\mu_2$ and variance $\sigma_2^2$. Let effect size $\delta$ be the difference between the experimental-group mean and the control-group mean, $\delta = \mu_2 - \mu_1$. The sample means and variances for the two independent samples are
\[ \bar{x}_1 = \frac{1}{n_1} \sum_{i=1}^{n_1} x_{1i} \quad \text{and} \quad s_1^2 = \frac{1}{n_1 - 1} \sum_{i=1}^{n_1} (x_{1i} - \bar{x}_1)^2 \]

\[ \bar{x}_2 = \frac{1}{n_2} \sum_{i=1}^{n_2} x_{2i} \quad \text{and} \quad s_2^2 = \frac{1}{n_2 - 1} \sum_{i=1}^{n_2} (x_{2i} - \bar{x}_2)^2 \]

where \( \bar{x}_j \) and \( s_j^2 \) are the respective sample means and sample variances of the two samples.

A two-sample means test involves testing the null hypothesis \( H_0: \mu_2 = \mu_1 \) versus the two-sided alternative hypothesis \( H_a: \mu_2 \neq \mu_1 \), the upper one-sided alternative \( H_a: \mu_2 > \mu_1 \), or the lower one-sided alternative \( H_a: \mu_2 < \mu_1 \).

The two-sample means test can be performed under four different assumptions: 1) population variances are known and not equal; 2) population variances are known and equal; 3) population variances are unknown and not equal; and 4) population variances are unknown and equal.

Let \( \sigma_D \) denote the standard deviation of the difference between the two sample means. The test statistic of the form

\[ TS = \frac{(\bar{x}_2 - \bar{x}_1) - (\mu_2 - \mu_1)}{\sigma_D} \]  

(1)

is used in each of the four cases described above. Each case, however, determines the functional form of \( \sigma_D \) and the sampling distribution of the test statistic (1) under the null hypothesis.

Let \( R = n_2 / n_1 \) denote the allocation ratio. Then \( n_2 = R \times n_1 \) and power can be viewed as a function of \( n_1 \). Therefore, for sample-size determination, the control-group sample size \( n_1 \) is computed first. The experimental-group size \( n_2 \) is then computed as \( R \times n_1 \), and the total sample size is computed as \( n = n_1 + n_2 \). By default, sample sizes are rounded to integer values; see Fractional sample sizes in [PSS-4] Unbalanced designs for details.

The following formulas are based on Armitage, Berry, and Matthews (2002); Chow et al. (2018); and Dixon and Massey (1983).

Methods and formulas are presented under the following headings:

- Known standard deviations
- Unknown standard deviations
- Unequal standard deviations
- Equal standard deviations

**Known standard deviations**

Below we present formulas for the computations that assume unequal standard deviations. When standard deviations are equal, the corresponding formulas are special cases of the formulas below with \( \sigma_1 = \sigma_2 = \sigma \).

When the standard deviations of the control and the experimental groups are known, the test statistic in (1) is a \( z \) test statistic

\[ z = \frac{(\bar{x}_2 - \bar{x}_1) - (\mu_2 - \mu_1)}{\sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}} \]

with \( \sigma_D = \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}} \). The sampling distribution of this test statistic under the null hypothesis is standard normal. The corresponding test is referred to as a \( z \) test.
Let \( \alpha \) be the significance level, \( \beta \) be the probability of a type II error, and \( z_{1-\alpha} \) and \( z_\beta \) be the \((1-\alpha)\)th and the \(\beta\)th quantiles of a standard normal distribution.

The power \( \pi = 1-\beta \) is computed using

\[
\pi = \begin{cases} 
\Phi \left( \frac{\delta}{\sigma_D} - z_{1-\alpha} \right) & \text{for an upper one-sided test} \\
\Phi \left( -\frac{\delta}{\sigma_D} - z_{1-\alpha} \right) & \text{for a lower one-sided test} \\
\Phi \left( \frac{\delta}{\sigma_D} - z_{1-\alpha}/2 \right) + \Phi \left( -\frac{\delta}{\sigma_D} - z_{1-\alpha}/2 \right) & \text{for a two-sided test}
\end{cases}
\]  

(2)

where \( \Phi(\cdot) \) is the cdf of a standard normal distribution.

For a one-sided test, the control-group sample size \( n_1 \) is computed as follows:

\[
n_1 = \left( \frac{z_{1-\alpha} - z_\beta}{\mu_2 - \mu_1} \right)^2 \left( \frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2} \right) 
\]  

(3)

For a one-sided test, if one of the group sizes is known, the other one is computed using the following formula. For example, to compute \( n_1 \) given \( n_2 \), we use the following formula:

\[
n_1 = \frac{\sigma_1^2}{\left( \frac{\mu_2 - \mu_1}{z_{1-\alpha} - z_\beta} \right)^2 - \frac{\sigma_2^2}{n_2}} 
\]  

(4)

For a two-sided test, sample sizes are computed by iteratively solving the two-sided power equation in (2). The default initial values for the iterative procedure are calculated from the respective equations (3) and (4), with \( \alpha \) replaced with \( \alpha/2 \).

The absolute value of the effect size for a one-sided test is obtained by inverting the corresponding one-sided power equation in (2):

\[
|\delta| = \sigma_D (z_{1-\alpha} - z_\beta)
\]

Note that the magnitude of the effect size is the same regardless of the direction of the test.

The experimental-group mean for a one-sided test is then computed as

\[
\mu_2 = \begin{cases} 
\mu_1 + (z_{1-\alpha} - z_\beta) \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}} & \text{when } \mu_2 > \mu_1 \\
\mu_1 - (z_{1-\alpha} - z_\beta) \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}} & \text{when } \mu_2 < \mu_1 
\end{cases}
\]

For a two-sided test, the experimental-group mean is computed by iteratively solving the two-sided power equation in (2) for \( \mu_2 \). The default initial value is obtained from the corresponding one-sided computation with \( \alpha/2 \).

**Unknown standard deviations**

When the standard deviations of the control group and the experimental group are unknown, the test statistic in (1) is a \( t \) test statistic

\[
t = \frac{(\bar{x}_2 - \bar{x}_1) - (\mu_2 - \mu_1)}{s_D}
\]
where $s_D$ is the estimated standard deviation of the sample mean difference. The sampling distribution of this test statistic under the null hypothesis is (approximately) a Student’s $t$ distribution with $\nu$ degrees of freedom. Parameters $\nu$ and $s_D$ are defined below, separately for the case of equal and unequal standard deviations.

Let $t_{\nu,\alpha}$ denote the $\alpha$th quantile of a Student’s $t$ distribution with $\nu$ degrees of freedom. Under the alternative hypothesis, the test statistic follows a noncentral Student’s $t$ distribution with $\nu$ degrees of freedom and noncentrality parameter $\lambda$.

The power is computed from the following equations:

$$
\pi = \begin{cases} 
1 - T_{\nu,\lambda} (t_{\nu,1-\alpha}) & \text{for an upper one-sided test} \\
T_{\nu,\lambda} (-t_{\nu,1-\alpha}) & \text{for a lower one-sided test} \\
1 - T_{\nu,\lambda} (t_{\nu,1-\alpha}/2) + T_{\nu,\lambda} (-t_{\nu,1-\alpha}/2) & \text{for a two-sided test}
\end{cases} \tag{5}
$$

In the equations above, $\lambda = |\mu_2 - \mu_1|/s_D$.

Sample sizes and the experimental-group mean are obtained by iteratively solving the nonlinear equation (5) for $n_1$, $n_2$, and $\mu_2$, respectively. For sample-size and effect-size computations, the default initial values for the iterative procedure are calculated using the corresponding formulas assuming known standard deviations from the previous subsection.

**Unequal standard deviations**

In the case of unequal standard deviations,

$$s_D = \sqrt{s_1^2/n_1 + s_2^2/n_2}$$

and the degrees of freedom $\nu$ of the test statistic is obtained by Satterthwaite’s formula:

$$\nu = \frac{\left(\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}\right)^2}{\frac{(s_1^2/n_1)^2}{n_1-1} + \frac{(s_2^2/n_2)^2}{n_2-1}}$$

The sampling distribution of the test statistic under the null hypothesis is an approximate Student’s $t$ distribution. We refer to the corresponding test as Satterthwaite’s $t$ test.

**Equal standard deviations**

In the case of equal standard deviations,

$$s_D = s_p \sqrt{1/n_1 + 1/n_2}$$

where $s_p = \left\{ \sum_{i=1}^{n_1} (x_{1i} - \bar{x}_1)^2 + \sum_{i=1}^{n_2} (x_{2i} - \bar{x}_2)^2 \right\} / (n_1 + n_2 - 2)$ is the pooled-sample standard deviation.

The degrees of freedom $\nu$ is

$$\nu = n_1 + n_2 - 2$$

The sampling distribution of the test statistic under the null hypothesis is exactly a Student’s $t$ distribution. We refer to the corresponding test as a two-sample $t$ test.
References


Also see

[PSS-2] power twomeans, cluster — Power analysis for a two-sample means test, CRD

[PSS-2] power — Power and sample-size analysis for hypothesis tests

[PSS-2] power oneway — Power analysis for one-way analysis of variance

[PSS-2] power twoway — Power analysis for two-way analysis of variance

[PSS-2] power, graph — Graph results from the power command

[PSS-2] power, table — Produce table of results from the power command

[PSS-3] ciwidth twomeans — Precision analysis for a two-means-difference CI

[PSS-5] Glossary

[R] ttest — t tests (mean-comparison tests)