

# Package ‘anscombiser’

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**Title** Create Datasets with Identical Summary Statistics

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**Description** Anscombe's quartet are a set of four two-variable datasets that have several common summary statistics but which have very different joint distributions. This becomes apparent when the data are plotted, which illustrates the importance of using graphical displays in Statistics. This package enables the creation of datasets that have identical marginal sample means and sample variances, sample correlation, least squares regression coefficients and coefficient of determination. The user supplies an initial dataset, which is shifted, scaled and rotated in order to achieve target summary statistics. The general shape of the initial dataset is retained. The target statistics can be supplied directly or calculated based on a user-supplied dataset. The 'datasauRus' package <https://cran.r-project.org/package=datasauRus> provides further examples of datasets that have markedly different scatter plots but share many sample summary statistics.

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**License** GPL (>= 2)

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<https://github.com/paulnorthrop/anscombiser>

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anscombiser-package     *anscombiser: Create Datasets with Identical Summary Statistics*

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## Description

Anscombe's quartet (Anscombe, 1973) are a set of four two-variable datasets that have several common summary statistics but which have very different joint distributions. This becomes apparent when the data are plotted, which illustrates the importance of using graphical displays in Statistics. This package enables the creation of datasets that have identical marginal sample means and sample variances, sample correlation, least squares regression coefficients and coefficient of determination. The user supplies an initial dataset, which is shifted, scaled and rotated in order to achieve target summary statistics. The general shape of the initial dataset is retained. The target statistics can be supplied directly or calculated based on a user-supplied dataset.

## Details

The main functions in `anscombiser` are

- `anscombise`, which modifies a user-supplied dataset so that it shares sample summary statistics with Anscombe's quartet.
- `mimic`, which modified a user-supplied dataset so that is shares sample summary statistics with another user-supplied dataset.

See `vignette("intro-to-anscombiser", package = "anscombiser")` for an overview of the package.

**Author(s)**

**Maintainer:** Paul J. Northrop <p.northrop@ucl.ac.uk> [copyright holder]

**References**

Anscombe, F. J. (1973). Graphs in Statistical Analysis. *The American Statistician* 27 (1): 17–21.  
[doi:10.1080/00031305.1973.10478966](https://doi.org/10.1080/00031305.1973.10478966)

**See Also**

[anscombise](#) and [mimic](#)

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anscombe

*Anscombe's Quartet Separated*

---

**Description**

Provides Anscombe's Quartet as separate data frames.

**Usage**

anscombe1

anscombe2

anscombe3

anscombe4

**Format**

All datasets are objects of class `data.frame` with 11 rows and 2 columns.

**Source**

Anscombe's Quartet of 'Identical' Simple Linear Regressions: `datasets::anscombe` in the `datasets` package. The  $i$ th dataset is `datasets::anscombe[, c(i, i + 4)]`.

**References**

Anscombe, Francis J. (1973). Graphs in statistical analysis. *The American Statistician*, **27**, 17-21.  
[doi:10.2307/2682899](https://doi.org/10.2307/2682899)

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anscombe

*Create new versions of Anscombe's quartet*


---

## Description

Modifies a dataset `x` so that it shares sample summary statistics with [Anscombe's quartet](#).

## Usage

```
anscombe(x, which = 1, idempotent = TRUE)
```

## Arguments

<code>x</code>	A numeric matrix or data frame. Each column contains observations on a different variable. Missing observations are not allowed.
<code>which</code>	An integer in <code>{1, 2, 3, 4}</code> . Which of Anscombe's datasets to use as the target dataset. Obviously, this makes very little difference.
<code>idempotent</code>	A logical scalar. If <code>idempotent = TRUE</code> then applying <code>anscombe</code> to one of the datasets in Anscombe's Quartet will return the dataset unchanged, apart from a change of <code>class</code> . If <code>idempotent = FALSE</code> then the returned dataset will be a rotated version of the original dataset, with the same summary statistics. See <b>Details</b> .

## Details

The input dataset `x` is modified by shifting, scaling and rotating it so that its sample mean and covariance matrix match those of the Anscombe quartet.

The rotation is based on the square root of the sample correlation matrix. If `idempotent = FALSE` then this square root is based on the Cholesky decomposition this matrix, using `chol`. If `idempotent = TRUE` the square root is based on the spectral decomposition of this matrix, using the output from `eigen`. This is a minimal rotation square root, which means that if the input data `x` already have the exactly/approximately the required summary statistics then the returned dataset is exactly/approximately the same as the target dataset.

## Value

An object of class `c("anscombe", "matrix", "array")` with `plot` and `print` methods. This returned dataset has the following summary statistics in common with Anscombe's quartet.

- The sample means of each variable.
- The sample variances of each variable.
- The sample correlation matrix.
- The estimated regression coefficients from least squares linear regressions of each variable on each other variable.

The target and new summary statistics are returned as attributes `old_stats` and `new_stats` and the chosen Anscombe's quartet dataset as an attribute `old_data`.

## See Also

[mimic](#) to modify a dataset to share sample summary statistics with another dataset.

[datasets::anscombe](#) for Anscombe's Quartet and [anscombe](#) for Anscombe's Quartet as 4 separate datasets.

[input\\_datasets](#): input1 to input8 for some input datasets of the same size as those in Anscombe's quartet.

## Examples

```
# Produce Anscombe-like datasets using input1 to input8

a1 <- anscombe(input1, idempotent = FALSE)
plot(a1)
a2 <- anscombe(input2)
plot(a2)
a3 <- anscombe(input3, idempotent = FALSE)
plot(a3)
a4 <- anscombe(input4, idempotent = FALSE)
plot(a4)
a5 <- anscombe(input5, idempotent = FALSE)
plot(a5)
a6 <- anscombe(input6)
plot(a6)
a7 <- anscombe(input7, idempotent = FALSE)
plot(a7)
a8 <- anscombe(input8, idempotent = FALSE)
plot(a8)

# Old faithful to new faithful
new_faithful <- anscombe(datasets::faithful, which = 4)
plot(new_faithful)
# Then check that the sample summary statistics are the same
plot(new_faithful, input = TRUE)

# Map of Italy
got_maps <- requireNamespace("maps", quietly = TRUE)
if (got_maps) {
  italy <- mapdata("Italy")
  new_italy <- anscombe(italy, which = 4)
  plot(new_italy)
}
```

---

anscombe\_gif

*Animation of several Anscombised datasets*

---

## Description

Create an animation to show datasets that share sample summary statistics with [Anscombe's quartet](#).

**Usage**

```
anscombe_gif(
  x,
  which = 1,
  idempotent = TRUE,
  theme_name = "classic",
  ease = "cubic-in-out",
  transition_length = 3,
  state_length = 1,
  wrap = TRUE
)
```

**Arguments**

**x** A list of input datasets. Each one must be a suitable argument `x` for [anscombe](#).

**which, idempotent** Vectors that provide the arguments of the same names to [anscombe](#) for each dataset. If necessary, `rep_len` is used to replicate these arguments so that they each have length `length(x)`.

**theme\_name** A character scalar used to set the `ggtheme`. One of "grey", "gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void" or "test".

**ease** A character scalar passed to `ease_aes` to control how the points move in transitioning from one dataset to the next.

**transition\_length, state\_length, wrap** Arguments passed to `transition_states`.

**Details**

For this function to work the packages [ggplot2](#) and [gganimate](#) must be installed.

**Value**

An object of class `c("gganim", "gg", "ggplot")` with an additional attribute `new_data` that is a data frame with 3 variables, `x`, `y` and `dataset` containing the datasets output from `anscombe`.

The returned object may be displayed using by typing its name, e.g., `anim` or saved as a GIF file using `anim_save`, e.g., `gganimate::anim_save("anscombe.gif", anim)`.

**See Also**

[anscombe](#) modifies a dataset so that it shares sample summary statistics with [Anscombe's quartet](#).

[input\\_datasets](#): `input1` to `input8` for some input datasets of the same size as those in [Anscombe's quartet](#).

**Examples**

```
# Animate some Anscombe-like datasets produced using input1 to input8
x <- list(input1, input2, input3, input4, input5, input6, input7, input8)
idem <- c(FALSE, TRUE, FALSE, FALSE, FALSE, TRUE, FALSE, FALSE)
```

```
anim <- anscombe_gif(x, idempotent = idem)
```

---

**get\_stats***Calculate Anscombe's summary statistics*

---

**Description**

Calculates a particular set of summary statistics for a dataset.

**Usage**

```
get_stats(x)
```

**Arguments**

**x** a numeric matrix or data frame with at least 2 columns/variables. Each column contains observations on a different variable. Missing observations are not allowed.

**Value**

A named list of summary statistics containing

- **n** The sample size.
- **means** The sample means of each variable.
- **variances** The sample means of each variable.
- **correlation** The sample correlation matrix.
- **intercepts,slopes,rsquared** Matrices whose (i,j)th entries are the estimated regression coefficients in a regression of  $x[, i]$  on  $x[, j]$  and the resulting coefficient of determination  $R^2$ .

**Examples**

```
get_stats(anscombe[, c(1, 5)])
```

---

input_datasets	<i>Input datasets for use by <code>anscombe()</code></i>
----------------	--

---

### Description

Provides input datasets from which `anscombe` will produce transformed datasets that behave like [Anscombe's quartet](#) of datasets, that is, with the same traditional statistical properties but different general behaviours. Use `plot(input1)`, for example, to see the behaviours of the datasets.

### Usage

input1

input2

input3

input4

input5

input6

input7

input8

### Format

All datasets are objects of class `matrix` (inherits from `array`) with 11 rows and 2 columns.

### Source

None. Created for use in `'anscombiser'`.

### References

Anscombe, Francis J. (1973). Graphs in statistical analysis. *The American Statistician*, **27**, 17-21.  
[doi:10.2307/2682899](https://doi.org/10.2307/2682899)

---

mapdata	<i>Extract longitude and latitude values</i>
---------	--

---

### Description

Extracts longitude and latitude values for a particular region from the world map supplied by the maps package.

### Usage

```
mapdata(region = ".", map = "world", exact = FALSE, ...)
```

### Arguments

region	Passed to <a href="#">map</a> as the argument regions.
map	Passed to <a href="#">map</a> as the argument database
exact	The argument exact passed to the <a href="#">map</a> function.
...	Additional arguments to be passed to <a href="#">map</a> .

### Value

A dataframe with two columns: long and lat for longitude and latitude.

### Examples

See the examples in [mimic](#).

---

mimic	<i>Modify a dataset to mimic another dataset</i>
-------	--

---

### Description

Modifies a dataset x so that it shares sample summary statistics with a target dataset x2.

### Usage

```
mimic(x, x2, idempotent = TRUE, ...)
```

## Arguments

<code>x, x2</code>	Numeric matrices or data frames. Each column contains observations on a different variable. Missing observations are not allowed. <code>get_stats(x2)</code> sets the target summary statistics. If <code>x2</code> is missing then <code>set_stats</code> is called with <code>d = ncol(x)</code> and any additional arguments supplied via <code>...</code> . This can be used to set target summary statistics (means, variances and/or correlations).
<code>idempotent</code>	A logical scalar. If <code>idempotent = TRUE</code> then <code>mimic(x, x)</code> returns <code>x</code> , apart from a change of <code>class</code> . If <code>idempotent = FALSE</code> then the returned dataset may be a rotated version of the original dataset, with the same summary statistics. See <b>Details</b> .
<code>...</code>	Additional arguments to be passed to <code>set_stats</code> .

## Details

The input dataset `x` is modified by shifting, scaling and rotating it so that its sample mean and covariance matrix match those of the target dataset `x2`.

The rotation is based on the square root of the sample correlation matrix. If `idempotent = FALSE` then this square root is based on the Cholesky decomposition this matrix, using `chol`. If `idempotent = TRUE` the square root is based on the spectral decomposition of this matrix, using the output from `eigen`. This is a minimal rotation square root, which means that if the input data `x` already have the exactly/approximately the required summary statistics then the returned dataset is exactly/approximately the same as the target dataset `x2`.

## Value

An object of class `c("anscombe", "matrix", "array")` with `plot` and `print` methods. This returned dataset has the following summary statistics in common with `x2`.

- The sample means of each variable.
- The sample variances of each variable.
- The sample correlation matrix.
- The estimated regression coefficients from least squares linear regressions of each variable on each other variable.

The target and new summary statistics are returned as attributes `old_stats` and `new_stats`. If `x2` is supplied then it is returned as a attribute `old_data`.

## See Also

`anscombi` modifies a dataset so that it shares sample summary statistics with [Anscombe's quartet](#).

## Examples

```
### 2D examples

# The UK and a dinosaur
got_maps <- requireNamespace("maps", quietly = TRUE)
got_datasauRus <- requireNamespace("datasauRus", quietly = TRUE)
```

```
if (got_maps && got_datasauRus) {
  library(maps)
  library(datasauRus)
  dino <- datasaurus_dozen_wide[, c("dino_x", "dino_y")]
  UK <- mapdata("UK")
  new_UK <- mimic(UK, dino)
  plot(new_UK)
}

# Trump and a dinosaur
if (got_datasauRus) {
  library(datasauRus)
  dino <- datasaurus_dozen_wide[, c("dino_x", "dino_y")]
  new_dino <- mimic(dino, trump)
  plot(new_dino)
}

## Examples of passing summary statistics

# The default is zero mean, unit variance and no correlation
new_faithful <- mimic(faithful)
plot(new_faithful)

# Change the correlation
mat <- matrix(c(1, -0.9, -0.9, 1), 2, 2)
new_faithful <- mimic(faithful, correlation = mat)
plot(new_faithful)

### A 3D example

new_randu <- mimic(datasets::randu, datasets::trees)
# The samples summary statistics are equal
get_stats(new_randu)
get_stats(datasets::trees)
```

---

mimic\_gif

*Animation of several mimicking datasets*

---

## Description

Create an animation to show datasets that mimic a target dataset x2.

## Usage

```
mimic_gif(
  x,
  x2,
  idempotent = TRUE,
  theme_name = "classic",
  ease = "cubic-in-out",
```

```

  transition_length = 3,
  state_length = 1,
  wrap = TRUE
)

```

### Arguments

<code>x</code>	A list of input datasets. Each one must be suitable argument <code>x</code> for <code>mimic</code> .
<code>x2</code>	A suitable argument <code>x2</code> for <code>mimic</code> .
<code>idempotent</code>	A logical vector that provides the argument of the same names to <code>mimic</code> for each dataset. If necessary, <code>rep_len</code> is used to replicate this argument so that it has length <code>length(x)</code> .
<code>theme_name</code>	A character scalar used to set the <code>ggtheme</code> . One of "grey", "gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void" or "test".
<code>ease</code>	A character scalar passed to <code>ease_aes</code> to control how the points move in transitioning from one dataset to the next.
<code>transition_length</code> , <code>state_length</code> , <code>wrap</code>	Arguments passed to <code>transition_states</code> .

### Details

For this function to work the packages `ggplot2` and `gganimate` must be installed.

### Value

An object of class `c("gganim", "gg", "ggplot")` with an additional attribute `new_data` that is a data frame with 3 variables, `x`, `y` and `dataset` containing the datasets output from `mimic`.

The returned object may be displayed using by typing its name, e.g., `anim` or saved as a GIF file using `anim_save`, e.g., `gganimate::anim_save("anscombe.gif", anim)`.

### See Also

`mimic` to modify a dataset to share sample summary statistics with another dataset.

`input_datasets`: `input1` to `input8` for some input datasets of the same size as those in Anscombe's quartet.

### Examples

```

# Create 8 datasets that mimic Anscombe's first dataset
x <- list(input1, input2, input3, input4, input5, input6, input7, input8)
anim <- mimic_gif(x, anscombe1)

```

---

plot.anscombe                      *Plot method for objects of class "anscombe"*

---

### Description

plot method for objects inheriting from class "anscombe".

### Usage

```
## S3 method for class 'anscombe'  
plot(x, input = FALSE, stats = TRUE, digits = 3, legend_args = list(), ...)
```

### Arguments

x	an object of class 'anscombe', a result of a call to <a href="#">anscombe</a> or <a href="#">mimic</a> .
input	A logical scalar. Should the old, input data, that is, the Anscombe's dataset chosen for <a href="#">anscombe</a> or the argument x2 to <a href="#">mimic</a> , be plotted? If old = FALSE then the new, output data are plotted. If old = TRUE then the old data are plotted.
stats	A logical scalar. Should the sample summary statistics n, means, variances and correlation be added to the plot?
digits	An integer. The argument digits passed to <a href="#">signif</a> to round the values of the statistics before adding them to the plot.
legend_args	A list of arguments to be passed to <a href="#">legend</a> when stats = TRUE, especially legend_args\$x to control the position of the legend.
...	Further arguments to be passed to <a href="#">plot</a>

### Details

This function is only applicable in 2 dimensions, that is, when `length(attr(x, "new_stats")$means) = 2`.

### Value

Nothing is returned.

### Examples

See the examples in [anscombe](#) and [mimic](#).

### See Also

[anscombe](#) and [mimic](#).

---

```
print.anscombe          Print method for objects of class "anscombe"
```

---

**Description**

print method for class "anscombe".

**Usage**

```
## S3 method for class 'anscombe'
print(x, ...)
```

**Arguments**

x                    an object of class "anscombe", a result of a call to [anscombise](#) or [mimic](#).  
 ...                  Additional optional arguments to be passed to [print](#).

**Details**

Just extracts the new dataset from x and prints it using [print](#).

**Value**

The argument x, invisibly.

**See Also**

[anscombise](#) and [mimic](#)

---

```
set_stats                Create a list of summary statistics
```

---

**Description**

Creates a list of summary statistics to pass to [mimic](#).

**Usage**

```
set_stats(d = 2, means = 0, variances = 1, correlation = diag(2))
```

**Arguments**

d                    An integer that is no smaller than 2.  
 means                A numeric vector of sample means.  
 variances            A numeric vector of positive sample variances.  
 correlation          A numeric correlation matrix. None of the off-diagonal entries in correlation are allowed to be equal to 1 in absolute value.

**Details**

The vectors means and variances are recycled using `rep_len` to have length `d`.

**Value**

A list containing the following components.

- means a d-vector of sample means.
- variances a d-vector sample variances.
- correlation a d by d correlation matrix.

**Examples**

```
# Uncorrelated with zero means and unit variances
set_stats()
# Sample correlation = 0.9
set_stats(correlation = matrix(c(1, 0.9, 0.9, 1), 2, 2))
```

---

trump	<i>Donald Trump</i>
-------	---------------------

---

**Description**

A dataset that provides an image of Donald Trump's face.

**Usage**

```
trump
```

**Format**

A matrix with 4885 rows and 2 columns: x and y.

**Source**

This image was created by Accentaur from the Noun Project. <https://thenounproject.com/term/donald-trump/727774/>

**Examples**

```
plot(trump)
```

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