

# Package ‘DemographicTable’

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**Type** Package

**Title** Creating Demographic Table

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**Description** Functions for creating demographic table with simple summary statistics, with optional comparison(s) over one or more groups. Numeric variables are summarized in means, standard deviations, medians, inter-quartile-ranges (IQR), skewness, Shapiro-Wilk normality test and ranges, and compared using two-sample t-test, Wilcoxon test, ANOVA and/or Kruskal-Wallis test. Logical and factor variables are summarized in counts and percentages and compared using chi-squared test and/or Fisher's exact test.

**License** GPL-2

**Encoding** UTF-8

**Imports** flextable, e1071, xtable

**Language** en-US

**Depends** R (>= 4.3.0)

**Suggests** officer, MASS

**RoxygenNote** 7.3.1

**NeedsCompilation** no

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**as\_flextab**.DemographicTable  
*Convert DemographicTable to flextab*

### Description

Convert a **DemographicTable** to **flextab** object.

### Usage

```
## S3 method for class 'DemographicTable'
as_flextab(x, ...)
```

### Arguments

x	a <b>DemographicTable</b> object
...	potential additional parameters, not currently in use

### Value

Function **as\_flextab**.DemographicTable returns a **flextab** object.

### Note

End user may use **set\_caption** to add a caption to the output demographic table.

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**class1List***First Class of Each Element in a Recursive Object*

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**Description**

First **class** of each element in a **recursive** object

**Usage**

```
class1List(x)
```

**Arguments**

x	a <b>recursive</b> object, e.g., <b>data.frame</b> or <b>list</b>
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**Value**

Function **class1List** returns a **list** of **character** elements. Each element is a collection of the names of the elements of each specific class.

**Examples**

```
class1List(esoph)
class1List(lm(Ozone ~ Wind + Temp, data = airquality))
```

---

**DemographicTable***Create Demographic Table*

---

**Description**

Create a demographic table with simple summary statistics, with optional comparison(s) over one or more groups.

**Usage**

```
DemographicTable(data, ...)

## S3 method for class 'data.frame'
DemographicTable(
  data,
  data.name = substitute(data),
  groups = NULL,
  keep_missing_group = TRUE,
  exclude = NULL,
  exclude_pattern,
```

```

    include,
    include_pattern,
    paired = FALSE,
    robust = TRUE,
    overall = TRUE,
    compare = TRUE,
    pairwise = 3L,
    ...
)

```

## Arguments

data	a <code>data.frame</code>
...	additional parameters, currently not in use
data.name	<code>character</code> scalar, or the argument call of <code>data</code> . A user-friendly name of the input <code>data</code> .
groups	<code>character</code> scalar or <code>vector</code> , the name(s) of sub-group(s) for which the summary statistics are to be provided. Default <code>NULL</code> indicating no sub-groups.
keep_missing_group	<code>logical</code> scalar. If <code>TRUE</code> (default), the subjects with missing group are put into a new group (' <code>.missing</code> '). If <code>FALSE</code> , these subjects are removed from group-wise summary statistics.
exclude	<code>character vector</code> , the name(s) of variable(s) to be excluded. Default <code>NULL</code> indicating no variable are to be excluded.
exclude_pattern	(optional) <code>character</code> scalar as <code>regex</code> (regular expression), the pattern of the names of the variable(s) to be excluded.
include	<code>character vector</code> , the name(s) of variable(s) to be included. Default <code>names(data)</code> indicating all variables are to be included.
include_pattern	(optional) <code>character</code> scalar as <code>regex</code> (regular expression), the pattern of the names of the variable(s) to be included.
paired	<code>logical</code> scalar, whether to perform paired test (default <code>FALSE</code> )
robust	<code>logical</code> scalar. If <code>TRUE</code> (default), use non-parametric methods for non-normally distributed <code>numeric</code> variables.
overall	<code>logical</code> scalar. If <code>TRUE</code> (default), a column of overall summary statistics will be provided.
compare	<code>logical</code> scalar. If <code>TRUE</code> (default), comparisons between group(s) will be made.
pairwise	<code>integer</code> scalar, minimum number of groups where pairwise comparisons need to be performed. Default <code>3L</code> .

## Details

A demographic table with simple summary statistics, with optional comparison(s) over one or more groups, is created.

`numeric` variables are summarized in means, standard deviations, medians, inter-quartile-ranges (IQR), skewness, *p*-value of Shapiro-Wilk normality test and ranges. If group is specified, they are compared using two-sample `t.test`, `wilcox.test` (Wilcoxon / Mann-Whitney), one-way `aov` (ANOVA) and/or `kruskal.test` (Kruskal-Wallis).

`logical` and `factor` variables are summarized in counts and percentages. If group is specified, they are compared using `prop.test` (chi-squared) and/or `fisher.test` (Fisher's exact).

## Value

Function `DemographicTable` returns an object of S3 class 'DemographicTable', which inherits from `matrix`.

## Examples

```
DemographicTable(esoph)
DemographicTable(ToothGrowth, groups = 'supp', include = 'len')
DemographicTable(ToothGrowth, groups = 'supp', include = 'len', paired = TRUE)
DemographicTable(ToothGrowth, groups = 'supp', include = 'len', compare = FALSE)
DemographicTable(warpbreaks, groups = c('wool', 'tension'))
DemographicTable(mtcars, groups = c('vs', 'am'), include = c('mpg', 'cyl', 'disp'))

# with missing value
DemographicTable(airquality, groups = 'Month', exclude = 'Day')
DemographicTable(MASS::survey, groups = 'Smoke', keep_missing_group = FALSE)
DemographicTable(MASS::survey, groups = 'Smoke', keep_missing_group = FALSE, useNA = 'always')

# write to Word file
library(flextable)
library(officer)
x = read_docx() |> body_add_flextable(value = as_flextable(DemographicTable(esoph)))
(out = file.path(tempdir(), 'demotable.docx'))
print(x, target = out)
# system(paste('open', out)) # works on Mac & Windows, but requires Microsoft Word
file.remove(out)
```

## Description

Obtain *p*-value from `shapiro.test`, taking into consideration of several exceptions.

## Usage

```
pval_shapiro(x, CLT = FALSE, ...)
```

## Arguments

x	<b>double</b> vector
CLT	<b>logical</b> scalar, whether to allow the use of Central Limit Theorem, default FALSE
...	additional parameters, currently not in use

## Details

Function **pval\_shapiro** provides a pseudo *p*-value for several exceptions of **shapiro.test**, serving as a criteria of whether robust statistics/tests need to be used

`length(x) < 3L` returns *p* = 0, indicating that robust methods are needed.

`length(x) > 5e3L` returns *p* = 1, indicating that no robust method is needed. For such large sample size, robust methods could be too slow.

`CLT & length(x) > 30L` returns *p* = 1, indicating that no robust method is needed because of the use of Central Limit Theorem.

`all x values are identical (or all.equal, to be implemented in future release)` returns *p* = 0, indicating that robust methods are needed.

**Otherwise** use the *p*-value from **shapiro.test**.

## Value

Function **pval\_shapiro** returns a **double** scalar.

## Examples

```
pval_shapiro(rnorm(5))
sapply(with(airquality, split(Ozone, f = Month)), FUN = pval_shapiro)
```

summaryText

*Summary Text*

## Description

Provide the summary text of an R object

## Usage

```
summaryText(x, fmt, ...)
## Default S3 method:
summaryText(x, fmt = "%.2f", ...)
## S3 method for class 'factor'
summaryText(x, fmt = "%.1f", useNA = c("no", "always"), ...)
```

```
## S3 method for class 'ordered'
summaryText(x, fmt = "%.1f", useNA = c("no", "always"), ...)

## S3 method for class 'character'
summaryText(x, ...)

## S3 method for class 'logical'
summaryText(x, fmt = "%.1f", ...)
```

## Arguments

x	an R object
fmt	<code>character</code> scalar, format string, see <a href="#">sprintf</a>
...	additional parameters, currently not in use
useNA	<code>character</code> scalar, 'no' (default) or 'always', see <a href="#">table</a>

## Value

Function `summaryText` returns a `character` scalar.

## Examples

```
x = rpois(n = 20L, lambda = 2)
x[sample.int(length(x), 3L)] = NA_integer_
summaryText(x)

# factor
x = state.region
x[2L] = NA_integer_
summaryText(x)

# binary
summaryText(c(TRUE, FALSE, TRUE, NA))
summaryText(c(TRUE, FALSE, TRUE))
summaryText(c(FALSE, FALSE, NA))
summaryText(c(FALSE, FALSE, FALSE))
summaryText(c(NA, NA, NA))
```

## Description

Write `DemographicTable` to LaTeX.

**Usage**

```
## S3 method for class 'DemographicTable'  
xtable(x, ...)
```

**Arguments**

x	a <a href="#">DemographicTable</a> object
...	additional parameters of <a href="#">xtable</a>

**Value**

Function [xtable.DemographicTable](#) returns an [xtable](#) object.

**Examples**

```
(tb = DemographicTable(ToothGrowth, groups = 'supp'))  
library(xtable)  
print(xtable(tb), sanitize.text.function = identity,  
      sanitize.colnames.function = NULL, include.rownames = FALSE)
```

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