

# Package ‘relimp’

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**Title** Relative Contribution of Effects in a Regression Model

**Author** David Firth with contributions from Heather Turner and John Fox

**Maintainer** David Firth <d.firth@warwick.ac.uk>

**URL** <http://warwick.ac.uk/relimp>

**Description** Functions to facilitate inference on the relative importance of predictors in a linear or generalized linear model, and a couple of useful Tcl/Tk widgets.

**Depends** R (>= 2.0.0)

**Suggests** tcltk, nnet, MASS, Rcmdr

**Imports** stats, utils

**License** GPL (>= 2)

**NeedsCompilation** no

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pickFrom

*Pick Subsets from a Vector*


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### Description

Provides a Tk dialog or a text-based menu for interactive selection of one or more subsets from a vector.

### Usage

```
pickFrom(vec, nsets = 1, return.indices = FALSE,
         setlabels = NULL, edit.setlabels = TRUE,
         subset = TRUE,
         warningText = "one or more selections empty",
         title = "Subset picker",
         items.label = "Pick from",
         labels.prompt = "Your label for this set",
         list.height = 20,
         items.scrollbar = TRUE,
         preserve.order = TRUE,
         graphics = TRUE,
         listFont = "Courier 12",
         labelFont = "Helvetica 11",
         windowPos = "+150+30")
```

### Arguments

<code>vec</code>	a vector
<code>nsets</code>	a positive integer, the number of subsets to be selected
<code>return.indices</code>	logical, whether indices (TRUE) or vector contents (FALSE) are to be returned
<code>setlabels</code>	a character vector of labels for the subsets
<code>edit.setlabels</code>	logical, determines whether a textbox is provided for editing the label of each subset
<code>subset</code>	logical, character or numeric vector indicating which elements of <code>vec</code> should be made available for selection. Default is to make all elements available.
<code>warningText</code>	character, text to use as a warning in situations where no selection is made into one or more of the specified sets
<code>title</code>	character, title of the Tk dialog window
<code>items.label</code>	character, a label for the set of items to be selected from
<code>labels.prompt</code>	character, a prompt for textual set label(s)
<code>list.height</code>	maximum number of elements of <code>vec</code> to display at once
<code>items.scrollbar</code>	logical, whether a scrollbar is to be provided when <code>vec</code> is longer than <code>list.height</code>

preserve.order	logical: should the order of items in vec be maintained in all of the returned subsets?
graphics	logical: should a dialog be used, if possible?
listFont	a Tk font specification for the items list and subsets
labelFont	a Tk font specification for the labels entrybox
windowPos	position of the Tk dialog, in pixels from top left of display

### Details

If graphics = TRUE and the tcltk package is operational, a Tk dialog is used, otherwise a text menu.

If return.indices is used together with subset, the indices returned relate to vec, not to vec[subset].

### Value

EITHER (in the case of a text menu or if the dialog is ended with "OK") a list, with nsets components. Each component is a selected sub-vector, or a numeric vector of indices for a selected sub-vector (if return.indices is TRUE). The component names are as specified in setlabels, or as specified interactively.

OR (if the dialog is ended either "Cancel" or the close-window control button is used) NULL.

### Author(s)

David Firth, with contributions from Heather Turner

### Examples

```
## These examples cannot be run by example() but should be OK when pasted
## into an interactive R session
## Not run:
pickFrom(c("apple", "banana", "plum", "grapefruit"), nsets = 2,
  preserve.order = FALSE,
  setlabels = c("Fruits I like", "Fruits I tolerate"))

## End(Not run)
## Not run:
## Type selections as e.g. 1:2, 4
pickFrom(c("apple", "banana", "plum", "grapefruit"), nsets = 2,
  preserve.order = FALSE,
  setlabels = c("Fruits I like", "Fruits I tolerate"),
  graphics = FALSE)

## End(Not run)
```

---

R.to.Tcl

*Convert a Character Vector to Tcl Format*

---

### Description

Converts a character vector into a brace-delimited Tcl list

### Usage

```
R.to.Tcl(character.vector)
```

### Arguments

```
character.vector  
    A character vector
```

### Value

A character vector of length 1

### Author(s)

David Firth, <d.firth@warwick.ac.uk>

### See Also

[Tcl.to.R](#)

### Examples

```
R.to.Tcl(c("apple", "banana"))
```

---

relimp

*Relative Importance of Predictors in a Regression Model*

---

### Description

Produces a summary of the relative importance of two predictors or two sets of predictors in a fitted model object.

### Usage

```
relimp(object, set1=NULL, set2=NULL, label1="set1", label2="set2",  
        subset=TRUE,  
        response.cat=NULL, ...)  
## S3 method for class 'relimp'  
print(x, digits=3, ...)
```

**Arguments**

object	A model object of class <code>lm</code> , <code>glm</code> , <code>coxph</code> , <code>survreg</code> , <code>multinom</code> , <code>polr</code> or <code>gls</code>
set1	An index or vector of indices for the effects to be included in the numerator of the comparison
set2	An index or vector of indices for the effects to be included in the denominator of the comparison
label1	A character string; mnemonic name for the variables in set1
label2	A character string; mnemonic name for the variables in set2
subset	Either a vector of numeric indices for the cases to be included in the standardization of effects, or a vector of logicals (TRUE for inclusion) whose length is the same as the number of rows in the model frame, <code>object\$model</code> . The default choice is to include all cases in the model frame.
response.cat	If object is of class <code>multinom</code> , this is a character string used to specify which regression is of interest (i.e., the regression which predicts the log odds on response cat versus the model's reference category). The <code>response.cat</code> argument should be an element of <code>object\$lab</code> ; or NULL if object is not of class <code>multinom</code> .
...	For models of class <code>glm</code> , one may additionally set the dispersion parameter for the family (for example, <code>dispersion=1.69</code> ). By default it is obtained from object. Supplying it here permits explicit allowance for over-dispersion, for example.
x	an object of class <code>relimp</code>
digits	The number of decimal places to be used in the printed summary. Default is 3.

**Details**

If `set1` and `set2` both have length 1, relative importance is measured by the ratio of the two standardized coefficients. Equivalently this is the ratio of the standard deviations of the two contributions to the linear predictor, and this provides the generalization to comparing two sets rather than just a pair of predictors.

The computed ratio is the square root of the variance-ratio quantity denoted as 'omega' in Silber, J H, Rosenbaum, P R and Ross, R N (1995). Estimated standard errors are calculated by the delta method, as described in that paper for example.

If `set1` and `set2` are unspecified, and if the `tcltk` package has been loaded, a dialog box is provided (by a call to `pickFrom`) for the choice of `set1` and `set2` from the available model coefficients.

**Value**

An object of class `relimp`, with at least the following components:

model	The call used to construct the model object summarized
sets	The two sets of indices specified as arguments
log.ratio	The natural logarithm of the ratio of effect standard deviations corresponding to the two sets specified
se.log.ratio	An estimated standard error for <code>log.ratio</code>

If dispersion was supplied as an argument, its value is stored as the dispersion component of the resultant object.

### Author(s)

David Firth <d.firth@warwick.ac.uk>

### References

Silber, J. H., Rosenbaum, P. R. and Ross, R N (1995) Comparing the Contributions of Groups of Predictors: Which Outcomes Vary with Hospital Rather than Patient Characteristics? *JASA* **90**, 7–18.

### See Also

[relreimp](#)

### Examples

```
set.seed(182) ## an arbitrary number, just for reproducibility
x <- rnorm(100)
z <- rnorm(100)
w <- rnorm(100)
y <- 3 + (2 * x) + z + w + rnorm(100)
test <- lm(y ~ x + z + w)
print(test)
relimp(test, 2, 3) # compares effects of x and z
relimp(test, 2, 3:4) # compares effect of x with that of (z,w) combined
##
## Data on housing and satisfaction, from Venables and Ripley
## -- multinomial logit model
library(MASS)
library(nnet)
data(housing)
house.mult <- multinom(Sat ~ Infl + Type + Cont, weights = Freq,
  data = housing)
relimp(house.mult, set1 = 2:3, set2 = 7, response.cat = "High")
```

---

relreimp

*Comparison of Relative Importances in a Multinomial Logit Model*

---

### Description

Produces a summary of the relative importance of two predictors or two sets of predictors in a fitted `multinom` model object, and compares relative importances across two of the fitted logit models.

### Usage

```
relreimp(object, set1=NULL, set2=NULL, label1="set1", label2="set2",
  subset=TRUE,
  response.cat1=NULL, response.cat2=NULL)
```

**Arguments**

<code>object</code>	A model object of class <code>multinom</code>
<code>set1</code>	An index or vector of indices for the effects to be included in the numerator of the comparison
<code>set2</code>	An index or vector of indices for the effects to be included in the denominator of the comparison
<code>label1</code>	A character string; mnemonic name for the variables in <code>set1</code>
<code>label2</code>	A character string; mnemonic name for the variables in <code>set2</code>
<code>subset</code>	Either a vector of numeric indices for the cases to be included in the standardization of effects, or a vector of logicals (TRUE for inclusion) whose length is the same as the number of rows in the model frame, <code>object\$model</code> . The default choice is to include all cases in the model frame.
<code>response.cat1</code>	A character string used to specify the first regression of interest (i.e., the regression which predicts the log odds on <code>response.cat1</code> versus the model's reference category). The <code>response.cat1</code> argument should be an element of <code>object\$lab</code> .
<code>response.cat2</code>	A character string used to specify the second regression of interest (i.e., the regression which predicts the log odds on <code>response.cat2</code> versus the model's reference category). The <code>response.cat2</code> argument should be an element of <code>object\$lab</code> .

**Details**

Computes a relative importance summary as described in [relimp](#), for each of the two regressions specified by `response.cat1` and `response.cat2` (relative to the same reference category); and computes the difference of those two relative importance summaries, along with an estimated standard error for that difference.

**Value**

An object of class `relreimp`, with at least the following components:

<code>model</code>	The call used to construct the model object summarized
<code>sets</code>	The two sets of indices specified as arguments
<code>response.category</code>	A character vector containing the specified <code>response.cat1</code> and <code>response.cat2</code>
<code>log.ratio</code>	The natural logarithm of the ratio of effect standard deviations corresponding to the two sets specified. A vector with three components: the first is for <code>response.cat1</code> versus the reference category, the second for <code>response.cat2</code> versus the reference category, the third is the difference.
<code>se.log.ratio</code>	Estimated standard errors for the elements of <code>log.ratio</code>

**Author(s)**

David Firth, <d.firth@warwick.ac.uk>

**See Also**[relimp](#)**Examples**

```
## Data on housing and satisfaction, from Venables and Ripley
library(MASS)
library(nnet)
data(housing)
house.mult <- multinom(Sat ~ Infl + Type + Cont, weights = Freq,
  data = housing)
relrelimp(house.mult, set1 = 2:3, set2 = 7,
  label1 = "Influence", label2 = "Contact",
  response.cat1 = "Medium", response.cat2 = "High")
## Computes the relative contribution of Influence and Contact in
## each of the two logistic regressions (Med/Low and High/Low), and
## compares those two relative-contribution measures.
```

---

`showData`*Display a Data Frame in a Tk Text Widget*

---

**Description**

Displays the contents of a data frame in a modeless Tk text window, for inspection. Objects not of class `data.frame`, for example objects of class `table`, or `matrix`, are coerced using `as.data.frame` prior to display.

**Usage**

```
showData(dataframe,
  colname.bgcolor = "grey50",
  rowname.bgcolor = "grey50",
  body.bgcolor = "white",
  colname.textcolor = "white",
  rowname.textcolor = "white",
  body.textcolor = "black",
  font = "Courier 12",
  maxheight = 30,
  maxwidth = 80,
  title = NULL,
  rowname.bar = "left",
  colname.bar = "top",
  rownumbers = FALSE,
  placement = "-20-40",
  suppress.X11.warnings = TRUE)
```

**Arguments**

dataframe	A data frame, or an object to which <code>as.data.frame()</code> can be validly applied
colname.bgcolor	A background colour for the variable-names panel
rowname.bgcolor	A background colour for the row-names panel
body.bgcolor	A background colour for the data
colname.textcolor	A colour for the variable names
rowname.textcolor	A colour for the row names
body.textcolor	A colour for the data
font	The text font used – should be a monospaced font
maxheight	The maximum number of rows to display
maxwidth	The maximum width of display, in characters
title	A title for the window. Default is to use the name of the dataframe as given in the call to <code>showData()</code>
rowname.bar	position of sidebar for row names, "left" or "right", or <code>c("left", "right")</code> , or NULL
colname.bar	position of column names, "top" or "bottom", or <code>c("top", "bottom")</code> , or NULL
rownnumbers	logical, whether row numbers should be displayed
placement	Position of the bottom right corner of the window
suppress.X11.warnings	logical, if TRUE then any X11 warnings are suppressed

**Value**

Invisibly returns the Tk window containing the displayed data frame.

**Note**

An error results if the printed representation of `dataframe` exceeds the maximum allowed width of 10000 characters; see [options](#).

Text can be copied from the Tk window to the system clipboard, using `<Control-C>` or via a right-click pop-up menu.

On some systems the window may take a few seconds to appear if the data frame is very large.

**Author(s)**

David Firth, `<d.firth@warwick.ac.uk>`; with Rcmdr-specific features contributed by John Fox

**Examples**

```
## This cannot be run by example() but should be OK when pasted
## into an interactive R session
## Not run:
data(mtcars)
showData(mtcars)
## End(Not run)
```

---

Tcl.to.R

*Convert a Tcl List to R Character Vector*

---

**Description**

Converts a brace-delimited list from Tcl into a character vector

**Usage**

```
Tcl.to.R(tcl.list)
```

**Arguments**

tcl.list            a character string

**Value**

a character vector

**Author(s)**

David Firth, <d.firth@warwick.ac.uk>

**See Also**

[R.to.Tcl](#)

**Examples**

```
Tcl.to.R("{apple} {banana} {pear}")
```

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