

Package ‘CB’

August 31, 2015

Title Cole Brokamp's Personal Functions

Version 0.1

Description Cole's personal R functions

Depends R (>= 3.1.2)

Imports parallel,
XML,
RCurl

License GPL

LazyData true

R topics documented:

CBapply	1
getPackages	2
LatLongToFIPS	3
ORGetter	3
save_pdf	4
tableSummary	4
tableTest	5
Index	6

CBapply	<i>CB's Custom Apply Function</i>
---------	-----------------------------------

Description

This function is a wrapper for `sapply` with `simplify=FALSE` and `USE.NAMES=TRUE`. It then `rbinds` via `do.call` to return `data.frame`. In order for the names to work properly, a function that returns a `data.frame` must be used (see example).

Usage

```
CBapply(X, FUN, output = "data.frame", num.cores = 1, ...)
```

Arguments

X	List of objects to apply over
FUN	Function to apply
output	Output type. Defaults to 'data.frame', but can also be set to 'list' to suppress rbinding of the list.
num.cores	Defaults to 1 and the base 'sapply' is used. If set to greater than one, then it is the number of cores used in parallel::mclapply().
...	Additional arguments to the function

Examples

```
X <- as.data.frame(matrix(runif(100),ncol=10))
names(X) <- LETTERS[1:10]
CBapply(X,mean)
# function must return a data.frame with named columns for column names to work
CBapply(X,function(x) data.frame('mean'=mean(x)))
```

getPackages

getPackages

Description

This function takes a package and returns a list of its dependencies. Good for downloading source files of packages to install on a R server where internet access is blocked.

Usage

```
getPackages(packs)
```

Arguments

packs	a quoted package name or list of package names
-------	--

Examples

```
## Not run:
# use this to get specifically named packages and their dependencies:
packages <- getPackages('pbapply')
# use this to get all packages installed on local machine and their dependencies:
# packages <- getPackages(row.names(installed.packages()))
# then download the packages:
download.packages(packages, destdir='.', type='source')

## End(Not run)
```

LatLongToFIPS

Converting Lat/Long Coords into FIPS code

Description

This function takes a latitude and longitude input numbers and returns the FIPS code by calling the Census Block Conversion API at the FCC.gov website. (See more details here: <http://www.fcc.gov/developers/census-block-conversions-api>)

Usage

```
LatLongToFIPS(latitude, longitude, census.year = "2010", showall = "false")
```

Arguments

latitude	Latitude coordinate
longitude	Longitude coordinate
census.year	Defaults to '2010'. Not tested on other years; shouldn't need to change as FIPS locations rarely change.
showall	Set to 'false' as default. Has to do with the FCC API; shouldn't need to change

Examples

```
LatLongToFIPS(latitude=39.135398, longitude=-84.519902)
```

ORGetter

Retreive Odds Ratio Table from Logistic GLM Objects

Description

This function returns a data.frame of the odds ratios and their 95

Usage

```
ORGetter(logistic.glm, digits = 2, sig.star = TRUE,  
show.intercept = FALSE)
```

Arguments

logistic.glm	A logistic GLM R object. If not an object of 'glm' and 'lm', it will stop with an error.
digits	Number of digits to round table
sig.star	Will return an extra column with a star if the confidence interval does not contain 1. Defaults to TRUE.
show.intercept	Will show the intercept and its confidence interval only if set to TRUE. Defaults to FALSE.

Examples

```
## Not run: x1 <- rnorm(100)
x2 <- rnorm(100)
y <- rbinom(100,1,prob=0.3)
logistic.model <- glm(y ~ x1 + x2,family='binomial')
ORGetter(logistic.model)

## End(Not run)
```

save_pdf

save_pdf

Description

Copies the graphics contents of current device to PDF (a wrapper for dev2.pdf). Default size is 8.5 x 11 in landscape mode.

Usage

```
save_pdf(file, width = 11, height = 8.5)
```

Arguments

file	filename to save the image
width	width of pdf image
height	height of pdf image

tableSummary

Summary Table

Description

This function summarizes numerical and dichotomous variables only. The summary number is either the mean of a numeric variable for the number and percentage of values that are the second of the two factors in a dichotomous variable. Missing values are removed before the summary statistic is calculated and the number of missing observations is also presented in the table.

Usage

```
tableSummary(x, digits.mean = 2, digits.percentage = 0)
```

Arguments

x	Vector of data which to summarize. Should be used for numerical and dichotomous variables only.
digits.mean	The mean is rounded and displayed using this many digits.
digits.percentage	The percentage is rounded and displayed using this many digits.

Examples

```
X <- data.frame('some.continuous'=runif(300), 'some.factor'=factor(rbinom(300,1,0.3)))
tableSummary(X$some.continuous)
# use CBapply to create a table
CBapply(X,tableSummary)
# specify the digits differently to change the display of the table
CBapply(X,tableSummary,digits.mean=3,digits.percentage=2)
```

tableTest	<i>Table Test</i>
-----------	-------------------

Description

This function summarizes and tests the differences of numerical and dichotomous variables only across some factor. The summary number is either the mean of a numeric variable for the number and percentage of values that are the second of the two factors in a dichotomous variable. Missing values are removed before the summary statistic, but the number missing is not reported. Furthermore, a p-value is reported testing the differences of the means or counts across the groups factor. The p-value is derived from an ANOVA for continuous variables or from a chi-squared test via monte-carlo simulation using 100,000 bootstrap replicates.

Usage

```
tableTest(x, group, digits.mean = 2, digits.percentage = 0)
```

Arguments

x	Vector of data which to summarize. Should be used for numerical and dichotomous variables only.
group	The factor for which to test the x variable across.
digits.mean	The mean is rounded and displayed using this many digits.
digits.percentage	The percentage is rounded and displayed using this many digits.

Examples

```
X <- data.frame('some.continuous'=runif(300), 'some.factor'=factor(rbinom(300,1,0.3)))
X$some.other.factor <- factor(rbinom(300,1,0.5))
tableTest(x=X$some.continuous,group=X$some.other.factor)
tableTest(x=X$some.factor,group=X$some.other.factor)
CBapply(X[,c('some.continuous','some.factor')],tableTest,group=X$some.other.factor)
```

Index

- *Topic **CBapply**
 - CBapply, [1](#)
 - getPackages, [2](#)
- *Topic **CB**
 - LatLongToFIPS, [3](#)
 - ORGetter, [3](#)
 - tableSummary, [4](#)
 - tableTest, [5](#)
- *Topic **FIPS**
 - LatLongToFIPS, [3](#)
- *Topic **OddsRatio**
 - ORGetter, [3](#)
- *Topic **summary**
 - tableSummary, [4](#)
- *Topic **table**
 - tableSummary, [4](#)
 - tableTest, [5](#)
- *Topic **test**
 - tableTest, [5](#)

CBapply, [1](#)

getPackages, [2](#)

LatLongToFIPS, [3](#)

ORGetter, [3](#)

save_pdf, [4](#)

tableSummary, [4](#)

tableTest, [5](#)