

# Package ‘CB’

November 4, 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,  
pbapply,  
plyr

**License** GPL

**LazyData** true

## R topics documented:

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| CBapply | <i>CB's Custom Apply Function</i> |
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## 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) # <- will return error
# function must return a data.frame with named columns for column names to work
CBapply(X,function(x) data.frame('mean'=mean(x)))
```

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getPackages

getPackages

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**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)
```

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LatLongToFIPS

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*Converting Lat/Long Coords into FIPS code*

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**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)
```

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ORGetter

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*Retreive Odds Ratio Table from Logistic GLM Objects*

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**Description**

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

**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)
```

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*save\_pdf*

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*save\_pdf*

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**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**

|                     |                            |
|---------------------|----------------------------|
| <code>file</code>   | filename to save the image |
| <code>width</code>  | width of pdf image         |
| <code>height</code> | height of pdf image        |

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*SPapply*

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*SPapply*

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**Description**

\*apply function for spatial point objects

**Usage**

```
SPapply(sp.object, FUN., ..., progress.bar = TRUE, id.row.names = FALSE)
```

**Arguments**

|                           |  |
|---------------------------|--|
| <code>sp.object</code>    | a <code>SpatialPoints</code> or <code>SpatialPointsDataFrame</code> object   |
| <code>FUN.</code>         | function to be applied; must take <code>sp.object</code> as first argument and must return a <code>data.frame</code> |
| <code>...</code>          | additional arguments passed to function  |
| <code>progress.bar</code> | logical, show progress bar?  |
| <code>id.row.names</code> | if TRUE, set row.names of output <code>data.frame</code> from <code>data\$id</code> of <code>sp.object</code>        |

**Value**

`data.frame` of all results from function applied to `sp.object`

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tableSummary

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*Summary Table*


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### 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 CBAppl to create a table
CBAppl(X,tableSummary)
# specify the digits differently to change the display of the table
CBAppl(X,tableSummary,digits.mean=3,digits.percentage=2)
```

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tableTest

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*Table Test*


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### 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)
```

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