Package 'csmtools'

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Type Package

Title R code in use in CSM solutions

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Description A collection of helpful functions that are widely or commonly used by the organization.

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LazyData TRUE

RoxygenNote 5.0.1

Imports data.table,magrittr

Suggests testthat

R topics documented:

csmtools-package *csmtools: A collection of useful code*

Description

csmtools is a collection of code that can be used in different projects to help prevent team members from reinventing the wheel for some things that others have already figured out.

Details

Bug reports, additions and enhancement requests are welcomed at: https://github.com/McClellandLegge/csmtools

dt_compare *Merge and compare columns of data.frames (data.tables)*

Description

Merge and compare columns of data.frames (data.tables)

Usage

```
dt_compare(x, y, compare = NULL, func = `-`, precision = 6, ...)
```

Arguments

x	A data.frame
y	A data.frame
compare	A character string or vector of shared column names
func	A binary function to compare the columns with, should be appropriate for the datatypes of the columns
precision	The precision of the comparison, is the digits argument to the round function
...	Any arguments to the merge function

Value

A data.frame

Examples

```
x <- iris[1:50,]
y <- iris[1:60,]
x$id <- seq(nrow(x))
y$id <- seq(nrow(y))
y$Sepal.Width = y$Sepal.Width + rnorm(n = nrow(y))

# can specify any arguments to 'merge'
res <- dt_compare(x, y, compare = c("Sepal.Width", "Sepal.Length"), by = "id", all.y = TRUE)
```

dt_reduce *Apply a row-wise Reduce*

Description

Apply a row-wise Reduce

Usage

```
dt_reduce(DT, FUN, ...)
```

Arguments

DT	A data.table
FUN	Any binary function
...	Quoted column names from DT

Details

Apply a row-wise reduce for a given function on a set of a data.table's columns. The main advantage of this function is that names can be passed to the function as vectors, eliminating the need to hard code differencing, etc. based on column names. Additionally, the output is specified by the user – often we want to perform a calculation and have vector output, something usaully implemented with an ugly unlist.

Value

A vector

Class will vary

Examples

```
library("data.table")
DT <- as.data.table(head(iris))
# basic differencing
dt_reduce(DT, `-`, "Sepal.Length", "Sepal.Width")

# paste columns together row-wise
dt_reduce(DT, paste, colnames(DT))

# calculate the mean
dt_reduce(DT, `+`, "Sepal.Length", "Sepal.Width", "Petal.Length") / nrow(DT)
```

file_size_filter *Filter files based on their size*

Description

Filter files based on their size

Usage

```
file_size_filter(x, size = 0, units = "B", include = FALSE)
```

Arguments

x	A character vector of filenames
size	A numeric vector, the size of the file
units	A character vector specifying the units. Options are B , KB, MB and GB. Must match the length of size if specifying more than one unit.
include	A boolean, include files of size size?

Details

Filters out the files that are less than (or less than or equal to) the size with units specified. If all files are filtered out, then a character vector of length 0 is returned.

Value

A character vector

Examples

```
x <- list.files(path = Sys.getenv("TEMP"), full.names = TRUE)[1]
file_size_filter(x, size = 1, units = "KB")
```

filter_files *Filter out file names based on criteria*

Description

Filter out file names based on criteria

Usage

```
filter_files(x, size = 0, units = "B", include = FALSE, simplify = TRUE)
```

Arguments

x	A character vector of filenames
size	A numeric vector, the size of the file
units	A character vector specifying the units. Options are B , KB, MB and GB. Must match the length of size if specifying more than one unit.
include	A boolean, include files of size size?
simplify	A boolean, should we create a data.table from the list of output?

Details

Automatically excludes any non-existant items

Value

A list or a data.table

Examples

```
# single file size and unit specification
x <- list.files(path = Sys.getenv("TEMP"), full.names = TRUE)[1:100]
filter_files(x, size = 1, units = "KB")

# multiple specifications
size <- c(0, rep(1, 4))
units <- c("B", "B", "KB", "MB", "GB")
res <- filter_files(x, size, units)
res$min_file_size <- factor(res$min_file_size, levels = paste(size, units))
table(res$min_file_size)
```

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floor *A function to extend the functionaliy of the base::floor function*

Description

A function to extend the functionaliy of the base::floor function

Usage

```
floor(x, digits = 0)
```

Arguments

x	A numeric
digits	The number of digits to the left of the decimal place to round to

Details

Usually used for purchase data when you need to floor the cents

Value

A numeric

Examples

```
x <- 99.9999
floor(x, 2)
# is equivalent to:
base::floor(100 * x)
```

`has_hive` *Check if the system has hive capabilities*

Description

Check if the system has hive capabilities

Usage

```
has_hive()
```

Details

Checks to see if the hive binaries are in the PATH variable

Value

A boolean

Examples

```
if(has_hive()) {
    print("yes")
} else {
    print("no")
}
```

`hive_datatypes` *Extract the R-datatypes for a hive table*

Description

Extract the R-datatypes for a hive table

Usage

```
hive_datatypes(schema, table_name)
```

Arguments

`schema`	A character string, the name of the hive schema
`table_name`	A character string, the name of the hive table

Details

For now the functions converts all number-y datatypes like integer, float, decimal to `numeric` and both date and string types to `character`.

Value

A `data.table` with columns:

- `name` The column name (`character`)
- `type` The R-datatype (`character`)

`hive_read` *Read a Hive table that is stored as a text file*

Description

Read a Hive table that is stored as a text file

Usage

```
hive_read(x, ...)
```

Arguments

`x`	A character string, the directory path of the hive table to read
`...`	Additional arguments to `fread`

Details

The function allows you to assign your own additional arguments to fread, but it defaults the separator to pipe ("|") and adds to the na.string to recognize the hive default.

Value

A `data.table`

`hread` *A function to perform a read of a hive table*

Description

A function to perform a read of a hive table

Usage

```
hread(table_name, schema, schema_loc, ...)
```

Arguments

table_name	A character string, the name of the hive table
schema	A character string, the name of the hive schema
schema_loc	A character string, the directory path of where the schema is located on the HDFS
...	Additional arguments to hive_read

Details

Will automatically read all files under the directory after finding the datatypes and column names from the hive metastore. Note that the schema can have a different physical location instead of being forced to have the schema_loc/schema naming convention.

Value

A data.table

Examples

```
## Not run:
schema_loc <- "/mapr/mapr03r/analytic_users/msmck/csm_synd_hive_schemas/csm_syndicated/"
hread("dictionary", "csm_syndicated", schema_loc)

## End(Not run)
```

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iri_week *A function to derive the IRI week for the date specified*

Description

A function to derive the IRI week for the date specified

Usage

```
iri_week(x, fmt = "%Y-%m-%d", ...)
```

Arguments

x	A date or character string
fmt	A date format
...	Additional arguments to as.Date including further arguments to be passed from or to other methods, including format for as.character and as.Date methods.

Value

A numeric

Examples

```
iri_week(Sys.Date())
iri_week("Dec. 12, 2016", "%b. %d, %Y")
```

is_defined *A function to perform a read of a hive table*

Description

A function to perform a read of a hive table

Usage

```
is_defined(..., .all = FALSE)
```

Arguments

...	A list of objects to test if they are null
.all	Boolean, should we return the tests for each individual element?

Details

Good for testing an input(s) to a function when it might be NULL

Examples

```
foo <- function(x = NULL, y = NULL) {
  if (is_defined(x, y)) {
    return(paste0(x, y))
  } else if (is_defined(x)) {
    return(x)
  } else if (any(is_defined(x, y, .all = TRUE))) {
    return("one is not null")
  }
}
foo(x = 1)
foo(x = 1, y = 2)
foo(y = 2)
```

make_compare_names *Make names for comparing two data sets*

Description

Make names for comparing two data sets

Usage

```
make_compare_names(compare, suffixes = c(".x", ".y"), sep = "_")
```

Arguments

compare	The column names to compare
suffixes	The suffixes for each set to use
sep	The separator between the names, the sep argument to the paste function

Value

A character vector

Examples

```
make_compare_names(compare = c("dollars", "units"), suffixes = c(".hive", ".sas"))
```

`ninja_load` *Load packages silently*

Description

Load packages silently

Usage

```
ninja_load(...)
```

Arguments

`...`	The quoted names of the packages you wish to load with deadly silence

Examples

```
# load some notoriously loud packages
## Not run:
ninja_load("data.table", "bit64")

## End(Not run)
```

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