# Package 'OuhscMunge'

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Title Data Manipulation Operations
<b>Description</b> Data manipulation operations frequently used in OUHSC BBMC projects.
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<pre>URL https://github.com/OuhscBbmc/OuhscMunge, http://ouhsc.edu/bbmc/</pre>
<pre>BugReports https://github.com/OuhscBbmc/OuhscMunge/issues</pre>
License GPL-2
LazyData TRUE
<b>Depends</b> $R(>=3.1.0)$
Imports devtools (>= 1.8.0), lubridate
Suggests RODBC, testthat
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R topics documented:
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clump\_month\_date

Assign date for a given year & month

#### **Description**

This accepts a date, but changes the day. Set/degrade/clump all the days within a month to the same day.

# Usage

```
clump_month_date(date_detailed, day_of_month = 15L)
```

#### **Arguments**

date\_detailed The Date value containing the desired year and month. The day will be over-

written. Required

day\_of\_month The factor label assigned to the missing value. Defaults to 15.

#### **Details**

We use this frequently to set/degrade/clump all the days to the middle of their respective month (ie, the 15th day). The midpoint of a month is usually the most appropriate summary location. It makes graphs more intuitive. Using the midpoint of month can also avoid problems with timezones. A date won't get nudged to a neighboring month accidentally.

#### Value

An array of Date values.

#### Note

A stop error will be thrown if date\_detailed is not a Date, or if day\_of\_month is not bounded by [1, 31]. Be careful that if you set a November date the 31st day, the result will be December 1st. Consequently, we recommend not setting the day to a value after the 28.

#### Author(s)

Will Beasley

# **Examples**

```
library(OuhscMunge)
detailed <- seq.Date(from=as.Date("2011-04-21"), to=as.Date("2011-07-14"), by="day")
clumped <- clump_month_date(detailed)
table(clumped)
# 2011-04-15 2011-05-15 2011-06-15 2011-07-15
# 10 31 30 14</pre>
```

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headstart\_utilities

Utilities for outputting characteristics of a datatset used it code.

# Description

These functions are used during the execution of a program. Rather they produce snippets that can be pasted into code, and help the developer avoid some typing.

#### Usage

```
column_rename_headstart( d )
column_class_headstart( d )
column_value_headstart( x )
```

#### **Arguments**

d A data.frame to describe.

x A vector to describe.

#### Value

Prints formatted code to the console.

#### Author(s)

Will Beasley

# **Examples**

```
column_rename_headstart(datasets::OrchardSprays)
column_class_headstart(datasets::OrchardSprays)
column_value_headstart(datasets::OrchardSprays$treatment)
```

OuhscMunge

Data manipulation operations frequently used in OUHSC BBMC projects. <a href="http://www.ouhsc.edu/bbmc/">http://www.ouhsc.edu/bbmc/</a>

#### **Description**

Thanks to Funders, including HRSA/ACF D89MC23154

OUHSC CCAN Independent Evaluation of the State of Oklahoma Competitive Maternal, Infant, and Early Childhood Home Visiting (MIECHV) Project., which evaluates MIECHV expansion and enhancement of Evidence-based Home Visitation programs in four Oklahoma counties.

#### **Details**

OuhscMunge.

#### Note

The release version will eventually be available through CRAN by running install.packages('OuhscMunge'). The most recent development version is available through GitHub by running devtools::install\_github (repo = 'OuhscBbmc/OuhscMunge') (make sure devtools is already installed). If you're having trouble with the package, please install the development version. If this doesn't solve your problem, please create a new issue, or email Will.

#### Author(s)

William Howard Beasley, University of Oklahoma Health Sciences Center, College of Medicine, Dept of Pediatrics, BBMC.

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#### **Examples**

```
## Not run:
# Install/update REDCapR with the release version from CRAN.
install.packages('OuhscMunge') #But it's not on CRAN yet.

# Install/update REDCapR with the development version from GitHub
#install.packages('devtools') #Uncomment if `devtools` isn't installed already.
devtools::install_github('OuhscBbmc/OuhscMunge')

## End(Not run)
```

replace\_nas\_with\_explicit

*Create explicit factor level for missing values.* 

# Description

Missing values are converted to a factor level. This explicit assignment can reduce the chances that missing values are inadvertantly ignored. It also allows the presence of a missing to become a predictor in models.

#### **Usage**

```
replace_nas_with_explicit(scores, new_na_label = "Unknown",
    create_factor = FALSE, add_unknown_level = FALSE)
```

# **Arguments**

An array of values, ideally either factor or character. Required

new\_na\_label The factor label assigned to the missing value. Defaults to Unknown.

create\_factor Converts scores into a factor, if it isn't one already. Defaults to FALSE.

add\_unknown\_level

Should a new factor level be created? (Specify TRUE if it already exists.) Defaults to FALSE.

#### Value

An array of values, where the NA values are now a factor level, with the label specified by the new\_na\_label value.

#### Note

The create\_factor parameter is respected only if scores isn't already a factor. Otherwise, levels without any values would be lost.

A stop error will be thrown if the operation fails to convert all the NA values.

#### Author(s)

Will Beasley

#### **Examples**

```
library(OuhscMunge) #Load the package into the current R session.
missing_indices <- c(3, 6, 8, 25)
# With a character variable:
a <- letters
a[missing_indices] <- NA_character_
a <- OuhscMunge::replace_nas_with_explicit(a)

# With a factor variable:
b <- factor(letters, levels=letters)
b[missing_indices] <- NA_character_
b <- OuhscMunge::replace_nas_with_explicit(b, add_unknown_level=TRUE)</pre>
```

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