

# Package ‘mde’

June 27, 2020

**Title** Missing Data Explorer

**Version** 0.2.1

**Description** Correct identification and handling of missing data is one of the most important steps in any analysis. To aid this process, 'mde' provides a very easy to use yet robust framework to quickly get an idea of where the missing data lies and therefore find the most appropriate action to take.

Graham WJ (2009) <doi:10.1146/annurev.psych.58.110405.085530>.

**License** GPL-3

**Depends** R(>= 3.6.0)

**Imports** dplyr(>= 0.8.9), tidyr(>= 1.0.3)

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.0

**URL** <https://github.com/Nelson-Gon/mde>

**BugReports** <https://github.com/Nelson-Gon/mde/issues>

**Suggests** knitr, rmarkdown, testthat, covr

**VignetteBuilder** knitr

**NeedsCompilation** no

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custom_na_recode	<i>Recode NA as another value using a function or a custom equation</i>
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## Description

Recode NA as another value using a function or a custom equation

## Usage

```
custom_na_recode(
  df,
  func = "mean",
  grouping_cols = NULL,
  across_columns = NULL
)
```

## Arguments

df	A valid R 'object' for which the percentage of missing values is required.
func	Function to use for the replacement e.g "mean". Defaults to mean.
grouping_cols	A character vector. If supplied, one can provide the columns by which to group the data.
across_columns	A character vector specifying across which columns recoding should be done #use all columns head(custom_na_recode(airquality,func="mean")) # use only a few columns head(custom_na_recode(airquality,func="mean",across_columns = c("Solar.R","Ozone"))) # use a function from another package #head(custom_na_recode(airquality, func=dplyr::lead)) some_data <- data.frame(ID=c("A1","A1","A1","A2","A2","A2"), A=c(5,NA,0,8,3,4), B=c(10,0,0,NA,5,6),C=c(1,NA,NA,25,7,8)) # grouping head(custom_na_recode(some_data,func = "mean", grouping_cols = "ID", across_columns = c("C", "A"))) head(custom_na_recode(some_data,func = "mean", grouping_cols = "ID"))

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drop_all_na	<i>Drop columns for which all values are NA</i>
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**Description**

Drop columns for which all values are NA

**Usage**

```
drop_all_na(df, grouping_cols = NULL)
```

**Arguments**

**df** A valid R 'object' for which the percentage of missing values is required.

**grouping\_cols** A character vector. If supplied, one can provide the columns by which to group the data.

**Examples**

```
test <- data.frame(ID= c("A","A","B","A","B"), Vals = c(rep(NA,4),2))
test2 <- data.frame(ID= c("A","A","B","A","B"), Vals = rep(NA, 5))
# drop columns where all values are NA
drop_all_na(test2)
# drop NAs only if all are NA for a given group, drops group too.
drop_all_na(test, "ID")
```

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drop_na_at	<i>Drop missing values at columns that match a given pattern</i>
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**Description**

Provides a simple yet efficient way to drop missing values("NA"s) at columns that match a given pattern.

**Usage**

```
drop_na_at(
  df,
  pattern_type = "contains",
  pattern = NULL,
  case_sensitive = FALSE,
  ...
)
```

**Arguments**

df	A data.frame object
pattern_type	One of "contains", "ends_with" or "starts_with"
pattern	The type of pattern to use when matching the pattern_type. The pattern is case sensitive
case_sensitive	Defaults to FALSE. Patterns are case insensitive if TRUE
...	Other params to other methods

**Value**

A data.frame object containing only columns that match the given pattern with the missing values removed.

**Examples**

```
head(drop_na_at(airquality, pattern_type = "starts_with", "O"))
```

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drop_na_if	<i>Condition based dropping of columns with missing values</i>
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**Description**

"drop\_na\_if" provides a simple way to drop columns with missing values if they meet certain criteria/conditions.

**Usage**

```
drop_na_if(df, sign = "gteq", percent_na = 50, keep_columns = NULL, ...)
```

**Arguments**

df	A data.frame object
sign	Character. One of gteq, lteq, lt, gt or eq which refer to greater than(gt) or equal(eq) or less than(lt) or equal to(eq) respectively.
percent_na	The percentage to use when dropping columns with missing values
keep_columns	Columns that should be kept despite meeting the target percent_na criterion(criteria)
...	Other arguments to "percent_missing"

**Value**

A data.frame object with columns that meet the target criteria dropped

**See Also**

[percent\\_missing](#)

**Examples**

```
head(drop_na_if(airquality, percent_na = 24))
#drop columns that have less than or equal to 4%
head(drop_na_if(airquality,sign="lteq", percent_na = 4))
# Drop all except with greater than or equal to 4% missing but keep Ozone
head(drop_na_if(airquality, sign="gteq",percent_na = 4, keep_columns = "Ozone"))
```

---

get\_na\_counts

*Add columnwise/groupwise counts of missing values*


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**Description**

This function takes a ‘data.frame’ object as an input and returns the corresponding ‘NA’ counts. ‘NA’ refers to R’s builtin missing data holder.

**Usage**

```
get_na_counts(x, grouping_cols = NULL)
```

**Arguments**

**x** A valid R ‘object’ for which ‘na\_counts’ are needed.

**grouping\_cols** A character vector. If supplied, one can provide the columns by which to group the data.

**Value**

An object of the same type as ‘x’ showing the respective number of missing values. If grouped is set to ‘TRUE’, the results are returned by group.

**Examples**

```
get_na_counts(airquality)
# Grouped counts
test <- data.frame(Subject = c("A","A","B","B"), res = c(NA,1,2,3),
ID = c("1","1","2","2"))
get_na_counts(test,grouping_cols =
c("ID", "Subject"))
```

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na_summary	<i>An all-in-one missingness report</i>
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**Description**

An all-in-one missingness report

**Usage**

```
na_summary(df, grouping_cols = NULL, sort_by = NULL, descending = FALSE, ...)
```

**Arguments**

df	A valid R ‘object’ for which the percentage of missing values is required.
grouping_cols	A character vector. If supplied, one can provide the columns by which to group the data.
sort_by	One of counts or percents. This determines whether the results are sorted by counts or percentages.
descending	Logical. Should missing values be sorted in decreasing order ie largest to smallest? Defaults to FALSE.
...	Arguments to other functions

**Examples**

```
na_summary(airquality)
# grouping
test2 <- data.frame(ID= c("A","A","B","A","B"),Vals = c(rep(NA,4),"No"),
ID2 = c("E","E","D","E","D"))
na_summary(test2,grouping_cols = c("ID","ID2"))
# sort summary
na_summary(airquality,sort_by = "percent_missing",descending = TRUE)
na_summary(airquality,sort_by = "percent_complete")
```

---

percent_missing	<i>Columnwise missingness percentages</i>
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**Description**

A convenient way to obtain percent missingness columnwise.

**Usage**

```
percent_missing(df, grouping_cols = NULL, exclude_cols = NULL)
```

**Arguments**

df	A valid R 'object' for which the percentage of missing values is required.
grouping_cols	A character vector. If supplied, one can provide the columns by which to group the data.
exclude_cols	A character vector indicating columns to exclude when returning results.

**Value**

An object of the same class as x showing the percentage of missing values.

**Examples**

```
test <- data.frame(ID= c("A","B","A","B","A","B","A"), Vals = c(NA,25,34,NA,67,NA,45))
percent_missing(test,grouping_cols = "ID")
percent_missing(airquality)
percent_missing(airquality,exclude_cols = c("Day","Temp"))
```

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recode_as_na	<i>Recode a value as NA</i>
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**Description**

This provides a convenient way to convert a number/value that should indeed be an "NA" to "NA". In otherwords, it converts a value to R's recognized NA.

**Usage**

```
recode_as_na(
  df,
  value = NULL,
  subset_cols = NULL,
  pattern_type = NULL,
  pattern = NULL,
  case_sensitive = FALSE,
  ...
)
```

**Arguments**

df	A data.frame object for which recoding is to be done.
value	The value to convert to 'NA'. We can for instance change "n/a" to 'NA' or any other value.
subset_cols	An optional character vector to define columns for which changes are required.
pattern_type	One of 'contains', 'starts_with' or 'ends_with'.
pattern	A character pattern to match
case_sensitive	Defaults to FALSE. Patterns are case insensitive if TRUE
...	Other arguments to other functions

**Value**

An object of the same class as x with values changed to 'NA'.

**Examples**

```
head(recode_as_na(airquality,value=c(67,118),pattern_type="starts_with",pattern="S|O"))
head(recode_as_na(airquality,value=c(41),pattern_type="ends_with",pattern="e"))
head(recode_as_na(airquality, value=41,subset_cols="Ozone"))
```

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recode_as_na_for	<i>Recode Values as NA if they meet defined criteria</i>
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**Description**

Recode Values as NA if they meet defined criteria

**Usage**

```
recode_as_na_for(df, criteria = "gt", value = 0, subset_cols = NULL)
```

**Arguments**

df	A data.frame object to manipulate
criteria	One of gt,gteq,lt,lteq to define greater than, greater than or equal to, less than or less than or equal to.
value	The value to convert to 'NA'. We can for instance change "n/a" to 'NA' or any other value.
subset_cols	An optional character vector for columns to manipulate.

**Value**

A data.frame object with the required changes.

**Examples**

```
recode_as_na_for(airquality,value=36, criteria = "gteq",
subset_cols = c("Ozone","Solar.R"))
```



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recode_na_as	<i>Replace missing values with another value</i>
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### Description

This provides a convenient way to recode "NA" as another value for instance "NaN", "n/a" or any other value a user wishes to use.

### Usage

```
recode_na_as(
  df,
  value = 0,
  subset_cols = NULL,
  pattern_type = NULL,
  pattern = NULL,
  case_sensitive = FALSE,
  ...
)
```

### Arguments

df	A data.frame object for which recoding is to be done.
value	The value to convert to 'NA'. We can for instance change "n/a" to 'NA' or any other value.
subset_cols	An optional character vector to define columns for which changes are required.
pattern_type	One of 'contains', 'starts_with' or 'ends_with'.
pattern	A character pattern to match
case_sensitive	Defaults to FALSE. Patterns are case insensitive if TRUE
...	Other arguments to other functions

### Value

An object of the same type as x with NAs replaced with the desired value.

### Examples

```
head(recode_na_as(airquality, "n/a"))
head(recode_na_as(airquality, subset_cols = "Ozone", value = "N/A"))
head(recode_na_as(airquality, value=0, pattern_type="starts_with",pattern="Solar"))
```

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recode_na_if	<i>Recode NA as another value with some conditions</i>
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**Description**

Recode NA as another value with some conditions

**Usage**

```
recode_na_if(df, grouping_cols = NULL, target_groups = NULL, replacement = 0)
```

**Arguments**

df	A data.frame object with missing values
grouping_cols	Character columns to use for grouping the data
target_groups	Character Recode NA as if and only if the grouping column is in this vector of values
replacement	Values to use to replace NAs for IDs that meet the requirements. Defaults to 0.

**Examples**

```
some_data <- data.frame(ID=c("A1", "A2", "A3", "A4"),
  A=c(5, NA, 0, 8), B=c(10, 0, 0, 1), C=c(1, NA, NA, 25))
# Replace NAs with 0s only for IDs in A2 and A3
recode_na_if(some_data, "ID", c("A2", "A3"), replacement=0)
```

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sort_by_missingness	<i>Sort Variables according to missingness</i>
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**Description**

Provides a useful way to sort the variables(columns) according to their missingness.

**Usage**

```
sort_by_missingness(x, sort_by = "counts", descending = FALSE, ...)
```

**Arguments**

x	A valid R 'object' for which 'na_counts' are needed.
sort_by	One of counts or percents. This determines whether the results are sorted by counts or percentages.
descending	Logical. Should missing values be sorted in decreasing order ie largest to smallest? Defaults to FALSE.
...	Other arguments to specific functions. See "See also below"

**Value**

A 'data.frame' object sorted by number/percentage of missing values

**See Also**

[get\\_na\\_counts](#) [percent\\_missing](#)

**Examples**

```
sort_by_missingness(airquality, sort_by = "counts")  
# sort by percents  
sort_by_missingness(airquality, sort_by="percents")  
# descending order  
sort_by_missingness(airquality, descend = TRUE)
```

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