Package 'sanityTracker'

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Type Package Title Keeps Track of all Performed Sanity Checks Version 0.1.0 Date 2020-04-14 Maintainer Marsel Scheer <scheer@freescience.de> **Description** During the preparation of data set(s) one usually performs some sanity checks. The idea is that irrespective of where the checks are performed, they are centralized by this package in order to list all at once with examples if a check failed. License GPL-3 **Encoding** UTF-8 LazyData true RoxygenNote 7.0.2 **Imports** data.table (>= 1.12.2), checkmate (>= 2.0.0) Suggests testthat, knitr, rmarkdown VignetteBuilder knitr URL https://github.com/MarselScheer/sanityTracker BugReports https://github.com/MarselScheer/sanityTracker/issues NeedsCompilation no

Author Marsel Scheer [aut, cre] Repository CRAN Date/Publication 2020-04-22 16:12:07 UTC

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.add_sanity_check Adds a sanity check to the list of already performed sanity checks

Description

NOTE the also add_sanity_check calls this function, the parameters are documented in add_sanity_check because that function gets exported.

Usage

```
.add_sanity_check(
  fail_vec,
  description,
  counter_meas,
  data,
  data_name,
  example_size,
  param_name,
  call,
  fail_callback,
  .fail_vec_str,
  .generated_desc
)
```

Arguments

fail_vec	see add_sanity_check
description	see add_sanity_check
counter_meas	see add_sanity_check
data	see add_sanity_check
data_name	see add_sanity_check
example_size	see add_sanity_check

param_name	see add_sanity_check
call	see add_sanity_check
fail_callback	see add_sanity_check
.fail_vec_str	should capture what was used originally for fail_vec.
.generated_des	c
	for convenience functions like sc_col_elements to provide additional informa- tion about the check.

Value

see add_sanity_check

add_sanity_check Adds a sanity check to the list of already performed sanity checks

Description

Adds a sanity check to the list of already performed sanity checks

Usage

```
add_sanity_check(
  fail_vec,
  description = "-",
  counter_meas = "-",
  data,
  data_name = checkmate::vname(x = data),
  example_size = 3,
  param_name = "-",
  call = h_deparsed_sys_call(which = -3),
  fail_callback
)
```

Arguments

fail_vec	logical vector where TRUE indicates that a fail has happend
description	(optional) of the sanity check. default is "-".
counter_meas	(optional) description of the counter measures that were applied to correct the problems. default is "-".
data	(optional) where the fails were found. Is used to store examples of failures. default is "-".
data_name	(optional) name of the data set that was used. defaults is the name of the object passed to data.
example_size	(optional) number failures to be extracted from the object passed to data. By default 3 random examples are extracted.

param_name	(optional) name of the parameter(s) that is used. This may be helpful for filtering the table of all performed sanity checks.
call	(optional) by default tracks the function that called add_sanity_check.
fail_callback	(optional) user-defined function that is called if any element of fail_vec is TRUE. This is helpful if an additional warning or error should be thrown or maybe a log-entry should be created.

Value

a list with three elements

entry_sanity_table invisibly the sanity check that is stored internally with the other sanity checks

fail_vec fail_vec as passed over to this function

fail TRUE if any element of fail is TRUE. Otherwise FALSE.

All performed sanity checks can be fetched via get_sanity_checks

Examples

```
d <- data.frame(person_id = 1:4, bmi = c(18,23,-1,35), age = 31:34)
dummy_call <- function(x) {</pre>
  add_sanity_check(
   x$bmi < 15,
   description = "bmi above 15",
   counter_meas = "none",
   data = x,
   param_name = "bmi")
  add_sanity_check(
   x$bmi > 30,
   description = "bmi below 30",
   counter_meas = "none")
}
dummy_call(x = d)
get_sanity_checks()
add_sanity_check(
   d$bmi < 15,
   description = "bmi above 15",
   fail_callback = warning)
```

clear_sanity_checks Removes all tracked sanity checks

Description

Removes all tracked sanity checks

Usage

clear_sanity_checks()

Description

Returns all performed sanity checks

Usage

get_sanity_checks()

Value

all sanity checks, i.e. a data.table with the following column

- **description** character that was provided by the user through the parameter description
- additional_desc character that provides additional information about the check that was generated by the convenience functions
- **data_name** name of the data set that passed to the function that performed the sanity check. This can also be specified by the user
- **n** a logical vector is the basis of all sanity checks. This is length of the logical vector that was used, which in general is the number of rows of the table that was checked
- n_fail how often the logical vector was TRUE
- **n_na** how often the logical vector was NA
- **counter_meas** character provided by the user about how a fail will be addressed. Note that this never happens inside a function of sanityTracker but is realized by the user after the check was performed. It is only for documentation when the results of the checks are displayed.
- fail_vec_str this captures how the actual logical vector of fails was build
- param_name usually generated by the convenience functions and it also may be a composition of more than one parameter name. However this parameter could also have been provided by the user
- call character of the call where the sanity check happend
- **example** if a check failed and the table is available, then some examples of rows that lead to the fail are stored here

See Also

add_sanity_check

h_add_sanity_check Wrapper for add_sanity_check for internal use

Description

The convenience function usually provide some defaults like description that can be overwritten by the user through the ... argument of the convenience function. This function manages to set those values that were NOT overwritten by the user through the ... argument and then call add_sanity_check.

Usage

```
h_add_sanity_check(
  ellipsis,
  fail_vec,
  .generated_desc,
  data,
  data_name = "",
  param_name = "",
  call = h_deparsed_sys_call(which = -2),
  .fail_vec_str = checkmate::vname(x = fail_vec)
)
```

Arguments

ellipsis	usually list() of the function that calls this function. It contains the parameters defined by the user for add_sanity_check.
fail_vec	logical vector where TRUE indicates that a fail has happend
.generated_des	c
	will be passed to .add_sanity_check if ellipsis does not contain a element with name 'description'
data	will be passed to .add_sanity_check if ellipsis does not contain a element with name 'data'
data_name	will be passed to .add_sanity_check if ellipsis does not contain a element with name 'data_name'
param_name	will be passed to .add_sanity_check if ellipsis does not contain a element with name 'param_name'
call	will be passed to .add_sanity_check if ellipsis does not contain a element with name 'call'
.fail_vec_str	usually not used by the user. Captures what was passed to fail_vec.

Value

see return value of add_sanity_check

h_collapse_char_vec

Examples

```
d <- data.frame(type = letters[1:4], nmb = 1:4)
# h_add_sanity_check is used on sc_col_elements()
sc_col_elements(object = d, col = "type", feasible_elements = letters[2:4])
get_sanity_checks()</pre>
```

h_collapse_char_vec Collapse a vector of characters to a string with separators

Description

Collapse a vector of characters to a string with separators

Usage

```
h_collapse_char_vec(v, collapse = ", ", qoute = "'")
```

Arguments

v	vector of chars to be collapsed
collapse	character that separates the elements in the returned object
qoute	character that surronds every element in \boldsymbol{v} in the returned object

Value

collapsed version of v

Examples

```
cat(sanityTracker:::h_collapse_char_vec(v = letters[1:4]))
```

h_complete_list	Extends a list with an named element if the element does not exist	ţ

Description

Extends a list with an named element if the element does not exist

Usage

h_complete_list(ell, name, value)

Arguments

ell	list to be extended (usually an ellipsis as list())
name	character with the name for the element to be added
value	value that will be stored in ell[[el_name]]

Value

if ell already contained the element name, then ell is returned without being modified. Otherwise, ell is returned extended by a new element with name name and value value.

Examples

```
ell <- list(a = 1, b = 2)
sanityTracker:::h_complete_list(ell = ell, name = "a", value = 100)
sanityTracker:::h_complete_list(ell = ell, name = "d", value = Inf)</pre>
```

h_deparsed_sys_call Simply converts a call into a character

Description

Simply converts a call into a character

Usage

```
h_deparsed_sys_call(which)
```

Arguments

which see sys.call. However the function bounds it by the number of encolsing environments.

Value

the call of the corresponding environment as character

sc_cols_bounded

Description

Checks that all elements from the specified columns are in a certain range

Usage

```
sc_cols_bounded(object, cols, rule = "(-Inf, Inf)", ...)
```

Arguments

object	table with a columns specified by cols
cols	vector of characters of columns that are checked against the specified range
rule	check as two numbers separated by a comma, enclosed by square brackets (end- point included) or parentheses (endpoint excluded). For example, " $[0, 3)$ " re- sults in all(x >= 0 & x < 3). The lower and upper bound may be omitted which is the equivalent of a negative or positive infinite bound, respectively. By defi- nition [0,] contains Inf, while [0,) does not. The same holds for the left (lower) boundary and -Inf. This explanation was copied from checkmate::qtest. That function is also the backbone of the this function.
	further parameters that are passed to add_sanity_check.

Value

list of logical vectors where TRUE indicates where the check failed. Every list entry represents one of the columns specified in cols. This might be helpful if one wants to apply a counter-measure

Examples

```
dummy_call <- function(x) {
   sc_cols_bounded(object = iris, cols = c("Sepal.Length", "Petal.Length"),
   rule = "[1, 7.9)")
}
dummy_call(x = d)
get_sanity_checks()</pre>
```

sc_cols_bounded_above Checks that all elements from the given columns are below a certain number

Description

Checks that all elements from the given columns are below a certain number

Usage

```
sc_cols_bounded_above(
   object,
   cols,
   upper_bound,
   include_upper_bound = TRUE,
   ...
)
```

Arguments

object	table with a columns specified by cols
cols	vector of characters of columns that are checked against the specified range
upper_bound	elements of the specified columns must be below this bound
include_upper_b	bound
	if TRUE (default), elements are allowed to be equal to the upper_bound
	further parameters that are passed to add_sanity_check.

Value

list of logical vectors where TRUE indicates where the check failed. Every list entry represents one of the columns specified in cols. This might be helpful if one wants to apply a counter-measure

sc_cols_bounded_below Checks that all elements from the given columns are above a certain number

Description

Checks that all elements from the given columns are above a certain number

sc_cols_non_NA

Usage

```
sc_cols_bounded_below(
   object,
   cols,
   lower_bound,
   include_lower_bound = TRUE,
   ...
)
```

Arguments

object	table with a columns specified by cols	
cols	vector of characters of columns that are checked against the specified range	
lower_bound	elements of the specified columns must be above this bound	
include_lower_bound		
	if TRUE (default), elements are allowed to be equal to the lower_bound	
	further parameters that are passed to add_sanity_check.	

Value

list of logical vectors where TRUE indicates where the check failed. Every list entry represents one of the columns specified in cols. This might be helpful if one wants to apply a counter-measure

Examples

```
d <- data.frame(a = c(0, 0.2, 3, Inf), b = c(1:4))
dummy_call <- function(x) {
    sc_cols_bounded_below(
        object = d, cols = c("a", "b"),
        lower_bound = 0.2,
        include_lower_bound = FALSE,
        description = "Measurements are expected to be bounded from below")
}
dummy_call(x = d)
get_sanity_checks()</pre>
```

sc_cols_non_NA Checks that all elements from the specified columns are not NA

Description

Checks that all elements from the specified columns are not NA

Usage

```
sc_cols_non_NA(object, cols = names(object), ..., unk_cols_callback = stop)
```

Arguments

object	table with a columns specified by cols	
cols	vector of characters of columns that are checked for NAs	
	further parameters that are passed to add_sanity_check.	
unk_cols_callback		
	user-defined function that is called if some of the cols are not contained in the object. This is helpful if an additional warning or error should be thrown or	
	object. This is helpful if an additional warning of error should be unown of	

maybe a log-entry should be created. Default is the function stop

Value

a list where every element is an object returned by add_sanity_check for each column specified in cols that exists in object

Examples

```
iris[c(1,3,5,7,9), 1] <- NA
dummy_call <- function(x) {</pre>
  sc_cols_non_NA(object = iris, description = "No NAs expected in iris")
}
dummy_call(x = iris)
get_sanity_checks()
```

sc_cols_positive Checks that all elements from the specified columns are positive

Description

Checks that all elements from the specified columns are positive

Usage

```
sc_cols_positive(object, cols, zero_feasible = TRUE, ...)
```

Arguments

object	table with a columns specified by cols
cols	vector of characters of columns that are checked against the specified range
zero_feasible	if zero is in the range or not
	further parameters that are passed to add_sanity_check.

Value

list of logical vectors where TRUE indicates where the check failed. Every list entry represents one of the columns specified in cols. This might be helpful if one wants to apply a counter-measure.

sc_cols_unique

Examples

```
d <- data.frame(a = c(0, 0.2, 3, Inf), b = c(1:4))
dummy_call <- function(x) {
    sc_cols_positive(d, cols = c("a", "b"), zero_feasible = FALSE,
    description = "Measurements are expected to be positive")
}
dummy_call(x = d)
get_sanity_checks()</pre>
```

sc_cols_unique Checks that the combination of the specified columns is unique

Description

Checks that the combination of the specified columns is unique

Usage

```
sc_cols_unique(object, cols = names(object), ...)
```

Arguments

object	table with a columns specified by cols
cols	vector of characters which combination is checked to be unique
	further parameters that are passed to add_sanity_check.

Value

see return object of add_sanity_check. Note that if a combination appears 3 times, then n_fail will increased by 3.

Examples

```
dummy_call <- function(x) {
  sc_cols_unique(
    object = x,
    cols = c("Species", "Sepal.Length",
        "Sepal.Width", "Petal.Length"))
}
dummy_call(x = iris)
get_sanity_checks()
get_sanity_checks()[["example"]]</pre>
```

sc_col_elements

Description

Checks that the elements of a column belong to a certain set

Usage

```
sc_col_elements(object, col, feasible_elements, ...)
```

Arguments

object	table with a column specified by col	
col	name as a character of the column which is checked	
feasible_elements		
	vector with characters that are feasible for col. Note that an element that is NA it is always counted as a fail if feasible_elements does not explicitly contains NA.	
	further parameters that are passed to add_sanity_check.	

Value

see return object of add_sanity_check

Examples

```
d <- data.frame(type = letters[1:4], nmb = 1:4)
dummy_call <- function(x) {
    sc_col_elements(object = d, col = "type", feasible_elements = letters[2:4])
}
dummy_call(x = d)
get_sanity_checks()</pre>
```

sc_left_join Performs various checks after a left-join was performed

Description

One check is that no rows were duplicated during merge and the other check is that no columns were duplicated during merge.

Usage

```
sc_left_join(joined, left, right, by, ..., find_nonunique_key = TRUE)
```

sc_left_join

Arguments

joined	the result of the left-join	
left	the left table used in the left-join	
right	the right table used in the left-join	
by	the variables used for the left-join	
	further parameters that are passed to add_sanity_check.	
find_nonunique_key		
	if TRUE a sanity-check is performed that finds keys (defined by by) that non-unique. However this can be a time-consuming step.	

Value

list with two elements for the two sanity checks performed by this function. The structure of each element is as the return object of add_sanity_check.

Examples

```
ab <- data.table::data.table(a = 1:4, b = letters[1:4])
abc <- data.table::data.table(a = c(1:4, 2), b = letters[1:5], c = rnorm(5))
j <- merge(x = ab, y = abc, by = "a")
dummy_call <- function() {
    sc_left_join(joined = j, left = ab, right = abc, by = "a",
    description = "Left join outcome to main population")
}
dummy_call()
get_sanity_checks()</pre>
```

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