Package ‘assertr’

February 23, 2018

Type Package
Title Assertive Programming for R Analysis Pipelines
Version 2.5
Description Provides functionality to assert conditions that have to be met so that errors in data used in analysis pipelines can fail quickly. Similar to 'stopifnot()' but more powerful, friendly, and easier for use in pipelines.

URL https://github.com/ropensci/assertr

BugReports https://github.com/ropensci/assertr/issues
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assert

Raises error if predicate is FALSE in any columns selected

Description

Meant for use in a data analysis pipeline, this function will just return the data it’s supplied if there are no FALSEs when the predicate is applied to every element of the columns indicated. If any element in any of the columns, when applied to the predicate, is FALSE, then this function will raise an error, effectively terminating the pipeline early.

Usage

```r
assert(data, predicate, ..., success_fun = success_continue, error_fun = error_stop)
```

```r
assert_(data, predicate, ..., .dots, success_fun = success_continue, error_fun = error_stop)
```

Arguments

- `data` A data frame
- `predicate` A function that returns FALSE when violated
- `...` Comma separated list of unquoted expressions. Uses dplyr’s `select` to select columns from data.
- `success_fun` Function to call if assertion passes. Defaults to returning data.
assert

  error_fun  Function to call if assertion fails. Defaults to printing a summary of all errors.
  .dots     Use assert_() to select columns using standard evaluation.

Details

  For examples of possible choices for the success_fun and error_fun parameters, run help("success_and_error_functions")

Value

  By default, the data is returned if predicate assertion is TRUE and and error is thrown if not. If a non-default success_fun or error_fun is used, the return values of these function will be returned.

Note

  See vignette("assertr") for how to use this in context

See Also

  verify insist assert_rows insist_rows

Examples

  # returns mtcars
  assert(mtcars, not_na, vs)

  # return mtcars
  assert(mtcars, not_na, mpg:carb)

  library(magrittr)    # for piping operator

  mtcars %>%
      assert(in_set(c(0,1)), vs)
  # anything here will run

  ## Not run:
  mtcars %>%
      assert(in_set(c(1, 2, 3, 4, 6), carb)
  # the assertion is untrue so
  ## nothing here will run
  ## End(Not run)
assertr package supplies a suite of functions designed to verify assumptions about data early in an analysis pipeline.

Details

See the assertr vignette or the documentation for more information.

You may also want to read the documentation for the functions that assertr provides:

- assert
- verify
- insist
- assert_rows
- insist_rows
- not_na
- in_set
- has_all_names
- is_uniq
- num_row_NAs
- maha_dist
- col_concat
- within_bounds
- within_n_sds
- within_n_mads
- success_and_error_functions
- chaining_functions

Examples

library(magrittr)  # for the piping operator
library(dplyr)

# this confirms that
# - that the dataset contains more than 10 observations
# - that the column for 'miles per gallon' (mpg) is a positive number
# - that the column for 'miles per gallon' (mpg) does not contain a datum
# - that is outside 4 standard deviations from its mean, and
# - that the am and vs columns (automatic/manual and v/straight engine,
assert_rows

# respectively contain 0s and 1s only
# - each row contains at most 2 NAs
# - each row's mahalanobis distance is within 10 median absolute deviations of
# all the distance (for outlier detection)

mtcars %>%
  verify(nrow(.) > 10) %>%
  verify(mpg > 0) %>%
  insist(within_n_sds(4), mpg) %>%
  assert(in_set(0,1), am, vs) %>%
  assert_rows(num_row_NAs, within_bounds(0,2), everything()) %>%
  insist_rows(maha_dist, within_n_mads(10), everything()) %>%
  group_by(cyl) %>%
  summarise(avg.mpg=mean(mpg))

---

assert_rows Raises error if predicate is FALSE for any row after applying row reduction function

Description

Meant for use in a data analysis pipeline, this function applies a function to a data frame that reduces each row to a single value. Then, a predicate function is applied to each of the row reduction values. If any of these predicate applications yield FALSE, this function will raise an error, effectively terminating the pipeline early. If there are no FALSEs, this function will just return the data that it was supplied for further use in later parts of the pipeline.

Usage

assert_rows(data, row_reduction_fn, predicate, ...,
  success_fun = success_continue, error_fun = error_stop)

assert_rows_(data, row_reduction_fn, predicate, ...,.dots,
  success_fun = success_continue, error_fun = error_stop)

Arguments

data A data frame
row_reduction_fn A function that returns a value for each row of the provided data frame
predicate A function that returns FALSE when violated
... Comma separated list of unquoted expressions. Uses dplyr's select to select columns from data.
success_fun Function to call if assertion passes. Defaults to returning data.
error_fun Function to call if assertion fails. Defaults to printing a summary of all errors.
.dots Use assert_rows_() to select columns using standard evaluation.
chaining_functions

Details
For examples of possible choices for the success_fun and error_fun parameters, run help("success_and_error_functions").

Value
By default, the data is returned if predicate assertion is TRUE and and error is thrown if not. If a non-default success_fun or error_fun is used, the return values of these function will be returned.

Note
See vignette("assertr") for how to use this in context

See Also
inspect_rows assert verify insist

Examples

# returns mtcars
assert_rows(mtcars, num_row_NAs, within_bounds(0,2), mpg:carb)

library(magrittr) # for piping operator

mtcars %>%
  assert_rows(rowSums, within_bounds(0,2), vs:am)
  # anything here will run

## Not run:
mtcars %>%
  assert_rows(rowSums, within_bounds(0,1), vs:am)
  # the assertion is untrue so
  # nothing here will run
## End(Not run)

chaining_functions  Chaining functions

Description
These functions are for starting and ending a sequence of assertr assertions and overriding the default behavior of assertr halting execution on the first error.

Usage

chain_start(data)

chain_end(data, success_fun = success_continue, error_fun = error_report)
Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>data</td>
<td>A data frame</td>
</tr>
<tr>
<td>success_fun</td>
<td>Function to call if assertion passes. Defaults to returning data.</td>
</tr>
<tr>
<td>error_fun</td>
<td>Function to call if assertion fails. Defaults to printing a summary of all errors.</td>
</tr>
</tbody>
</table>

Details

For more information, read the relevant section in this package’s vignette using, vignette("assertr")

For examples of possible choices for the `success_fun` and `error_fun` parameters, run `help("success_and_error_functions")`

Examples

```r
library(magrittr)

mtcars %>%
  chain_start() %>%
  verify(nrow(mtcars) > 10) %>%
  verify(mpg > 0) %>%
  insist(within_n_sds(4), mpg) %>%
  assert(in_set(0:1), am, vs) %>%
  chain_end()
```

Description

This function will return a vector, with the same length as the number of rows of the provided data frame. Each element of the vector will be it’s corresponding row with all of its values (one for each column) "pasted" together in a string.

Usage

```r
col_concat(data, sep = "")
```

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>data</td>
<td>A data frame</td>
</tr>
<tr>
<td>sep</td>
<td>A string to separate the columns with (default: &quot;)&quot;)</td>
</tr>
</tbody>
</table>

Value

A vector of rows concatenated into strings

See Also

`paste`
Examples

```r
col_concat(mtcars)

library(magrittr) # for piping operator

# you can use "assert_rows"", "is_uniq", and this function to
# check if joint duplicates (across different columns) appear
# in a data frame
## Not run:
mtcars %>%
  assert_rows(col_concat, is_uniq, mpg, hp)
  # fails because the first two rows are jointly duplicates
  # on these two columns

## End(Not run)

mtcars %>%
  assert_rows(col_concat, is_uniq, mpg, hp, wt) # ok
```

---

`has_all_names`  
*Returns TRUE if data.frame or list has specified names*

Description

This function checks parent frame environment for existence of names. This is meant to be used with `assertr`'s `verify` function to check for the existence of specific column names in a `data.frame` that is piped to `verify`. It can also work on a non-`data.frame` list.

Usage

```r
has_all_names(...)```

Arguments

...  
A arbitrary amount of quoted names to check for

Value

TRUE is all names exist, FALSE if not

See Also

`exists`
Examples

```r
verify(mtcars, has_all_names("mpg", "wt", "qsec"))
library(magrittr) # for pipe operator
## Not run:
mtcars %>%
  verify(has_all_names("mpgg")) # fails
## End(Not run)
mpgg <- "something"

mtcars %>%
  verify(exists("mpgg")) # passes but big mistake
## Not run:
mtcars %>%
  verify(has_all_names("mpgg")) # correctly fails
## End(Not run)
```

---

**insist**

*Raises error if dynamically created predicate is FALSE in any columns selected*

**Description**

Meant for use in a data analysis pipeline, this function applies a predicate generating function to each of the columns indicated. It will then use these predicates to check every element of those columns. If any of these predicate applications yield FALSE, this function will raise an error, effectively terminating the pipeline early. If there are no FALSES, this function will just return the data that it was supplied for further use in later parts of the pipeline.

**Usage**

```r
insist(data, predicate_generator, ..., success_fun = success_continue,
       error_fun = error_stop)

insist_(data, predicate_generator, ..., .dots, success_fun = success_continue,
         error_fun = error_stop)
```

**Arguments**

- `data` A data frame
predicate_generator
A function that is applied to each of the column vectors selected. This will produce, for every column, a true predicate function to be applied to every element in the column vectors selected.

... Comma separated list of unquoted expressions. Uses dplyr's select to select columns from data.

success_fun Function to call if assertion passes. Defaults to returning data.

error_fun Function to call if assertion fails. Defaults to printing a summary of all errors.

dots Use insist() to select columns using standard evaluation.

Details
For examples of possible choices for the success_fun and error_fun parameters, run help("success_and_error_functions")

Value
By default, the data is returned if dynamically created predicate assertion is TRUE and an error is thrown if not. If a non-default success_fun or error_fun is used, the return values of these functions will be returned.

Note
See vignette("assertr") for how to use this in context

See Also
assert verify insist_rows assert_rows

Examples

insist(iris, within_n_sds(3), Sepal.Length)  # returns iris

library(magrittr)

iris %>%
  insist(within_n_sds(4), Sepal.Length:Petal.Width)
# anything here will run

## Not run:
iris %>%
  insist(within_n_sds(3), Sepal.Length:Petal.Width)
# datum at index 16 of 'Sepal.Width' vector is (4.4)
# is outside 3 standard deviations from the mean of Sepal.Width.
# The check fails, raises a fatal error, and the pipeline
# is terminated so nothing after this statement will run
## End(Not run)
insist_rows

Raises error if dynamically created predicate is FALSE for any row after applying row reduction function

Description

Meant for use in a data analysis pipeline, this function applies a function to a data frame that reduces each row to a single value. Then, a predicate generating function is applied to row reduction values. It will then use these predicates to check each of the row reduction values. If any of these predicate applications yield FALSE, this function will raise an error, effectively terminating the pipeline early. If there are no FALSEs, this function will just return the data that it was supplied for further use in later parts of the pipeline.

Usage

insist_rows(data, row_reduction_fn, predicate_generator, ..., success_fun = success_continue, error_fun = error_stop)

insist_rows_(data, row_reduction_fn, predicate_generator, ..., .dots, success_fun = success_continue, error_fun = error_stop)

Arguments

data A data frame
row_reduction_fn A function that returns a value for each row of the provided data frame
predicate_generator A function that is applied to the results of the row reduction function. This will produce a true predicate function to be applied to every element in the vector that the row reduction function returns.
... Comma separated list of unquoted expressions. Uses dplyr's select to select columns from data.
success_fun Function to call if assertion passes. Defaults to returning data.
error_fun Function to call if assertion fails. Defaults to printing a summary of all errors.
.dots Use insist_rows_() to select columns using standard evaluation.

Details

For examples of possible choices for the success_fun and error_fun parameters, run help("success_and_error_functions")

Value

By default, the data is returned if dynamically created predicate assertion is TRUE and and error is thrown if not. If a non-default success_fun or error_fun is used, the return values of these function will be returned.
in_set

Returns TRUE if value in set

Description
This function returns a predicate function that will take a single value and return TRUE if the value is a member of the set of objects supplied. This doesn’t actually check the membership of anything—it only returns a function that actually does the checking when called with a value. This is a convenience function meant to return a predicate function to be used in an assertr assertion.

Usage
in_set(..., allow.na = TRUE)

Arguments
... objects that make up the set
allow.na A logical indicating whether NAs (including NaNs) should be permitted (default TRUE)

Examples
# returns mtcars
insist_rows(mtcars, maha_dist, within_n_mads(30), mpg:carb)

library(magrittr) # for piping operator

mtcars %>%
  insist_rows(maha_dist, within_n_mads(10), vs:am)
  # anything here will run

## Not run:
mtcars %>
  insist_rows(maha_dist, within_n_mads(1), everything())
  # the assertion is untrue so
  # nothing here will run
## End(Not run)

Note
See vignette("assertr") for how to use this in context

See Also
inspect_rows assert verify
**is_uniq**

Returns **TRUE** where no elements appear more than once

**Description**

This function is meant to take only a vector. It relies heavily on the `duplicated` function where it can be thought of as the inverse. Where this function differs, though—besides being only meant for one vector or column—is that it marks the first occurrence of a duplicated value as "non unique", as well.

**Usage**

```r
is_uniq(x, allow.na = FALSE)
```

**Value**

A function that takes one value and returns **TRUE** if the value is in the set defined by the arguments supplied by `in_set` and **FALSE** otherwise.

**See Also**

`%in%`

**Examples**

```r
predicate <- in_set(3, 4)
predicate(4)

## is equivalent to

in_set(3, 4)(3)

# the remainder of division by 2 is always 0 or 1
rem <- 10 %% 2
in_set(0, 1)(rem)

## this is meant to be used as a predicate in an assert statement
assert(mtcars, in_set(3, 4, 5), gear)

## or in a pipeline, like this was meant for
library(magrittr)

mtcars %>%
  assert(in_set(3, 4, 5), gear) %>%
  assert(in_set(0, 1), vs, am)
```
Arguments

- x: A vector to check for unique elements in
- allow.na: A logical indicating whether NAs should be preserved as missing values in the return value (FALSE) or if they should be treated just like any other value (TRUE) (default is FALSE)

Value

A vector of the same length where the corresponding element is TRUE if the element only appears once in the vector and FALSE otherwise

See Also

duplicated

Examples

```r
is_uniq(1:10)
```

```r
## not run:
# returns FALSE where a "5" appears
is_uniq(c(1:10, 5))
## End(not run)
```

```r
library(magrittr)
```

```r
## not run:
# this fails 4 times
mtcars %>% assert(is_uniq, qsec)
## End(not run)
```

---

**maha_dist**

*Computes mahalanobis distance for each row of data frame*

Description

This function will return a vector, with the same length as the number of rows of the provided data frame, corresponding to the average mahalanobis distances of each row from the whole data set.

Usage

```r
maha_dist(data, keep.NA = TRUE, robust = FALSE, stringsAsFactors = FALSE)
```
**maha_dist**

**Arguments**

- **data** A data frame
- **keep.NA** Ensure that every row with missing data remains NA in the output? TRUE by default.
- **robust** Attempt to compute mahalanobis distance based on robust covariance matrix? FALSE by default
- **stringsAsFactors** Convert non-factor string columns into factors? FALSE by default

**Details**

This is useful for finding anomalous observations, row-wise.

It will convert any categorical variables in the data frame into numerics as long as they are factors. For example, in order for a character column to be used as a component in the distance calculations, it must either be a factor, or converted to a factor by using the `stringsAsFactors` parameter.

**Value**

A vector of observation-wise mahalanobis distances.

**See Also**

`insist_rows`

**Examples**

```r
maha_dist(mtcars)
maha_dist(iris, robust=TRUE)

library(magrittr)      # for piping operator
library(dplyr)         # for "everything()" function

# using every column from mtcars, compute mahalanobis distance
# for each observation, and ensure that each distance is within 10 median absolute deviations from the median
mtcars %>%
  insist_rows(maha_dist, within_n_mads(10), everything())
  ## anything here will run
```
not_na

Returns TRUE if value is not NA

Description

This is the inverse of `is.na`. This is a convenience function meant to be used as a predicate in an `assertr` assertion.

Usage

```r
not_na(x, allow.NaN = FALSE)
```

Arguments

- `x`: A R object that supports `is.na` and `is.nan`
- `allow.NaN`: A logical indicating whether NaNs should be allowed (default FALSE)

Value

A vector of the same length that is TRUE when the element is not NA and FALSE otherwise

See Also

- `is.na`
- `is.nan`

Examples

```r
not_na(NA)
not_na(2.8)
not_na("tree")
not_na(c(1, 2, NA, 4))
```

num_row_NAs

Counts number of NAs in each row

Description

This function will return a vector, with the same length as the number of rows of the provided data frame, corresponding to the number of missing values in each row

Usage

```r
num_row_NAs(data, allow.NaN = FALSE)
```
Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>data</td>
<td>A data frame</td>
</tr>
<tr>
<td>allow.NaN</td>
<td>Treat NaN like NA (by counting it). FALSE by default</td>
</tr>
</tbody>
</table>

Value

A vector of number of missing values in each row

See Also

is.na is.nan not_na

Examples

```r
d # using every column from mtcars, make sure there are at most # 2 NAs in each row. If there are any more than two, error out mtcars %>%
  assert_rows(num_row_NAs, within_bounds(0,2), everything())

## anything here will run
```

Description

`print` method for class "assertr_assert_error" This prints the error message and the entire two-column `data.frame` holding the indexes and values of the offending data.

Usage

```r
# S3 method for class 'assertr_assert_error'
print(x, ...)
```

Arguments

- `x` An assertr_assert_error object
- `...` Further arguments passed to or from other methods
success_and_error_functions

See Also

summary.assertr_assert_error

print.assertr_verify_error

Printing assertr's verify errors

Description

'summary' method for class "assertr_verify_error"

Usage

## S3 method for class 'assertr_verify_error'
print(x, ...)

Arguments

x An assertr_verify_error object.
...
Further arguments passed to or from other methods

See Also

summary.assertr_verify_error

success_and_error_functions

Success and error functions

Description

The behavior of functions like assert, assert_rows, insist, insist_rows, verify when the assertion passes or fails is configurable via the success_fun and error_fun parameters, respectively. The success_fun parameter takes a function that takes the data passed to the assertion function as a parameter. You can write your own success handler function, but there are two provided by this package:

• success_continue - just returns the data that was passed into the assertion function
• success_logical - returns TRUE

The error_fun parameter takes a function that takes the data passed to the assertion function as a parameter. You can write your own error handler function, but there are a few provided by this package:

• error_stop - Prints a summary of the errors and halts execution.
success_and_error_functions

- **error_report** - Prints all the information available about the errors in a "tidy" data.frame (including information such as the name of the predicate used, the offending value, etc...) and halts execution.

- **error_append** - Attaches the errors to a special attribute of data and returns the data. This is chiefly to allow assertr errors to be accumulated in a pipeline so that all assertions can have a chance to be checked and so that all the errors can be displayed at the end of the chain.

- **error_return** - Returns the raw object containing all the errors

- **error_df_return** - Returns a "tidy" data.frame containing all the errors, including informations such as the name of the predicate used, the offending value, etc...

- **error_logical** - returns FALSE

- **just_warn** - Prints a summary of the errors but does not halt execution, it just issues a warning.

- **warn_report** - Prints all the information available about the errors but does not halt execution, it just issues a warning.

**Usage**

```r
success_logical(data, ...)
success_continue(data, ...)
error_stop(errors, data = NULL, warn = FALSE, ...)
just_warn(errors, data = NULL)
error_report(errors, data = NULL, warn = FALSE, ...)
warn_report(errors, data = NULL)
error_append(errors, data = NULL)
error_return(errors, data = NULL)
error_df_return(errors, data = NULL)
error_logical(errors, data = NULL, ...)
```

**Arguments**

- `data` A data frame
- `...` Further arguments passed to or from other methods
- `errors` A list of objects of class assertr_errors
- `warn` If TRUE, assertr will issue a warning instead of an error
summary.assertr_assert_error

Summarizing assertr’s assert errors

Description

‘summary’ method for class "assertr_assert_error" This prints the error message and the first five rows of the two-column ‘data.frame’ holding the indexes and values of the offending data.

Usage

```r
## S3 method for class 'assertr_assert_error'
summary(object, ...)
```

Arguments

- `object` An assertr_assert_error object
- `...` Additional arguments affecting the summary produced

See Also

- `print.assertr_assert_error`

summary.assertr_verify_error

Summarizing assertr’s verify errors

Description

‘summary’ method for class "assertr_verify_error"

Usage

```r
## S3 method for class 'assertr_verify_error'
summary(object, ...)
```

Arguments

- `object` An assertr_verify_error object
- `...` Additional arguments affecting the summary produced

See Also

- `print.assertr_verify_error`
verify

Raises error if expression is FALSE anywhere

Description

Meant for use in a data analysis pipeline, this function will just return the data it's supplied if all the logicals in the expression supplied are TRUE. If at least one is FALSE, this function will raise an error, effectively terminating the pipeline early.

Usage

```r
verify(data, expr, success_fun = success_continue, error_fun = error_stop)
```

Arguments

- `data`: A data frame, list, or environment
- `expr`: A logical expression
- `success_fun`: Function to call if assertion passes. Defaults to returning `data`.
- `error_fun`: Function to call if assertion fails. Defaults to printing a summary of all errors.

Details

For examples of possible choices for the `success_fun` and `error_fun` parameters, run `help("success_and_error_functions")`.

Value

By default, the `data` is returned if predicate assertion is TRUE and an error is thrown if not. If a non-default `success_fun` or `error_fun` is used, the return values of these function will be returned.

Note

See `vignette("assertr")` for how to use this in context.

See Also

- `assert`
- `insist`

Examples

```r
verify(mtcars, drat > 2)  # returns mtcars
## Not run:
verify(mtcars, drat > 3)  # produces error
## End(Not run)

library(magrittr)  # for piping operator
```
## not run:
mtcars %>%
  verify(drat > 3) %>%
  # anything here will not run
## End(not run)

mtcars %>%
  verify(nrow(mtcars) > 2)
  # anything here will run

alist <- list(a=c(1,2,3), b=c(4,5,6))
verify(alist, length(a) > 2)
verify(alist, length(a) > 2 && length(b) > 2)
verify(alist, a > 0 & b > 2)

## not run:
alist %>%
  verify(alist, length(a) > 5)
  # nothing here will run
## End(not run)

---

**within_bounds**  
*Creates bounds checking predicate*

### Description
This function returns a predicate function that will take a numeric value or vector and return TRUE if the value(s) is/are within the bounds set. This does not actually check the bounds of anything—it only returns a function that actually does the checking when called with a number. This is a convenience function meant to return a predicate function to be used in an `assertr` assertion.

### Usage
```r
within_bounds(lower_bound, upper_bound, include.lower = TRUE, include.upper = TRUE, allow.na = TRUE)
```

### Arguments
- **lower_bound**: The lowest permitted value
- **upper_bound**: The upper permitted value
- **include.lower**: A logical indicating whether lower bound should be inclusive (default TRUE)
- **include.upper**: A logical indicating whether upper bound should be inclusive (default TRUE)
- **allow.na**: A logical indicating whether NAs (including NaNs) should be permitted (default TRUE)
**Value**

A function that takes numeric value or numeric vector and returns TRUE if the value(s) is/are within the bounds defined by the arguments supplied by `within_bounds` and FALSE otherwise.

**Examples**

```r
predicate <- within_bounds(3, 4)
predicate(pi)

# is equivalent to
within_bounds(3, 4)(pi)

# a correlation coefficient must always be between 0 and 1
coeff <- cor.test(c(1, 2, 3), c(.5, 2.4, 4))[["estimate"]]
within_bounds(0, 1)(coeff)

## check for positive number
positivep <- within_bounds(0, Inf, include.lower=FALSE)

## this is meant to be used as a predicate in an assert statement
assert(mtcars, within_bounds(4, 8), cyl)

## or in a pipeline
library(magrittr)

mtcars %>%
  assert(within_bounds(4, 8), cyl)
```

---

**within_n_mads**

*Return a function to create robust z-score checking predicate*

**Description**

This function takes one argument, the number of median absolute deviations within which to accept a particular data point. This is generally more useful than its sister function `within_n_sds` because it is more robust to the presence of outliers. It is therefore better suited to identify potentially erroneous data points.

**Usage**

```r
within_n_mads(n, ...)
```

**Arguments**

- `n` The number of median absolute deviations from the median within which to accept a datum
- `...` Additional arguments to be passed to `within_bounds`
within_n_sds

Details
As an example, if '2' is passed into this function, this will return a function that takes a vector and figures out the bounds of two median absolute deviations (MADs) from the median. That function will then return a within_bounds function that can then be applied to a single datum. If the datum is within two MADs of the median of the vector given to the function returned by this function, it will return TRUE. If not, FALSE.

This function isn’t meant to be used on its own, although it can. Rather, this function is meant to be used with the insist function to search for potentially erroneous data points in a data set.

Value
A function that takes a vector and returns a within_bounds predicate based on the MAD of that vector.

See Also
within_n_sds

Examples

```r
test.vector <- rnorm(100, mean=100, sd=20)
within.one.mad <- within_n_mads(1)
custom.bounds.checker <- within_one.mad(test.vector)
custom.bounds.checker(105)  # returns TRUE
custom.bounds.checker(40)   # returns FALSE

# same as
within.n_mads(1)(test.vector)(40)  # returns FALSE

within.n_mads(2)(test.vector)(as.numeric(NA))  # returns TRUE
# because, by default, within_bounds() will accept
# NA values. If we want to reject NAs, we have to
# provide extra arguments to this function
within.n_mads(2, allow.na=FALSE)(test.vector)(as.numeric(NA))  # returns FALSE

# or in a pipeline, like this was meant for
library(magrittr)
iris %>%
  insist(within.n_mads(5), Sepal.Length)
```

within_n_sds  
Return a function to create z-score checking predicate
Description

This function takes one argument, the number of standard deviations within which to accept a particular data point.

Usage

within_n_sds(n, ...)

Arguments

- `n`: The number of standard deviations from the mean within which to accept a datum

... Additional arguments to be passed to `within_bounds`

Details

As an example, if '2' is passed into this function, this will return a function that takes a vector and figures out the bounds of two standard deviations from the mean. That function will then return a `within_bounds` function that can then be applied to a single datum. If the datum is within two standard deviations of the mean of the vector given to the function returned by this function, it will return TRUE. If not, FALSE.

This function isn’t meant to be used on its own, although it can. Rather, this function is meant to be used with the `insist` function to search for potentially erroneous data points in a data set.

Value

A function that takes a vector and returns a `within_bounds` predicate based on the standard deviation of that vector.

See Also

`within_n_mads`

Examples

test.vector <- rnorm(100, mean=100, sd=20)

within.one.sd <- within_n_sds(1)
custom.bounds.checker <- within.one.sd(test.vector)
custom.bounds.checker(105)  # returns TRUE
custom.bounds.checker(40)   # returns FALSE

# same as
within.n.sds(1)(test.vector)(40)  # returns FALSE

within.n.sds(2)(test.vector)(as.numeric(NA))  # returns TRUE
# because, by default, within_bounds() will accept
# NA values. If we want to reject NAs, we have to
# provide extra arguments to this function
within.n.sds(2, allow.na=FALSE)(test.vector)(as.numeric(NA))  # returns FALSE
# or in a pipeline, like this was meant for

library(magrittr)

iris %>%
  insist(within_n_sds($), Sepal.Length)
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