Package 'cleanepi'

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Title Clean and Standardize Epidemiological Data

Version 1.1.1

Description Cleaning and standardizing tabular data package, tailored specifically for curating epidemiological data. It streamlines various data cleaning tasks that are typically expected when working with datasets in epidemiology. It returns the processed data in the same format, and generates a comprehensive report detailing the outcomes of each cleaning task.

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BugReports https://github.com/epiverse-trace/cleanepi/issues

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Contents

`add_to_dictionary` *Add an element to the data dictionary*

Description

Add an element to the data dictionary

Usage

```
add_to_dictionary(dictionary, option, value, grp, order = NULL)
```

Arguments

`dictionary`	A `<data.frame>` with the dictionary associated with the input data. This is expected to be compatible with the **matchmaker** package and must contain the following four columns: `options` This column contains the current values used to represent the different groups in the input data frame (required). `values` The values that will be used to replace the current options (required). `grp` The name of the columns where every option belongs to (required). `orders` This defines the user-defined order of different options (optional).
`option`	A `<vector>` of characters with the new options that need to be added to the dictionary.
`value`	A `<vector>` of characters with the values to be used when replacing the new options.
`grp`	A `<vector>` of characters with the name of the column that contains the option of interest.
`order`	A `<vector>` of numeric values with the order of the new option.

Value

A `<data.frame>`. This is the new data dictionary with an additional line that contains the details about the new options.

Examples

```
test <- add_to_dictionary(
  dictionary = readRDS(
    system.file("extdata", "test_dict.RDS", package = "cleanepi")
  ),
  option = "ml",
  value = "male",
  grp = "gender",
  order = NULL
 )
```

add_to_report *Add an element to the report object*

Description

Add an element to the report object

Usage

```
add_to_report(x, key, value)
```

Arguments

x	A <data.frame> or <linelist>
key	A <character> with the name of the cleaning operation
value	The object to add to the report object

Value

The input <data.frame> or <linelist> with an additional element to the report.

Examples

```
# scan through the data
scan_res <- scan_data(
  data = readRDS(system.file("extdata", "test_df.RDS", package = "cleanepi"))
)

# Perform data cleaning
cleaned_data <- clean_data(
  data = readRDS(
    system.file("extdata", "test_df.RDS", package = "cleanepi")
  ),
  to_numeric = list(target_columns = "sex", lang = "en"),
  dictionary = NULL
)

# add the data scanning result to the report
cleaned_data <- add_to_report(
  x = cleaned_data,
  key = "scanning_result",
  value = scan_res
)
```

check_date_sequence *Checks whether the order in a sequence of date events is chronological. order.*

Description

Checks whether a date sequence in a vector of specified columns is in chronological order or not.

Usage

```
check_date_sequence(data, target_columns)
```

Arguments

data	The input <data.frame> or <linelist>
target_columns	A <vector> of column names for events. Users should specify at least 2 column names in the expected order. For example: target_columns = c("date_symptoms_onset", "date_hospitalization", "date_death"). When the input data is a <linelist> object, this parameter can be set to linelist_tags to apply the date sequence checking exclusively to the tagged columns. The date values in the target columns should be in the ISO8601 format, e.g., 2024-12-31. Otherwise, use the standardize_dates() function to standardize the target columns.

Value

The input dataset. When found, the incorrect date sequences will be stored in the report and can be accessed using the print_report() function as shown in the example below.

Examples

```
# import the data
data <- readRDS(system.file("extdata", "test_df.RDS", package = "cleanepi"))

# standardize the date values
data <- data %>%
  standardize_dates(
    target_columns  = c("date_first_pcr_positive_test", "date.of.admission"),
    error_tolerance = 0.4,
    format = NULL,
    timeframe = NULL
  )

# check whether all admission dates come after the test dates
good_date_sequence <- check_date_sequence(
  data = data,
  target_columns = c("date_first_pcr_positive_test", "date.of.admission")
)

# display rows where admission dates do not come after the test dates
```

```
print_report(
  data = good_date_sequence,
  what = "incorrect_date_sequence"
)
```

`check_subject_ids` *Check whether the subject IDs comply with the expected format. When incorrect IDs are found, the function sends a warning and the user can call the* `correct_subject_ids` *function to correct them.*

Description

Check whether the subject IDs comply with the expected format. When incorrect IDs are found, the function sends a warning and the user can call the `correct_subject_ids` function to correct them.

Usage

```
check_subject_ids(
  data,
  target_columns,
  prefix = NULL,
  suffix = NULL,
  range = NULL,
  nchar = NULL
)
```

Arguments

`data`	The input <data.frame> or <linelist>
`target_columns`	A <vector> of column names with the subject ids.
`prefix`	A <character> with the expected prefix used in the subject IDs
`suffix`	A <character> with the expected suffix used in the subject IDs
`range`	A <vector> with the expected range of numbers in the subject IDs
`nchar`	An <integer> that represents the expected number of characters in the subject ids.

Value

The input dataset with a warning if incorrect subject ids were found

Examples

```
data <- readRDS(
  system.file("extdata", "test_df.RDS", package = "cleanepi")
)

# make first and last subject IDs the same
data$study_id[10] <- data$study_id[1]

# set subject ID number 9 to NA
data$study_id[9] <- NA

# detect the incorrect subject ids i.e. IDs that do not have any or both of
# the followings:
# - starts with 'PS',
# - ends with 'P2',
# - has a number within 1 and 100,
# - contains 7 characters.
dat <- check_subject_ids(
  data = data,
  target_columns = "study_id",
  prefix = "PS",
  suffix = "P2",
  range = c(1, 100),
  nchar = 7
)

# display rows with invalid subject ids
print_report(dat, "incorrect_subject_id")
```

---

clean_data *Clean and standardize data*

Description

Cleans up messy data frames by performing several operations. These include among others: cleaning of column names, detecting and removing duplicates, empty records and columns, constant columns, replacing missing values by NA, converting character columns into dates when they contain a certain number of date values, detecting subject IDs with wrong formats, etc.

Usage

```
clean_data(data, ...)
```

Arguments

data	The input <data.frame> or <linelist>
...	A <list> of cleaning operations to be applied on the input data. The acceptable arguments for ... are:

`standardize_column_names` A <list> with the arguments needed to standardize the column names. The elements of this list are the input for the `standardize_column_names` function.

`replace_missing_values` A <list> of parameters to be used when replacing the missing values by NA. The elements of the list are the inputs for the `replace_missing_values` function.

`remove_duplicates` A <list> with the arguments that define the columns and other parameters to be considered when looking for duplicates. They are the input values for the `remove_duplicates` function.

`remove_constants` A <list> with the parameters that define whether to remove constant data or not. The values are the input for the `remove_constants` function.

`standardize_dates` A <list> of parameters that will be used to standardize the date values from the input data. They represent the input values for the `standardize_dates` function.

`standardize_subject_ids` A <list> of parameters that are needed to check the IDs that comply with the expect format. These arguments are the input values of the `check_subject_ids`.

`to_numeric` A <list> with the parameters needed to convert the specified columns into numeric. When provided, the parameters will be the input values for the `convert_to_numeric`.

`dictionary` A <data.frame> that will be used to substitute the current values in the specified columns the those in the dictionary. It is the main argument for the `clean_using_dictionary` function.

`check_date_sequence` A <list> of arguments to be used when determining whether the sequence of date events is respected across all rows of the input data. The value in this list are the input for the `check_date_sequence` function.

Value

The cleaned input data according to the user-specified parameters. This is associated with a data cleaning report that can be accessed using `attr(cleaned_data, "report")`

Examples

```
# Parameters for column names standardization: rename all column names if
# applicable
standardize_column_names <- list(keep = NULL, rename = NULL)

# parameters to remove constant columns, empty rows and columns: remove rows
# and columns with 100% constant data
remove_constants <- list(cutoff = 1)

# Parameters for substituting missing values ("-99") with NA
replace_missing_values <- list(target_columns = NULL, na_strings = "-99")

# Parameters for duplicates removal across all columns
remove_duplicates <- list(target_columns = NULL)
```

```
# Parameters for the conversion of Date columns into "%Y-%m-%d" format
standardize_dates <- list(
  target_columns = NULL,
  error_tolerance = 0.4,
  format = NULL,
  timeframe = as.Date(c("1973-05-29", "2023-05-29")),
  orders = list(
    world_named_months = c("Ybd", "dby"),
    world_digit_months = c("dmy", "Ymd"),
    US_formats = c("Omdy", "YOmd")
  )
)

# Parameters to check whether the subject IDs comply with the expected format
standardize_subject_ids <- list(
  target_columns = "study_id",
  prefix = "PS",
  suffix = "P2",
  range = c(1, 100),
  nchar = 7
)

# convert the 'sex' column into numeric
to_numeric <- list(target_columns = "sex", lang = "en")

# the dictionary-based cleaning will not be performed here
dictionary = NULL

# no need to check if the sequence of date events is correct
check_date_sequence <- NULL

# perform the data cleaning
cleaned_data <- clean_data(
  data = readRDS(
    system.file("extdata", "test_df.RDS", package = "cleanepi")
  ),
  standardize_column_names = standardize_column_names,
  remove_constants = remove_constants,
  replace_missing_values = replace_missing_values,
  remove_duplicates = remove_duplicates,
  standardize_dates = standardize_dates,
  standardize_subject_ids = standardize_subject_ids,
  to_numeric = to_numeric,
  dictionary = NULL,
  check_date_sequence = NULL
)
```

clean_using_dictionary *Perform dictionary-based cleaning*

Description

Perform dictionary-based cleaning

Usage

```
clean_using_dictionary(data, dictionary)
```

Arguments

data	The input <data.frame> or <linelist>
dictionary	A <data.frame> with the dictionary associated with the input data. This is expected to be compatible with the **matchmaker** package and must contain the following four columns: `options` This column contains the current values used to represent the different groups in the input data frame (required). `values` The values that will be used to replace the current options (required). `grp` The name of the columns where every option belongs to (required). `orders` This defines the user-defined order of different options (optional).

Value

A <data.frame> or <linelist> where the target options have been replaced with their corresponding values in the columns specified in the data dictionary.

Examples

```
data <- readRDS(
  system.file("extdata", "messy_data.RDS", package = "cleanepi")
)
dictionary <- readRDS(
  system.file("extdata", "test_dict.RDS", package = "cleanepi")
)

# adding an option that is not defined in the dictionary to the 'gender'
# column
data$gender[2] <- "homme"
cleaned_df <- clean_using_dictionary(
  data = data,
  dictionary = dictionary
)

# print the report
print_report(cleaned_df, "misspelled_values")
```

common_na_strings *Common strings representing missing values*

Description

This vector contains common values of NA (missing) and is intended for use within {cleanepi} functions `replace_missing_values()`. The current list of strings used can be found by printing out `common_na_strings`. It serves as a helpful tool to explore your data for possible missing values. However, I strongly caution against using this to replace `NA` values without meticulously examining the incidence for each case. Please note that `common_na_strings` utilizes `\\` around the "?", ".", and "*" characters to prevent their wildcard

Usage

```
common_na_strings
```

Format

A vector of 35 character strings.

Source

This vector is a combination of `naniar::common_na_strings` (https://github.com/njtierney/naniar/) and other strings found in the literature.

convert_numeric_to_date *Convert numeric to date*

Description

Convert numeric to date

Usage

```
convert_numeric_to_date(data, target_columns, ref_date, forward = TRUE)
```

Arguments

`data`	The input `<data.frame>` or `<linelist>`
`target_columns`	A `<vector>` of columns names to be converted from numeric to date. When the input data is a `<linelist>` object, this parameter can be set to `linelist_tags` to apply the conversion exclusively to the tagged columns.
`ref_date`	A `<Date>` value with reference date. This can also be a character string with the name of the reference column.
`forward`	A `<logical>` to indicate whether the counts started after the reference date (`TRUE`) or not (`FALSE`). The default is `TRUE`.

Value

A <data.frame> or <linelist> where the column of interest are updated

Examples

```
data <- readRDS(system.file("extdata", "test_df1.RDS", package = "cleanepi"))
data <- convert_numeric_to_date(
  data = data,
  target_columns = "recruited_on_day",
  ref_date = as.Date("2022-10-13"),
  forward = TRUE
)
```

---

convert_to_numeric *Convert columns into numeric*

Description

When this function is invoked without specifying the column names to be converted, the target columns are the ones returned by the scan_data() function. Furthermore, it identifies columns where the proportion of numeric values is at least twice the percentage of character values and performs the conversion in them. The function internally makes call of the main function from the **numberize** package.

Usage

```
convert_to_numeric(data, target_columns = NULL, lang = c("en", "fr", "es"))
```

Arguments

data	The input <data.frame> or <linelist>
target_columns	A <vector> of the target column names. When the input data is a <linelist> object, this parameter can be set to linelist_tags to apply the conversion exclusively to the tagged columns. .
lang	A <character> with the text's language. Currently one of "en", "fr", "es".

Value

A <data.frame> or <linelist> wherein all the specified or detected columns have been transformed into numeric format after the conversion process.

Examples

```
data <- readRDS(
  system.file("extdata", "messy_data.RDS", package = "cleanepi")
)

# convert the 'age' column into numeric
dat <- convert_to_numeric(
  data = data,
  target_columns = "age",
  lang = "en"
)

# print the report from this operation
print_report(data = dat, "converted_into_numeric")
```

`correct_misspelled_values`

Correct misspelled values by using approximate string matching techniques to compare them against the expected values.

Description

Correct misspelled values by using approximate string matching techniques to compare them against the expected values.

Usage

```
correct_misspelled_values(
  data,
  target_columns,
  wordlist,
  max_distance = 1,
  confirm = rlang::is_interactive(),
  ...
)
```

Arguments

`data`	The input <data.frame> or <linelist>
`target_columns`	A <vector> of the target column names. When the input data is a <linelist> object, this parameter can be set to `linelist_tags` to apply the fuzzy matching exclusively to the tagged columns.
`wordlist`	A <vector> of characters with the words to match to the detected misspelled values.
`max_distance`	An <integer> for the maximum distance allowed for detecting a spelling mistakes from the `wordlist`. The distance is the generalized Levenshtein edit distance (see `adist()`). Default is 1.

confirm	A <logical> that determines whether to show the user a menu of spelling corrections. If TRUE and using R interactively then the user will have the option to review the proposed spelling corrections. This argument is useful for turning off the menu() when rlang::is_interactive() returns TRUE but not wanting to prompt the user e.g. devtools::run_examples().
...	dots Extra arguments to pass to adist().

Details

When used interactively (see interactive()) the user is presented a menu to ensure that the words detected using approximate string matching are not false positives and the user can decided whether to proceed with the spelling corrections. In non-interactive sessions all misspelled values are replaced by their closest values within the provided vector of expected values.

If multiple words supplied in the wordlist equally match a word in the data and confirm is TRUE the user is presented a menu to choose the replacement word. If it is not used interactively multiple equal matches throws a warning.

Value

The corrected input data according to the user-specified wordlist.

Examples

```
df <- data.frame(
  case_type = c("confirmed", "confermed", "probable", "susspected"),
  outcome = c("died", "recoverd", "did", "recovered")
)
df
correct_misspelled_values(
  data = df,
  target_columns = c("case_type", "outcome"),
  wordlist = c("confirmed", "probable", "suspected", "died", "recovered"),
  confirm = FALSE
)
```

---

correct_subject_ids *Correct the wrong subject IDs based on the user-provided values.*

Description

After detecting incorrect subject IDs from the check_subject_ids() function, use this function to provide the correct IDs and perform the substitution.

Usage

```
correct_subject_ids(data, target_columns, correction_table)
```

Arguments

data	The input <data.frame> or <linelist>
target_columns	A <vector> of column names with the subject ids.
correction_table	A <data.frame> with the following two columns: **from** a column with the wrong subject IDs **to** a column with the values to be used to substitute the incorrect ids.

Value

The input dataset where all subject ids comply with the expected format.

Examples

```
data <- readRDS(
  system.file("extdata", "test_df.RDS", package = "cleanepi")
)
# detect the incorrect subject ids i.e. IDs that do not have any or both of
# the followings:
# - starts with 'PS',
# - ends with 'P2',
# - has a number within 1 and 100,
# - contains 7 characters.
dat <- check_subject_ids(
  data = data,
  target_columns = "study_id",
  prefix = "PS",
  suffix = "P2",
  range = c(1, 100),
  nchar = 7
)

# display rows with invalid subject ids
print_report(dat, "incorrect_subject_id")

# generate the correction table
correction_table <- data.frame(
  from = c("P0005P2", "PB500P2", "PS004P2-1"),
  to = c("PB005P2", "PB050P2", "PS004P2")
)

# perform the correction
dat <- correct_subject_ids(
  data = dat,
  target_columns = "study_id",
  correction_table = correction_table
)
```

`find_duplicates` *Identify and return duplicated rows in a data frame or linelist.*

Description

Identify and return duplicated rows in a data frame or linelist.

Usage

```
find_duplicates(data, target_columns = NULL)
```

Arguments

`data`	The input <data.frame> or <linelist>.
`target_columns`	A <vector> of columns names or indices to consider when looking for duplicates. When the input data is a <linelist> object, this parameter can be set to `linelist_tags` from which duplicates to be removed. Its default value is `NULL`, which considers duplicates across all columns.

Value

A <data.frame> or <linelist> of all duplicated rows with following 2 additional columns:

row_id The indices of the duplicated rows from the input data. Users can choose from these indices, which row they consider as redundant in each group of duplicates.

group_id a unique identifier associated to each group of duplicates.

Examples

```
data <- readRDS(
  system.file("extdata", "test_linelist.RDS", package = "cleanepi")
)

# find duplicates across the following columns: "dt_onset", "dt_report",
# "sex", and "outcome"
dups <- find_duplicates(
  data = data,
  target_columns = c("dt_onset", "dt_report", "sex", "outcome")
)

# print the detected duplicates
print_report(dups, "found_duplicates")
```

get_default_params *Set and return* clean_data *default parameters*

Description

When clean_data() function is called without any argument, these default values provided to the function's arguments will be applied on the input data. By default, operations that require the target columns to be specified by the user will not be performed. The default cleaning operations include: i) standardizing column names, ii) detecting and removing duplicates, and iii) removing constant data.

Usage

```
get_default_params()
```

Value

A <list> of the default cleaning parameters.

Examples

```
default_params <- get_default_params()
```

print_report *Generate report from data cleaning operations*

Description

Generate report from data cleaning operations

Usage

```
print_report(
  data,
  what = NULL,
  print = FALSE,
  report_title = "{cleanepi} data cleaning report",
  output_file_name = NULL,
  format = "html"
)
```

Arguments

data	A <data.frame> or <linelist> object returned from the clean_data or the main functions of each data cleaning module.
what	A <character> with the name of the specific data cleaning report which would be displayed. The possible values are: incorrect_date_sequence To display rows with the incorrect date sequences colnames To display the column names before and after cleaning converted_into_numeric To display the names of the columns that that have been converted into numeric date_standardization To display rows in the cleaned data with date values that are outside of the specified time frame, and rows with date values that comply with multiple formats misspelled_values To display the detected misspelled values removed_duplicates To display the duplicated rows that have been removed found_duplicates To display the duplicated rows constant_data To display the constant data i.e. constant columns, empty rows and columns missing_values_replaced_at To display the names of the columns where the missing value strings have been replaced with NA incorrect_subject_id To display the missing, duplicated and invalid subject subject IDs
print	A <logical> that specifies whether to to open the report in your browser in the form of a HTML file or no. Default is FALSE.
report_title	A <character> with the title that will appear on the report
output_file_name	A <character> used to specify the name of the report file, excluding any file extension. If no file name is supplied, one will be automatically generated with the format cleanepi_report_YYMMDD_HHMMSS.
format	A <character> with the file format of the report. Currently only "html" is supported.

Value

A <character> containing the name and path of the saved report

Examples

```
data <- readRDS(system.file("extdata", "test_df.RDS", package = "cleanepi"))
test_dictionary <- readRDS(
  system.file("extdata", "test_dictionary.RDS", package = "cleanepi")
)

# scan through the data
scan_res <- scan_data(data)

# Perform data cleaning
```

```
cleaned_data <- data %>%
 standardize_column_names(keep = NULL, rename = c("DOB" = "dateOfBirth")) %>%
 replace_missing_values(target_columns = NULL, na_strings = "-99") %>%
 remove_constants(cutoff = 1.0) %>%
 remove_duplicates(target_columns = NULL) %>%
 standardize_dates(
   target_columns = NULL,
   error_tolerance = 0.4,
   format = NULL,
   timeframe = as.Date(c("1973-05-29", "2023-05-29"))
 ) %>%
 check_subject_ids(
   target_columns = "study_id",
   prefix = "PS",
   suffix = "P2",
   range = c(1L, 100L),
   nchar = 7L
 ) %>%
 convert_to_numeric(target_columns = "sex", lang = "en") %>%
 clean_using_dictionary(dictionary = test_dictionary)

# add the data scanning result to the report
cleaned_data <- add_to_report(
  x = cleaned_data,
  key = "scanning_result",
  value = scan_res
)

# save a report in the current directory using the previously-created objects
print_report(
  data = cleaned_data,
  report_title = "{cleanepi} data cleaning report",
  output_file_name = NULL,
  format = "html",
  print = TRUE
)
```

`remove_constants` *Remove constant data, including empty rows, empty columns, and columns with constant values.*

Description

The function iteratively removes constant data until none remain. It records details of the removed constant data as a data frame within the report object.

Usage

```
remove_constants(data, cutoff = 1)
```

Arguments

data	The input <data.frame> or <linelist>
cutoff	A <numeric> value specifying the cut-off for removing constant data. The possible values vary between 0 (excluded) and 1 (included). The default is 1 i.e. remove rows and columns with 100% constant data.

Value

The input dataset where the constant data is filtered out based on specified cut-off.

Examples

```
data <- readRDS(system.file("extdata", "test_df.RDS", package = "cleanepi"))

# introduce an empty column
data$empty_column <- NA
# inject some missing values across some columns
data$study_id[3] = NA_character_
data$date.of.admission[3] = NA_character_
data$date.of.admission[4] = NA_character_
data$dateOfBirth[3] = NA_character_
data$dateOfBirth[4] = NA_character_
data$dateOfBirth[5] = NA_character_

# with cutoff = 1, line 3, 4, and 5 are not removed
cleaned_df <- remove_constants(
  data = data,
  cutoff = 1
)

# drop rows or columns with a percentage of constant values
# equal to or more than 50%
cleaned_df <- remove_constants(
  data = cleaned_df,
  cutoff = 0.5
)

# drop rows or columns with a percentage of constant values
# equal to or more than 25%
cleaned_df <- remove_constants(
  data = cleaned_df,
  cutoff = 0.25
)

# drop rows or columns with a percentage of constant values
# equal to or more than 15%
cleaned_df <- remove_constants(
  data = cleaned_df,
  cutoff = 0.15
)
```

```
# check the report to see what has happened
print_report(cleaned_df, "constant_data")
```

`remove_duplicates` *Remove duplicates*

Description

When removing duplicates, users can specify a set columns to consider with the `target_columns` argument.

Usage

```
remove_duplicates(data, target_columns = NULL)
```

Arguments

`data`	The input <data.frame> or <linelist>.
`target_columns`	A <vector> of column names to use when looking for duplicates. When the input data is a `linelist` object, this parameter can be set to `linelist_tags` if you wish to look for duplicates on tagged columns only. Default is `NULL`.

Details

Caveat: In many epidemiological datasets, multiple rows may share the same value in one or more columns without being true duplicates. For example, several individuals might have the same symptom onset date and admission date. Be cautious when using this function—especially when applying it to a single target column—to avoid incorrect identification or removal of valid entries.

Value

The input data <data.frame> or <linelist> without the duplicated rows identified from all or the specified columns.

Examples

```
data <- readRDS(
  system.file("extdata", "test_linelist.RDS", package = "cleanepi")
)
no_dups <- remove_duplicates(
  data = data,
  target_columns = "linelist_tags"
)

# print the removed duplicates
print_report(no_dups, "removed_duplicates")

# print the detected duplicates
print_report(no_dups, "found_duplicates")
```

replace_missing_values *Replace missing values with* NA

Description

Replace missing values with NA

Usage

```
replace_missing_values(
  data,
  target_columns = NULL,
  na_strings = cleanepi::common_na_strings
)
```

Arguments

data	The input <data.frame> or <linelist>
target_columns	A <vector> of column names. If provided, missing values will be substituted only in the specified columns. When the input data is a <linelist> object, this parameter can be set to linelist_tags to replace missing values with NA in the tagged columns only.
na_strings	A <vector> of characters that represent the missing values in the columns of interest. By default, it utilizes cleanepi::common_na_strings. However, if the missing values string in the columns of interest is not included in this predefined vector, it can be used as the value for this argument.

Value

The input data where missing values are replaced by NA.

Examples

```
data <- readRDS(
  system.file("extdata", "test_df.RDS", package = "cleanepi")
)

# replace all occurrences of '-99' with NA
cleaned_data <- replace_missing_values(
  data = data,
  target_columns = NULL,
  na_strings = "-99"
)

# print the names of the columns where the replacement occurred
print_report(cleaned_data, "missing_values_replaced_at")
```

scan_data *Scan through a data frame and return the proportion of* missing, numeric, Date, character, logical *values.*

Description

The function checks for the existence of character columns in the data. When found, it reports back the proportion of the data types mentioned above in those columns. See the details section to know more about how it works.

Usage

```
scan_data(data, format = "proportion")
```

Arguments

data	The input <data.frame> or <linelist>
format	A <character> with the format in which the output of the data scanning result will be returned. The function returns the proportions of the different data types by default. Other possible values are: percentage and fraction to return the percentage or the fraction of the data types respectively.

Details

How does it work? The <character> columns are identified first. If no <character> columns are found, the function returns a message.

For each <character> column, the function counts:

1. The number of missing values (NA).
2. The number of numeric values. A process is initiated to detect valid dates among these numeric values using lubridate::as_date() and date_guess() functions. If valid dates are found, a warning is triggered to alert about ambiguous numeric values potentially representing dates. **Note:** A date is considered valid if it falls within the range from today's date to 50 years in the past.
3. The detection of <Date> values from non-numeric data using the date_guess() function. The total date count includes dates from today's from both numeric and non-numeric values. Due to overlap, the sum of counts across rows in the scanning result may exceed 1.
4. The count of <logical> values.

Remaining values are categorized as <character>.

Value

A <data.frame> if the input data contains columns of type character. It invisibly returns NA otherwise. The returned data frame will have the same number of rows as the number of character columns, and six columns representing their column names, proportion of missing, numeric, date, character, and logical values.

Examples

```
# scan through a data frame of character columns only
scan_result <- scan_data(
  data = readRDS(
    system.file("extdata", "messy_data.RDS", package = "cleanepi")
  )
)

# scan through a data frame with two character columns
scan_result <- scan_data(
  data = readRDS(system.file("extdata", "test_linelist.RDS",
                             package = "cleanepi"))
)

# scan through a data frame with no character columns
data(iris)
iris[["fct"]] <- as.factor(sample(c("gray", "orange"), nrow(iris),
                          replace = TRUE))
iris[["lgl"]] <- sample(c(TRUE, FALSE), nrow(iris), replace = TRUE)
iris[["date"]] <- as.Date(seq.Date(from = as.Date("2024-01-01"),
                                   to = as.Date("2024-08-30"),
                                   length.out = nrow(iris)))
iris[["posit_ct"]] <- as.POSIXct(iris[["date"]])
scan_result <- scan_data(data = iris)
```

`standardize_column_names` *Standardize column names of a data frame or line list*

Description

All columns names will be reformatted to snake_case. When the conversion to snakecase does not work as expected, use the `keep` and/or `rename` arguments to reformat the column name properly.

Usage

```
standardize_column_names(data, keep = NULL, rename = NULL)
```

Arguments

`data`	The input `<data.frame>` or `<linelist>`.
`keep`	A `<vector>` of column names to maintain as they are. When dealing with a `<linelist>`, this can be set to `linelist_tags`, to maintain the tagged column names. The Default is `NULL`.
`rename`	A named `<vector>` of column names to be renamed. This should be in the form of `c(new_name1 = "old_name1", new_name2 = "old_name2")` for example.

Value

A <data.frame> or <linelist> with easy to work with column names.

Examples

```
# do not rename 'date.of.admission'
cleaned_data <- standardize_column_names(
  data = readRDS(
    system.file("extdata", "test_df.RDS", package = "cleanepi")
  ),
  keep = "date.of.admission"
)

# do not rename 'date.of.admission', but rename 'dateOfBirth' and 'sex' to
# 'DOB' and 'gender' respectively
cleaned_data <- standardize_column_names(
  data = readRDS(
    system.file("extdata", "test_df.RDS", package = "cleanepi")
  ),
  keep = "date.of.admission",
  rename = c(DOB = "dateOfBirth", gender = "sex")
)

# print the report
print_report(
  data = cleaned_data,
  what = "colnames"
)
```

---

standardize_dates *Standardize date variables*

Description

When the format of the values in a column and/or the target columns are not defined, we strongly recommend checking a few converted dates manually to make sure that the dates extracted from a character vector or a factor are correct.

Usage

```
standardize_dates(
  data,
  target_columns = NULL,
  format = NULL,
  timeframe = NULL,
  error_tolerance = 0.4,
 orders = list(world_named_months = c("Ybd", "dby"), world_digit_months = c("dmy",
```

```
    "Ymd"), US_formats = c("Omdy", "YOmd"))
)
```

Arguments

data	The input <data.frame> or <linelist>
target_columns	A <vector> of the target date column names. When the input data is a <linelist> object, this parameter can be set to linelist_tags if you wish to standardize the date columns across tagged columns only. Default is NULL.
format	A <vector> of the expected formats in the date values from the date columns. Default is NULL.
timeframe	A <vector> of 2 values of type <Date>. If provided, date values that do not fall within this timeframe will be set to NA.
error_tolerance	A <numeric> between 0 and 1 indicating the proportion of entries which cannot be identified as dates to be tolerated; if this proportion is exceeded, the original vector is returned, and a message is issued; defaults to 0.4 (40 percent).
orders	A <list> or <vector> of characters with the date codes for fine-grained parsing of dates. This allows for parsing of mixed dates. If a <list> is supplied, that <list> will be used for successive tries in parsing. When this is not provided (orders = NULL), the function will use the following order defined in the guesser:

```
list(
  quarter_partial_dates = c("Y", "Ym", "Yq"),
 world_digit_months = c("Yq", "ymd", "ydm", "dmy", "mdy", "myd", "dym",
                        "Ymd", "Ydm", "dmY", "mdY", "mYd", "dYm"),
 world_named_months = c("dby", "dyb", "bdy", "byd", "ybd", "ydb",
                        "dbY", "dYb", "bdY", "bYd", "Ybd", "Ydb"),
  us_format = c("Omdy", "YOmd")
)
```

Details

Check for the presence of date values that could have multiple formats from the $multi_format_dates element of the report.

Converting ambiguous character strings to dates is difficult for many reasons:

- dates may not use the standard Ymd format
- within the same variable, dates may follow different formats
- dates may be mixed with things that are not dates
- the behavior of as.Date in the presence of non-date is hard to predict, sometimes returning NA, sometimes issuing an error.

This function tries to address all the above issues. Dates with the following format should be automatically detected, irrespective of separators (e.g. "-", " ", "/") and surrounding text:

- "19 09 2018"
- "2018 09 19"
- "19 Sep 2018"
- "2018 Sep 19"
- "Sep 19 2018"

How it works:

This function relies heavily on `lubridate::parse_date_time()`, which is an extremely flexible date parser that works well for consistent date formats, but can quickly become unwieldy and may produce spurious results. `standardize_dates()` will use a list of formats in the `orders` argument to run `parse_date_time()` with each format vector separately and take the first correctly parsed date from all the trials.

With the default orders shown above, the dates 03 Jan 2018, 07/03/1982, and 08/20/85 are correctly interpreted as 2018-01-03, 1982-03-07, and 1985-08-20. The examples section will show how you can manipulate the `orders` to be customized for your situation.

Value

The input dataset where the date columns have been standardized. The date values that are out of the specified timeframe will be reported in the report. Similarly, date values that comply with multiple formats will also be featured in the report object.

Examples

```
x <- c("03 Jan 2018", "07/03/1982", "08/20/85")
# The below will coerce values where the month is written in letters only
# into Date.
as.Date(lubridate::parse_date_time(x, orders = c("Ybd", "dby")))

# coerce values where the month is written in letters or numbers into Date.
as.Date(lubridate::parse_date_time(x, orders = c("dmy", "Ymd")))

# How to use standardize_dates()
data <- readRDS(system.file("extdata", "test_df.RDS", package = "cleanepi"))

# convert values in the 'date.of.admission' column into "%Y-%m-%d"
# format
dat <- standardize_dates(
  data = data,
  target_columns = "date.of.admission",
  format = NULL,
  timeframe = as.Date(c("2021-01-01", "2021-12-01")),
  error_tolerance = 0.4,
  orders = list(
    world_named_months = c("Ybd", "dby"),
    world_digit_months = c("dmy", "Ymd"),
    US_format = c("Omdy", "YOmd")
  )
)
```

```
# print the report
print_report(dat, "date_standardization")
```

`timespan` *Calculate time span between dates*

Description

Calculate time span between dates

Usage

```
timespan(
  data,
  target_column = NULL,
  end_date = Sys.Date(),
  span_unit = c("years", "months", "weeks", "days"),
  span_column_name = "span",
  span_remainder_unit = NULL
)
```

Arguments

`data`	The input <data.frame> or <linelist>
`target_column`	A <vector> of character used to specify the name of the date column of interest. The values in this column should be of type <Date> in ISO8601 format, e.g., 2024-01-31.
`end_date`	The end date. It can be either a <character> that is the name of another column of type <Date> from the input data or a <vector> of Dates or a single <Date> value. This should also be in the ISO8601 format, e.g., 2024-01-31. Default is today's date `Sys.Date()`.
`span_unit`	A <character> that specifies the units in which the time span between the dates will be returned. The possible units are: 'years', 'months', 'weeks' or 'days'.
`span_column_name`	A <character> that specifies the name of the new column to be used to store the calculated time span in the input data frame.
`span_remainder_unit`	A <character> that specifies the unit in which the remainder of the time span should be calculated. May be one of "months", "weeks", and "days". Remainders requested in the same unit as the age will return values of 0. Default is `NULL` for decimal time span.

Value

The input <data.frame> with one or two additional columns:

span or any other name chosen by the user. This will contain the calculated time span in the desired units.

"*_remainder" a column with the number of the remaining days or weeks or months depending on the value of the 'span_remainder_unit' parameter. The star represents here the value of the 'span_column_name' argument.

Examples

```
# In the below example, this function is used to calculate patient's age from
# their dates of birth

# import the data, replace missing values with NA and convert date into ISO
# format
data <- readRDS(system.file("extdata", "test_df.RDS", package = "cleanepi"))
data <- data %>%
  replace_missing_values(target_columns = "dateOfBirth",
                         na_strings = "-99") %>%
  standardize_dates(target_columns = "dateOfBirth",
                    error_tolerance = 0.0)

# calculate the age in 'years' and return the remainder in 'months'
age <- timespan(
  data = data,
  target_column = "dateOfBirth",
  end_date = Sys.Date(),
  span_unit = "years",
  span_column_name = "age_in_years",
  span_remainder_unit = "months"
)
```

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