

Package ‘admiral’

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Type Package

Title ADaM in R Asset Library

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Description A toolbox for programming Clinical Data Interchange Standards Consortium (CDISC) compliant Analysis Data Model (ADaM) datasets in R. ADaM datasets are a mandatory part of any New Drug or Biologics License Application submitted to the United States Food and Drug Administration (FDA). Analysis derivations are implemented in accordance with the “Analysis Data Model Implementation Guide” (CDISC Analysis Data Model Team, 2021, <<https://www.cdisc.org/standards/foundational/adam/adamig-v1-3-release-package>>).

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admiral_adlb	<i>Lab Analysis Dataset</i>
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Description

An example of lab analysis dataset

Usage

admiral_adlb

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 3779 rows and 111 columns.

Source

Derived from the `adlb` template, then further filtered due to dataset size by the following USUB-JIDs: 01-701-1015, 01-701-1023, 01-701-1028, 01-701-1033, 01-701-1034, 01-701-1047, 01-701-1097, 01-705-1186, 01-705-1292, 01-705-1310, 01-708-1286

See Also

Other datasets: [admiral_ads1](#), [ex_single](#), [example_qs](#), [queries_mh](#), [queries](#)

admiral_ads1	<i>Subject Level Analysis Dataset</i>
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Description

An example subject level analysis dataset

Usage

```
admiral_ads1
```

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 306 rows and 50 columns.

Source

Derived from the `dm` and `ds` datasets using `{admiral}` (https://github.com/pharmaverse/admiral/blob/main/inst/templates/ad_ads1.R)

See Also

Other datasets: [admiral_adlb](#), [ex_single](#), [example_qs](#), [queries_mh](#), [queries](#)

`assert_db_requirements`*Check required parameters for a basket*

Description

If a basket (SMQ, SDG, ...) are requested, the version and a function to access the database must be provided. The function checks these requirements.

Usage

```
assert_db_requirements(  
    version,  
    version_arg_name,  
    fun,  
    fun_arg_name,  
    queries,  
    i  
)
```

Arguments

<code>version</code>	Version provided by user
<code>version_arg_name</code>	Name of the argument providing the version
<code>fun</code>	Function provided by user
<code>fun_arg_name</code>	Name of the argument providing the function
<code>queries</code>	Queries provide by user
<code>i</code>	Index of query being checked

Value

An error is issued if `version` or `fun` is null.

See Also

Other Advanced Functions: [assert_terms\(\)](#), [assert_valid_queries\(\)](#), [extend_source_datasets\(\)](#), [filter_date_sources\(\)](#), [format.basket_select\(\)](#), [list_tte_source_objects\(\)](#), [params\(\)](#), [validate_basket_select\(\)](#), [validate_query\(\)](#)

 assert_terms

Asserts Requirements for Terms for Queries

Description

The function checks the requirements for terms for queries provided by the user. The terms could have been provided directly in the query definition or via a user provided function for accessing a SMQ or SDG database.

Usage

```
assert_terms(
  terms,
  expect_query_name = FALSE,
  expect_query_id = FALSE,
  source_text
)
```

Arguments

terms	Terms provided by user
expect_query_name	Is the QUERY_NAME column expected?
expect_query_id	Is the QUERY_ID column expected?
source_text	Text describing the source of the terms, e.g., "the data frame provided for the definition element".

Value

An error is issued if

- terms is not a data frame,
- terms has zero observations,
- the TERM_LEVEL variable is not in terms,
- neither the TERM_NAME nor the TERM_ID variable is in terms,
- expect_query_name == TRUE and the QUERY_NAME variable is not in terms,
- expect_query_id == TRUE and the QUERY_ID variable is not in terms,

See Also

[create_query_data\(\)](#), [query\(\)](#)

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_valid_queries\(\)](#), [extend_source_datasets\(\)](#), [filter_date_sources\(\)](#), [format.basket_select\(\)](#), [list_tte_source_objects\(\)](#), [params\(\)](#), [validate_basket_select\(\)](#), [validate_query\(\)](#)

Examples

```
try(
  assert_terms(
    terms = 42,
    source_text = "object provided by the `definition` element"
  )
)
```

assert_valid_queries *Verify if a Dataset Has the Required Format as Queries Dataset.*

Description

Verify if a Dataset Has the Required Format as Queries Dataset.

Usage

```
assert_valid_queries(queries, queries_name)
```

Arguments

queries A data.frame.
queries_name Name of the queries dataset, a string.

Details

Check if the dataset has the following columns

- VAR_PREFIX, e.g., SMQ01, CQ12
- QUERY_NAME, non NA, must be unique per each VAR_PREFIX
- QUERY_ID, could be NA, must be unique per each VAR_PREFIX
- QUERY_SCOPE, 'BROAD', 'NARROW', or NA
- QUERY_SCOPE_NUM, 1, 2, or NA
- TERM_LEVEL, e.g., "AEDECOD", "AELLT", "AELLTCD", ...
- TERM_NAME, character, could be NA only at those observations where TERM_ID is non-NA
- TERM_ID, integer, could be NA only at those observations where TERM_NAME is non-NA

Value

The function throws an error if any of the requirements not met.

See Also

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_terms\(\)](#), [extend_source_datasets\(\)](#), [filter_date_sources\(\)](#), [format.basket_select\(\)](#), [list_tte_source_objects\(\)](#), [params\(\)](#), [validate_basket_select\(\)](#), [validate_query\(\)](#)

Examples

```
data("queries")
assert_valid_queries(queries, "queries")
```

atoxgr_criteria_ctcv4 *Metadata Holding Grading Criteria for NCI-CTCAEv4*

Description

Metadata Holding Grading Criteria for NCI-CTCAEv4

Usage

```
atoxgr_criteria_ctcv4
```

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 40 rows and 13 columns.

Details

This metadata has its origin in the ADLB Grading Spec Excel file which ships with {admiral} and can be accessed using `system.file("adlb_grading/adlb_grading_spec.xlsx", package = "admiral")` in sheet = "NCICTCAEv4". The dataset contained in there has the following columns:

- SOC: variable to hold the SOC of the lab test criteria.
- TERM: variable to hold the term describing the criteria applied to a particular lab test, eg. 'Anemia' or 'INR Increased'. Note: the variable is case insensitive.
- Grade 1: Criteria defining lab value as Grade 1.
- Grade 2: Criteria defining lab value as Grade 2.
- Grade 3: Criteria defining lab value as Grade 3.
- Grade 4: Criteria defining lab value as Grade 4.
- Grade 5: Criteria defining lab value as Grade 5.
- Definition: Holds the definition of the lab test abnormality.
- GRADE_CRITERIA_CODE: variable to hold code that creates grade based on defined criteria.
- SI_UNIT_CHECK: variable to hold unit of particular lab test. Used to check against input data if criteria is based on absolute values.
- VAR_CHECK: List of variables required to implement lab grade criteria. Use to check against input data.
- DIRECTION: variable to hold the direction of the abnormality of a particular lab test value. 'L' is for LOW values, 'H' is for HIGH values. Note: the variable is case insensitive.
- COMMENT: Holds any information regarding rationale behind implementation of grading criteria.

Note: Variables SOC, TERM, Grade 1, Grade 2, Grade 3, Grade 4, Grade 5, Definition are from the source document on NCI-CTC website defining the grading criteria. **Common Terminology Criteria for Adverse Events (CTCAE)v4.0** From these variables only 'TERM' is used in the admiral code, the rest are for information and traceability only.

See Also

Other metadata: [atoxgr_criteria_ctcv5](#), [dose_freq_lookup](#)

atoxgr_criteria_ctcv5 *Metadata Holding Grading Criteria for NCI-CTCAEv5*

Description

Metadata Holding Grading Criteria for NCI-CTCAEv5

Usage

```
atoxgr_criteria_ctcv5
```

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 37 rows and 13 columns.

Details

This metadata has its origin in the ADLB Grading Spec Excel file which ships with {admiral} and can be accessed using `system.file("adlb_grading/adlb_grading_spec.xlsx", package = "admiral")` in sheet = "NCICTCAEv5". The dataset contained in there has the following columns:

- SOC: variable to hold the SOC of the lab test criteria.
- TERM: variable to hold the term describing the criteria applied to a particular lab test, eg. 'Anemia' or 'INR Increased'. Note: the variable is case insensitive.
- Grade 1: Criteria defining lab value as Grade 1.
- Grade 2: Criteria defining lab value as Grade 2.
- Grade 3: Criteria defining lab value as Grade 3.
- Grade 4: Criteria defining lab value as Grade 4.
- Grade 5: Criteria defining lab value as Grade 5.
- Definition: Holds the definition of the lab test abnormality.
- GRADE_CRITERIA_CODE: variable to hold code that creates grade based on defined criteria.
- SI_UNIT_CHECK: variable to hold unit of particular lab test. Used to check against input data if criteria is based on absolute values.
- VAR_CHECK: List of variables required to implement lab grade criteria. Use to check against input data.
- DIRECTION: variable to hold the direction of the abnormality of a particular lab test value. 'L' is for LOW values, 'H' is for HIGH values. Note: the variable is case insensitive.
- COMMENT: Holds any information regarding rationale behind implementation of grading criteria.

Note: Variables SOC, TERM, Grade 1, Grade 2, Grade 3, Grade 4, Grade 5, Definition are from the source document on NCI-CTC website defining the grading criteria. **Common Terminology Criteria for Adverse Events (CTCAE)v5.0** From these variables only 'TERM' is used in the admiral code, the rest are for information and traceability only.

See Also

Other metadata: [atoxgr_criteria_ctcv4](#), [dose_freq_lookup](#)

basket_select	<i>Create a basket_select object</i>
---------------	--------------------------------------

Description

Create a basket_select object

Usage

```
basket_select(name = NULL, id = NULL, scope = NULL, type)
```

Arguments

name	Name of the query used to select the definition of the query from the company database.
id	Identifier of the query used to select the definition of the query from the company database.
scope	Scope of the query used to select the definition of the query from the company database. <i>Permitted Values:</i> "BROAD", "NARROW", NA_character_
type	The type argument expects a character scalar. It is passed to the company specific <code>get_terms()</code> function such that the function can determine which sort of basket is requested

Details

Exactly one of name or id must be specified.

Value

An object of class basket_select.

See Also

[create_query_data\(\)](#), [query\(\)](#)

Source Objects: [censor_source\(\)](#), [date_source\(\)](#), [death_event](#), [dthcaus_source\(\)](#), [event_source\(\)](#), [event\(\)](#), [query\(\)](#), [tte_source\(\)](#)

call_derivation	<i>Call a Single Derivation Multiple Times</i>
-----------------	--

Description

Call a single derivation multiple times with some parameters/arguments being fixed across iterations and others varying.

Usage

```
call_derivation(dataset = NULL, derivation, variable_params, ...)
```

Arguments

dataset	The input dataset
derivation	The derivation function to call A function that performs a specific derivation is expected. A derivation adds variables or observations to a dataset. The first argument of a derivation must expect a dataset and the derivation must return a dataset. The function must provide the dataset argument and all arguments specified in the <code>params()</code> objects passed to the <code>variable_params</code> and <code>...</code> argument. Please note that it is not possible to specify <code>{dplyr}</code> functions like <code>mutate()</code> or <code>summarize()</code> .
variable_params	A list of function arguments that are different across iterations. Each set of function arguments must be created using <code>params()</code> .
...	Any number of <i>named</i> function arguments that stay the same across iterations. If a function argument is specified both inside <code>variable_params</code> and <code>...</code> then the value in <code>variable_params</code> overwrites the one in <code>...</code>

Value

The input dataset with additional records/variables added depending on which derivation has been used.

See Also

[params\(\)](#)

Higher Order Functions: [derivation_slice\(\)](#), [restrict_derivation\(\)](#), [slice_derivation\(\)](#)

Examples

```
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data(admiral_ae)
data(admiral_adsl)
```

```

adae <-
  select(admiral_ae[sample(1:nrow(admiral_ae), 1000), ], USUBJID, AESTDTC, AEENDTC) %>%
  derive_vars_merged(
    dataset_add = admiral_ads1,
    new_vars = exprs(TRTSDT, TRTEDT),
    by_vars = exprs(USUBJID)
  )

## While `derive_vars_dt()` can only add one variable at a time, using `call_derivation()`
## one can add multiple variables in one go
call_derivation(
  dataset = adae,
  derivation = derive_vars_dt,
  variable_params = list(
    params(dtc = AESTDTC, date_imputation = "first", new_vars_prefix = "AST"),
    params(dtc = AEENDTC, date_imputation = "last", new_vars_prefix = "AEN")
  ),
  min_dates = exprs(TRTSDT),
  max_dates = exprs(TRTEDT)
)

## The above call using `call_derivation()` is equivalent to the following
adae %>%
  derive_vars_dt(
    new_vars_prefix = "AST",
    dtc = AESTDTC,
    date_imputation = "first",
    min_dates = exprs(TRTSDT),
    max_dates = exprs(TRTEDT)
  ) %>%
  derive_vars_dt(
    new_vars_prefix = "AEN",
    dtc = AEENDTC,
    date_imputation = "last",
    min_dates = exprs(TRTSDT),
    max_dates = exprs(TRTEDT)
  )

```

call_user_fun

Calls a Function Provided by the User

Description

Calls a function provided by the user and adds the function call to the error message if the call fails.

Usage

```
call_user_fun(call)
```

Arguments

call Call to be executed

Value

The return value of the function call

See Also

Utilities used within Derivation functions: [extract_unit\(\)](#), [get_not_mapped\(\)](#), [signal_duplicate_records\(\)](#)

Examples

```
call_user_fun(compute_bmi(
  height = 172,
  weight = 60
))

try(call_user_fun(compute_bmi(
  height = 172,
  weight = "hallo"
)))
```

censor_source *Create a censor_source Object*

Description

censor_source objects are used to define censorings as input for the `derive_param_tte()` function.

Usage

```
censor_source(
  dataset_name,
  filter = NULL,
  date,
  censor = 1,
  set_values_to = NULL
)
```

Arguments

dataset_name The name of the source dataset
The name refers to the dataset provided by the `source_datasets` parameter of `derive_param_tte()`.

filter An unquoted condition for selecting the observations from dataset which are events or possible censoring time points.

date	A variable providing the date of the event or censoring. A date, or a datetime can be specified. An unquoted symbol is expected. Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object.
censor	Censoring value CDISC strongly recommends using 0 for events and positive integers for censoring.
set_values_to	A named list returned by <code>exprs()</code> defining the variables to be set for the event or censoring, e.g. <code>exprs(EVENTDESC = "DEATH", SRCDOM = "ADSL", SRCVAR = "DTHDT")</code> . The values must be a symbol, a character string, a numeric value, or NA.

Value

An object of class `censor_source`, inheriting from class `tte_source`

See Also

[derive_param_tte\(\)](#), [event_source\(\)](#)

Source Objects: [basket_select\(\)](#), [date_source\(\)](#), [death_event](#), [dthcaus_source\(\)](#), [event_source\(\)](#), [event\(\)](#), [query\(\)](#), [tte_source\(\)](#)

Examples

```
# Last study date known alive censor

censor_source(
  dataset_name = "adsl",
  date = LSTALVDT,
  set_values_to = exprs(
    EVNTDESC = "ALIVE",
    SRCDOM = "ADSL",
    SRCVAR = "LSTALVDT"
  )
)
```

 chr2vars

Turn a Character Vector into a List of Expressions

Description

Turn a character vector into a list of expressions

Usage

```
chr2vars(chr)
```

Arguments

chr A character vector

Value

A list of expressions as returned by [exprs\(\)](#)

See Also

Other utils_quo: [negate_vars\(\)](#)

Examples

```
chr2vars(c("USUBJID", "AVAL"))
```

compute_bmi

Compute Body Mass Index (BMI)

Description

Computes BMI from height and weight

Usage

```
compute_bmi(height, weight)
```

Arguments

height HEIGHT value
It is expected that HEIGHT is in cm.
Permitted Values: numeric vector

weight WEIGHT value
It is expected that WEIGHT is in kg.
Permitted Values: numeric vector

Details

Usually this computation function can not be used with %>%.

Value

The BMI (Body Mass Index Area) in kg/m².

See Also

BDS-Findings Functions that returns a vector: [compute_bsa\(\)](#), [compute_framingham\(\)](#), [compute_map\(\)](#), [compute_qtc\(\)](#), [compute_qual_imputation_dec\(\)](#), [compute_qual_imputation\(\)](#), [compute_rr\(\)](#), [compute_scale\(\)](#)

Examples

```
compute_bmi(height = 170, weight = 75)
```

compute_bsa	<i>Compute Body Surface Area (BSA)</i>
-------------	--

Description

Computes BSA from height and weight making use of the specified derivation method

Usage

```
compute_bsa(height = height, weight = weight, method)
```

Arguments

height	HEIGHT value It is expected that HEIGHT is in cm. Permitted Values: numeric vector
weight	WEIGHT value It is expected that WEIGHT is in kg. Permitted Values: numeric vector
method	Derivation method to use: Mosteller: $\sqrt{\text{height} * \text{weight} / 3600}$ DuBois-DuBois: $0.20247 * (\text{height}/100) ^ 0.725 * \text{weight} ^ 0.425$ Haycock: $0.024265 * \text{height} ^ 0.3964 * \text{weight} ^ 0.5378$ Gehan-George: $0.0235 * \text{height} ^ 0.42246 * \text{weight} ^ 0.51456$ Boyd: $0.0003207 * (\text{height} ^ 0.3) * (1000 * \text{weight}) ^ (0.7285 - (0.0188 * \log_{10}(1000 * \text{weight})))$ Fujimoto: $0.008883 * \text{height} ^ 0.663 * \text{weight} ^ 0.444$ Takahira: $0.007241 * \text{height} ^ 0.725 * \text{weight} ^ 0.425$ Permitted Values: character value

Details

Usually this computation function can not be used with %>%.

Value

The BSA (Body Surface Area) in m².

See Also

BDS-Findings Functions that returns a vector: [compute_bmi\(\)](#), [compute_framingham\(\)](#), [compute_map\(\)](#), [compute_qtc\(\)](#), [compute_qual_imputation_dec\(\)](#), [compute_qual_imputation\(\)](#), [compute_rr\(\)](#), [compute_scale\(\)](#)

Examples

```
# Derive BSA by the Mosteller method
compute_bsa(
  height = 170,
  weight = 75,
  method = "Mosteller"
)

# Derive BSA by the DuBois & DuBois method
compute_bsa(
  height = c(170, 185),
  weight = c(75, 90),
  method = "DuBois-DuBois"
)
```

compute_dtf

Derive the Date Imputation Flag

Description

Derive the date imputation flag ('--DTF') comparing a date character vector ('--DTC') with a Date vector ('--DT').

Usage

```
compute_dtf(dtc, dt)
```

Arguments

dtc	The date character vector ('--DTC'). A character date is expected in a format like yyyy-mm-ddThh:mm:ss (partial or complete).
dt	The Date vector to compare. A date object is expected.

Details

Usually this computation function can not be used with %>%.

Value

The date imputation flag ('--DTF') (character value of 'D', 'M', 'Y' or NA)

See Also

Date/Time Computation Functions that returns a vector: [compute_duration\(\)](#), [compute_tmf\(\)](#), [convert_date_to_dtm\(\)](#), [convert_dtc_to_dtm\(\)](#), [convert_dtc_to_dt\(\)](#), [impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Examples

```
compute_dtf(dtc = "2019-07", dt = as.Date("2019-07-18"))
compute_dtf(dtc = "2019", dt = as.Date("2019-07-18"))
```

compute_duration	<i>Compute Duration</i>
------------------	-------------------------

Description

Compute duration between two dates, e.g., duration of an adverse event, relative day, age, ...

Usage

```
compute_duration(
  start_date,
  end_date,
  in_unit = "days",
  out_unit = "days",
  floor_in = TRUE,
  add_one = TRUE,
  trunc_out = FALSE
)
```

Arguments

start_date	The start date A date or date-time object is expected. Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object. Refer to <code>convert_dtc_to_dt()</code> to obtain a vector of imputed dates.
end_date	The end date A date or date-time object is expected. Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object. Refer to <code>convert_dtc_to_dt()</code> to obtain a vector of imputed dates.
in_unit	Input unit See <code>floor_in</code> and <code>add_one</code> parameter for details. Default: 'days' Permitted Values: 'years', 'months', 'days', 'hours', 'minutes', 'min', 'seconds', 'sec'
out_unit	Output unit The duration is derived in the specified unit Default: 'days' Permitted Values: 'years', 'months', 'weeks', 'days', 'hours', 'minutes', 'min', 'seconds', 'sec'

floor_in	<p>Round down input dates?</p> <p>The input dates are round down with respect to the input unit, e.g., if the input unit is 'days', the time of the input dates is ignored.</p> <p>Default: 'TRUE'</p> <p>Permitted Values: TRUE, FALSE</p>
add_one	<p>Add one input unit?</p> <p>If the duration is non-negative, one input unit is added. i.e., the duration can not be zero.</p> <p>Default: TRUE</p> <p>Permitted Values: TRUE, FALSE</p>
trunc_out	<p>Return integer part</p> <p>The fractional part of the duration (in output unit) is removed, i.e., the integer part is returned.</p> <p>Default: FALSE</p> <p>Permitted Values: TRUE, FALSE</p>

Details

The output is a numeric vector providing the duration as time from start to end date in the specified unit. If the end date is before the start date, the duration is negative.

Value

The duration between the two date in the specified unit

See Also

Date/Time Computation Functions that returns a vector: [compute_dtf\(\)](#), [compute_tmf\(\)](#), [convert_date_to_dtm\(\)](#), [convert_dtc_to_dtm\(\)](#), [convert_dtc_to_dt\(\)](#), [impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Examples

```
library(lubridate)

# Derive duration in days (integer), i.e., relative day
compute_duration(
  start_date = ymd_hms("2020-12-06T15:00:00"),
  end_date = ymd_hms("2020-12-24T08:15:00")
)

# Derive duration in days (float)
compute_duration(
  start_date = ymd_hms("2020-12-06T15:00:00"),
  end_date = ymd_hms("2020-12-24T08:15:00"),
  floor_in = FALSE,
  add_one = FALSE
)

# Derive age in years
```

```

compute_duration(
  start_date = ymd("1984-09-06"),
  end_date = ymd("2020-02-24"),
  trunc_out = TRUE,
  out_unit = "years",
  add_one = FALSE
)

# Derive duration in hours
compute_duration(
  start_date = ymd_hms("2020-12-06T9:00:00"),
  end_date = ymd_hms("2020-12-06T13:30:00"),
  out_unit = "hours",
  floor_in = FALSE,
  add_one = FALSE,
)

```

compute_framingham	<i>Compute Framingham Heart Study Cardiovascular Disease 10-Year Risk Score</i>
--------------------	---

Description

Computes Framingham Heart Study Cardiovascular Disease 10-Year Risk Score (FCVD101) based on systolic blood pressure, total serum cholesterol (mg/dL), HDL serum cholesterol (mg/dL), sex, smoking status, diabetic status, and treated for hypertension flag.

Usage

```
compute_framingham(sysbp, chol, cholhdl, age, sex, smokefl, diabetfl, trthypfl)
```

Arguments

sysbp	Systolic blood pressure A numeric vector is expected.
chol	Total serum cholesterol (mg/dL) A numeric vector is expected.
cholhdl	HDL serum cholesterol (mg/dL) A numeric vector is expected.
age	Age (years) A numeric vector is expected.
sex	Gender A character vector is expected. Expected Values: 'M' 'F'
smokefl	Smoking Status A character vector is expected. Expected Values: 'Y' 'N'

diabetfl Diabetic Status
 A character vector is expected. Expected Values: 'Y' 'N'

trthypfl Treated for hypertension status
 A character vector is expected. Expected Values: 'Y' 'N'

Details

The predicted probability of having cardiovascular disease (CVD) within 10-years according to Framingham formula. See AHA Journal article General Cardiovascular Risk Profile for Use in Primary Care for reference.

For Women:

	Factor	Amount
	Age	2.32888
	Total Chol	1.20904
	HDL Chol	-0.70833
	Sys BP	2.76157
	Sys BP + Hypertension Meds	2.82263
	Smoker	0.52873
	Non-Smoker	0
	Diabetic	0.69154
	Not Diabetic	0
	Average Risk	26.1931
	Risk Period	0.95012

For Men:

	Factor	Amount
	Age	3.06117
	Total Chol	1.12370
	HDL Chol	-0.93263
	Sys BP	1.93303
	Sys BP + Hypertension Meds	2.99881
	Smoker	.65451
	Non-Smoker	0
	Diabetic	0.57367
	Not Diabetic	0
	Average Risk	23.9802
	Risk Period	0.88936

The equation for calculating risk:

$$RiskFactors = (\log(Age)*AgeFactor) + (\log(TotalChol)*TotalCholFactor) + (\log(CholHDL)*CholHDLFactor)$$

$$Risk = 100 * (1 - RiskPeriodFactor^{exp(RiskFactors)})$$

Value

A numeric vector of Framingham values

See Also

[derive_param_framingham\(\)](#)

BDS-Findings Functions that returns a vector: [compute_bmi\(\)](#), [compute_bsa\(\)](#), [compute_map\(\)](#), [compute_qtc\(\)](#), [compute_qual_imputation_dec\(\)](#), [compute_qual_imputation\(\)](#), [compute_rr\(\)](#), [compute_scale\(\)](#)

Examples

```
compute_framingham(
  sysbp = 133, chol = 216.16, cholhdl = 54.91, age = 53,
  sex = "M", smokefl = "N", diabetfl = "N", trthypfl = "N"
)
```

```
compute_framingham(
  sysbp = 161, chol = 186.39, cholhdl = 64.19, age = 52,
  sex = "F", smokefl = "Y", diabetfl = "N", trthypfl = "Y"
)
```

compute_map

Compute Mean Arterial Pressure (MAP)

Description

Computes mean arterial pressure (MAP) based on diastolic and systolic blood pressure. Optionally heart rate can be used as well.

Usage

```
compute_map(diabp, sysbp, hr = NULL)
```

Arguments

diabp	Diastolic blood pressure A numeric vector is expected.
sysbp	Systolic blood pressure A numeric vector is expected.
hr	Heart rate A numeric vector or NULL is expected.

Details

$$\frac{2DIABP + SYSBP}{3}$$

if it is based on diastolic and systolic blood pressure and

$$DIABP + 0.01e^{4.14 - \frac{40.74}{HR}} (SYSBP - DIABP)$$

if it is based on diastolic, systolic blood pressure, and heart rate.

Usually this computation function can not be used with %>%.

Value

A numeric vector of MAP values

See Also

BDS-Findings Functions that returns a vector: [compute_bmi\(\)](#), [compute_bsa\(\)](#), [compute_framingham\(\)](#), [compute_qtc\(\)](#), [compute_qual_imputation_dec\(\)](#), [compute_qual_imputation\(\)](#), [compute_rr\(\)](#), [compute_scale\(\)](#)

Examples

```
# Compute MAP based on diastolic and systolic blood pressure
compute_map(diabp = 51, sysbp = 121)

# Compute MAP based on diastolic and systolic blood pressure and heart rate
compute_map(diabp = 51, sysbp = 121, hr = 59)
```

compute_qtc

Compute Corrected QT

Description

Computes corrected QT using Bazett's, Fridericia's or Sagie's formula.

Usage

```
compute_qtc(qt, rr, method)
```

Arguments

qt	QT interval A numeric vector is expected. It is expected that QT is measured in msec.
rr	RR interval A numeric vector is expected. It is expected that RR is measured in msec.
method	Method used to QT correction Permitted Values: "Bazett", "Fridericia", "Sagie"

Details

Depending on the chosen method one of the following formulae is used.

Bazett:

$$\frac{QT}{\sqrt{\frac{RR}{1000}}}$$

Fridericia:

$$\frac{QT}{\sqrt[3]{\frac{RR}{1000}}}$$

Sagie:

$$1000 \left(\frac{QT}{1000} + 0.154 \left(1 - \frac{RR}{1000} \right) \right)$$

Usually this computation function can not be used with %>%.

Value

QT interval in msec

See Also

BDS-Findings Functions that returns a vector: [compute_bmi\(\)](#), [compute_bsa\(\)](#), [compute_framingham\(\)](#), [compute_map\(\)](#), [compute_qual_imputation_dec\(\)](#), [compute_qual_imputation\(\)](#), [compute_rr\(\)](#), [compute_scale\(\)](#)

Examples

```
compute_qtc(qt = 350, rr = 56.54, method = "Bazett")
```

```
compute_qtc(qt = 350, rr = 56.54, method = "Fridericia")
```

```
compute_qtc(qt = 350, rr = 56.54, method = "Sagie")
```

```
compute_qual_imputation
```

Function to Impute Values When Qualifier Exists in Character Result

Description

Derive an imputed value

Usage

```
compute_qual_imputation(character_value, imputation_type = 1, factor = 0)
```

Arguments

character_value	Character version of value to be imputed
imputation_type	(default value=1) Valid Values: 1: Strip <, >, = and convert to numeric. 2: imputation_type=1 and if the character value contains a < or >, the number of of decimals associated with the character value is found and then a factor of $1/10^{(\text{number of decimals} + 1)}$ will be added/subtracted from the numeric value. If no decimals exists, a factor of $1/10$ will be added/subtracted from the value.
factor	Numeric value (default=0), when using imputation_type = 1, this value can be added or subtracted when the qualifier is removed.

Value

The imputed value

See Also

BDS-Findings Functions that returns a vector: [compute_bmi\(\)](#), [compute_bsa\(\)](#), [compute_framingham\(\)](#), [compute_map\(\)](#), [compute_qtc\(\)](#), [compute_qual_imputation_dec\(\)](#), [compute_rr\(\)](#), [compute_scale\(\)](#)

Examples

```
compute_qual_imputation("<40")
```

```
compute_qual_imputation_dec
```

Compute Factor for Value Imputations When Character Value Contains < or >

Description

Function to compute factor for value imputation when character value contains < or >. The factor is calculated using the number of decimals. If there are no decimals, the factor is 1, otherwise the factor = $1/10^{\text{decimal place}}$. For example, the factor for 100 = 1, the factor for 5.4 = $1/10^1$, the factor for 5.44 = $1/10^2$. This results in no additional false precision added to the value. This is an intermediate function.

Usage

```
compute_qual_imputation_dec(character_value_decimal)
```

Arguments

character_value_decimal	Character value to determine decimal precision
-------------------------	--

Details

Derive an imputed value

Value

Decimal precision value to add or subtract

See Also

BDS-Findings Functions that returns a vector: [compute_bmi\(\)](#), [compute_bsa\(\)](#), [compute_framingham\(\)](#), [compute_map\(\)](#), [compute_qtc\(\)](#), [compute_qual_imputation\(\)](#), [compute_rr\(\)](#), [compute_scale\(\)](#)

Examples

```
compute_qual_imputation_dec("<40.1")
```

compute_rr

Compute RR Interval From Heart Rate

Description

Computes RR interval from heart rate.

Usage

```
compute_rr(hr)
```

Arguments

hr	Heart rate
	A numeric vector is expected. It is expected that heart rate is measured in beats/min.

Details

Usually this computation function can not be used with %>%.

Value

RR interval in msec:

$$\frac{60000}{HR}$$

See Also

BDS-Findings Functions that returns a vector: [compute_bmi\(\)](#), [compute_bsa\(\)](#), [compute_framingham\(\)](#), [compute_map\(\)](#), [compute_qtc\(\)](#), [compute_qual_imputation_dec\(\)](#), [compute_qual_imputation\(\)](#), [compute_scale\(\)](#)

Examples

```
compute_rr(hr = 70.14)
```

compute_scale	<i>Compute Scale Parameters</i>
---------------	---------------------------------

Description

Computes the average of a set of source values and transforms the result from the source range to the target range. For example, for calculating the average of a set of questionnaire response scores and re-coding the average response to obtain a subscale score.

Usage

```
compute_scale(
  source,
  source_range = NULL,
  target_range = NULL,
  flip_direction = FALSE,
  min_n = 1
)
```

Arguments

source	A vector of values to be scaled A numeric vector is expected.
source_range	The permitted source range A numeric vector containing two elements is expected, representing the lower and upper bounds of the permitted source range. Alternatively, if no argument is specified for source_range and target_range, no transformation will be performed.
target_range	The target range A numeric vector containing two elements is expected, representing the lower and upper bounds of the target range. Alternatively, if no argument is specified for source_range and target_range, no transformation will be performed.
flip_direction	Flip direction of the scale? The transformed values will be reversed within the target range, e.g. within the range 0 to 100, 25 would be reversed to 75. This argument will be ignored if source_range and target_range aren't specified. Default: FALSE Permitted Values: TRUE, FALSE

min_n Minimum number of values for computation
 The minimum number of non-missing values in source for the computation to be carried out. If the number of non-missing values is below min_n, the result will be set to missing, i.e. NA.
 A positive integer is expected.
 Default: 1

Details

Returns a numeric value. If source contains less than min_n values, the result is set to NA. If source_range and target_range aren't specified, the mean will be computed without any transformation being performed.

Value

The average of source transformed to the target range or NA if source doesn't contain min_n values.

See Also

BDS-Findings Functions that returns a vector: [compute_bmi\(\)](#), [compute_bsa\(\)](#), [compute_framingham\(\)](#), [compute_map\(\)](#), [compute_qtc\(\)](#), [compute_qual_imputation_dec\(\)](#), [compute_qual_imputation\(\)](#), [compute_rr\(\)](#)

Examples

```
compute_scale(
  source = c(1, 4, 3, 5),
  source_range = c(1, 5),
  target_range = c(0, 100),
  flip_direction = TRUE,
  min_n = 3
)
```

compute_tmf

Derive the Time Imputation Flag

Description

Derive the time imputation flag ('--TMF') comparing a date character vector ('--DTC') with a Datetime vector ('--DTM').

Usage

```
compute_tmf(dtc, dtm, ignore_seconds_flag = FALSE)
```

Arguments

<code>dtc</code>	The date character vector ('--DTC'). A character date is expected in a format like yyyy-mm-ddThh:mm:ss (partial or complete).
<code>dtm</code>	The Date vector to compare ('--DTM'). A datetime object is expected.
<code>ignore_seconds_flag</code>	ADaM IG states that given SDTM ('--DTC') variable, if only hours and minutes are ever collected, and seconds are imputed in ('--DTM') as 00, then it is not necessary to set ('--TMF') to 'S'. A user can set this to TRUE so the 'S' Flag is dropped from ('--TMF'). A logical value Default: FALSE

Details

Usually this computation function can not be used with %>%.

Value

The time imputation flag ('--TMF') (character value of 'H', 'M', 'S' or NA)

See Also

Date/Time Computation Functions that returns a vector: [compute_dtf\(\)](#), [compute_duration\(\)](#), [convert_date_to_dtm\(\)](#), [convert_dtc_to_dtm\(\)](#), [convert_dtc_to_dt\(\)](#), [impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Examples

```
compute_tmf(dtc = "2019-07-18T15:25", dtm = as.POSIXct("2019-07-18T15:25:00"))
compute_tmf(dtc = "2019-07-18T15", dtm = as.POSIXct("2019-07-18T15:25:00"))
compute_tmf(dtc = "2019-07-18", dtm = as.POSIXct("2019-07-18"))
```

consolidate_metadata *Consolidate Multiple Meta Datasets Into a Single One*

Description

The purpose of the function is to consolidate multiple meta datasets into a single one. For example, from global and project specific parameter mappings a single lookup table can be created.

Usage

```
consolidate_metadata(
  datasets,
  key_vars,
  source_var = SOURCE,
  check_vars = "warning",
  check_keys = "error"
)
```

Arguments

datasets	List of datasets to consolidate <i>Permitted Values:</i> A named list of datasets
key_vars	Key variables The specified variables must be a unique of all input datasets. <i>Permitted Values:</i> A list of variables created by <code>exprs()</code>
source_var	Source variable The specified variable is added to the output dataset. It is set the name of the dataset the observation is originating from. <i>Permitted Values:</i> A symbol
check_vars	Check variables? If "message", "warning", or "error" is specified, a message is issued if the variable names differ across the input datasets (datasets). <i>Permitted Values:</i> "none", "message", "warning", "error"
check_keys	Check keys? If "warning" or "error" is specified, a message is issued if the key variables (key_vars) are not a unique key in all of the input datasets (datasets). <i>Permitted Values:</i> "none", "warning", "error"

Details

All observations of the input datasets are put together into a single dataset. If a by group (defined by `key_vars`) exists in more than one of the input datasets, the observation from the last dataset is selected.

Value

A dataset which contains one row for each by group occurring in any of the input datasets.

See Also

Creating auxiliary datasets: [create_period_dataset\(\)](#), [create_query_data\(\)](#), [create_single_dose_dataset\(\)](#)

Examples

```
library(tibble)
glob_ranges <- tribble(
  ~PARAMCD, ~ANRLO, ~ANRHI,
  "PULSE",   60,   100,
  "SYSBP",   90,   130,
  "DIABP",   60,   80
)
proj_ranges <- tribble(
  ~PARAMCD, ~ANRLO, ~ANRHI,
  "SYSBP",   100,   140,
  "DIABP",   70,   90
)
stud_ranges <- tribble(
  ~PARAMCD, ~ANRLO, ~ANRHI,
  "BMI",     18,   25
)

consolidate_metadata(
  datasets = list(
    global = glob_ranges,
    project = proj_ranges,
    study = stud_ranges
  ),
  key_vars = exprs(PARAMCD)
)
```

convert_blanks_to_na *Convert Blank Strings Into NAs*

Description

Turn SAS blank strings into proper R NAs.

Usage

```
convert_blanks_to_na(x)

## Default S3 method:
convert_blanks_to_na(x)

## S3 method for class 'character'
convert_blanks_to_na(x)

## S3 method for class 'list'
convert_blanks_to_na(x)

## S3 method for class 'data.frame'
convert_blanks_to_na(x)
```

Arguments

x Any R object

Details

The default methods simply returns its input unchanged. The character method turns every instance of "" into NA_character_ while preserving *all* attributes. When given a data frame as input the function keeps all non-character columns as is and applies the just described logic to character columns. Once again all attributes such as labels are preserved.

Value

An object of the same class as the input

See Also

Utilities for Formatting Observations: [convert_na_to_blanks\(\)](#), [yn_to_numeric\(\)](#)

Examples

```
library(tibble)

convert_blanks_to_na(c("a", "b", "", "d", ""))

df <- tibble(
  a = structure(c("a", "b", "", "c"), label = "A"),
  b = structure(c(1, NA, 21, 9), label = "B"),
  c = structure(c(TRUE, FALSE, TRUE, TRUE), label = "C"),
  d = structure(c("", "", "s", "q"), label = "D")
)
print(df)
convert_blanks_to_na(df)
```

convert_date_to_dtm *Convert a Date into a Datetime Object*

Description

Convert a date (datetime, date, or date character) into a Date vector (usually '--DTM').

Usage

```
convert_date_to_dtm(
  dt,
  highest_imputation = "h",
  date_imputation = "first",
  time_imputation = "first",
  min_dates = NULL,
```

```

max_dates = NULL,
preserve = FALSE
)

```

Arguments

- dt** The date to convert.
A date or character date is expected in a format like yyyy-mm-ddThh:mm:ss.
- highest_imputation**
Highest imputation level
The `highest_imputation` argument controls which components of the DTC value are imputed if they are missing. All components up to the specified level are imputed.
If a component at a higher level than the highest imputation level is missing, `NA_character_` is returned. For example, for `highest_imputation = "D"` `"2020"` results in `NA_character_` because the month is missing.
If `"n"` is specified, no imputation is performed, i.e., if any component is missing, `NA_character_` is returned.
If `"Y"` is specified, `date_imputation` should be `"first"` or `"last"` and `min_dates` or `max_dates` should be specified respectively. Otherwise, `NA_character_` is returned if the year component is missing.
Default: `"h"`
Permitted Values: `"Y"` (year, highest level), `"M"` (month), `"D"` (day), `"h"` (hour), `"m"` (minute), `"s"` (second), `"n"` (none, lowest level)
- date_imputation**
The value to impute the day/month when a datepart is missing.
A character value is expected, either as a
- format with month and day specified as `"mm-dd"`: e.g. `"06-15"` for the 15th of June (The year can not be specified; for imputing the year `"first"` or `"last"` together with `min_dates` or `max_dates` argument can be used (see examples).),
 - or as a keyword: `"first"`, `"mid"`, `"last"` to impute to the first/mid/last day/month.
- The argument is ignored if `highest_imputation` is less than `"D"`.
Default: `"first"`.
- time_imputation**
The value to impute the time when a timepart is missing.
A character value is expected, either as a
- format with hour, min and sec specified as `"hh:mm:ss"`: e.g. `"00:00:00"` for the start of the day,
 - or as a keyword: `"first"`, `"last"` to impute to the start/end of a day.
- The argument is ignored if `highest_imputation = "n"`.
Default: `"first"`.

min_dates Minimum dates

A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example

```
impute_dtc_dtm(
  "2020-11",
  min_dates = list(
    ymd_hms("2020-12-06T12:12:12"),
    ymd_hms("2020-11-11T11:11:11")
  ),
  highest_imputation = "M"
)
```

returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).

For date variables (not datetime) in the list the time is imputed to "00:00:00". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.

max_dates Maximum dates

A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.

For date variables (not datetime) in the list the time is imputed to "23:59:59". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.

preserve Preserve day if month is missing and day is present

For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "mid").

Permitted Values: TRUE, FALSE

Default: FALSE

Details

Usually this computation function can not be used with %>%.

Value

A datetime object

See Also

Date/Time Computation Functions that returns a vector: [compute_dtf\(\)](#), [compute_duration\(\)](#), [compute_tmf\(\)](#), [convert_dtc_to_dtm\(\)](#), [convert_dtc_to_dt\(\)](#), [impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Examples

```
convert_date_to_dtm("2019-07-18T15:25:00")
convert_date_to_dtm(Sys.time())
convert_date_to_dtm(as.Date("2019-07-18"), time_imputation = "23:59:59")
convert_date_to_dtm("2019-07-18", time_imputation = "23:59:59")
convert_date_to_dtm("2019-07-18")
```

convert_dtc_to_dt	<i>Convert a Date Character Vector into a Date Object</i>
-------------------	---

Description

Convert a date character vector (usually '-DTC') into a Date vector (usually '-DT').

Usage

```
convert_dtc_to_dt(
  dtc,
  highest_imputation = "n",
  date_imputation = "first",
  min_dates = NULL,
  max_dates = NULL,
  preserve = FALSE
)
```

Arguments

dtc	The -DTC date to convert.
highest_imputation	<p>Highest imputation level</p> <p>The highest_imputation argument controls which components of the DTC value are imputed if they are missing. All components up to the specified level are imputed.</p> <p>If a component at a higher level than the highest imputation level is missing, NA_character_ is returned. For example, for highest_imputation = "D" "2020" results in NA_character_ because the month is missing.</p> <p>If "n" is specified no imputation is performed, i.e., if any component is missing, NA_character_ is returned.</p> <p>If "Y" is specified, date_imputation should be "first" or "last" and min_dates or max_dates should be specified respectively. Otherwise, NA_character_ is returned if the year component is missing.</p>

Default: "n"

Permitted Values: "Y" (year, highest level), "M" (month), "D" (day), "n" (none, lowest level)

date_imputation

The value to impute the day/month when a datepart is missing.

A character value is expected, either as a

- format with month and day specified as "mm-dd": e.g. "06-15" for the 15th of June (The year can not be specified; for imputing the year "first" or "last" together with min_dates or max_dates argument can be used (see examples).),
- or as a keyword: "first", "mid", "last" to impute to the first/mid/last day/month.

The argument is ignored if highest_imputation is less than "D".

Default: "first"

min_dates

Minimum dates

A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example

```
impute_dtc_dtm(
  "2020-11",
  min_dates = list(
    ymd_hms("2020-12-06T12:12:12"),
    ymd_hms("2020-11-11T11:11:11")
  ),
  highest_imputation = "M"
)
```

returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).

max_dates

Maximum dates

A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.

preserve

Preserve day if month is missing and day is present

For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "MID").

Permitted Values: TRUE, FALSE

Default: FALSE

Details

Usually this computation function can not be used with %>%.

Value

a date object

See Also

Date/Time Computation Functions that returns a vector: [compute_dtf\(\)](#), [compute_duration\(\)](#), [compute_tmf\(\)](#), [convert_date_to_dtm\(\)](#), [convert_dtc_to_dtm\(\)](#), [impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Examples

```
convert_dtc_to_dt("2019-07-18")
convert_dtc_to_dt("2019-07")
```

convert_dtc_to_dtm	<i>Convert a Date Character Vector into a Datetime Object</i>
--------------------	---

Description

Convert a date character vector (usually '--DTC') into a Date vector (usually '--DTM').

Usage

```
convert_dtc_to_dtm(
  dtc,
  highest_imputation = "h",
  date_imputation = "first",
  time_imputation = "first",
  min_dates = NULL,
  max_dates = NULL,
  preserve = FALSE
)
```

Arguments

dtc	The '--DTC' date to convert.
highest_imputation	<p>Highest imputation level</p> <p>The highest_imputation argument controls which components of the DTC value are imputed if they are missing. All components up to the specified level are imputed.</p> <p>If a component at a higher level than the highest imputation level is missing, NA_character_ is returned. For example, for highest_imputation = "D" "2020" results in NA_character_ because the month is missing.</p>

If "n" is specified, no imputation is performed, i.e., if any component is missing, NA_character_ is returned.

If "Y" is specified, date_imputation should be "first" or "last" and min_dates or max_dates should be specified respectively. Otherwise, NA_character_ is returned if the year component is missing.

Default: "h"

Permitted Values: "Y" (year, highest level), "M" (month), "D" (day), "h" (hour), "m" (minute), "s" (second), "n" (none, lowest level)

date_imputation

The value to impute the day/month when a datepart is missing.

A character value is expected, either as a

- format with month and day specified as "mm-dd": e.g. "06-15" for the 15th of June (The year can not be specified; for imputing the year "first" or "last" together with min_dates or max_dates argument can be used (see examples).),
- or as a keyword: "first", "mid", "last" to impute to the first/mid/last day/month.

The argument is ignored if highest_imputation is less than "D".

Default: "first".

time_imputation

The value to impute the time when a timepart is missing.

A character value is expected, either as a

- format with hour, min and sec specified as "hh:mm:ss": e.g. "00:00:00" for the start of the day,
- or as a keyword: "first", "last" to impute to the start/end of a day.

The argument is ignored if highest_imputation = "n".

Default: "first".

min_dates

Minimum dates

A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example

```
impute_dtc_dtm(
  "2020-11",
  min_dates = list(
    ymd_hms("2020-12-06T12:12:12"),
    ymd_hms("2020-11-11T11:11:11")
  ),
  highest_imputation = "M"
)
```

returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12"

is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).

For date variables (not datetime) in the list the time is imputed to "00:00:00". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.

max_dates	<p>Maximum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.</p> <p>For date variables (not datetime) in the list the time is imputed to "23:59:59". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.</p>
preserve	<p>Preserve day if month is missing and day is present</p> <p>For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "mid").</p> <p>Permitted Values: TRUE, FALSE</p> <p><i>Default:</i> FALSE</p>

Details

Usually this computation function can not be used with %>%.

Value

A datetime object

See Also

Date/Time Computation Functions that returns a vector: [compute_dtf\(\)](#), [compute_duration\(\)](#), [compute_tmf\(\)](#), [convert_date_to_dtm\(\)](#), [convert_dtc_to_dt\(\)](#), [impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Examples

```
convert_dtc_to_dtm("2019-07-18T15:25:00")
convert_dtc_to_dtm("2019-07-18T00:00:00") # note Time = 00:00:00 is not printed
convert_dtc_to_dtm("2019-07-18")
```

convert_na_to_blanks *Convert NAs Into Blank Strings*

Description

Turn NAs to blank strings .

Usage

```
convert_na_to_blanks(x)

## Default S3 method:
convert_na_to_blanks(x)

## S3 method for class 'character'
convert_na_to_blanks(x)

## S3 method for class 'list'
convert_na_to_blanks(x)

## S3 method for class 'data.frame'
convert_na_to_blanks(x)
```

Arguments

x Any R object

Details

The default methods simply returns its input unchanged. The character method turns every instance of `NA_character_` or `NA` into `""` while preserving *all* attributes. When given a data frame as input the function keeps all non-character columns as is and applies the just described logic to character all attributes such as labels are preserved.

Value

An object of the same class as the input

See Also

Utilities for Formatting Observations: [convert_blanks_to_na\(\)](#), [yn_to_numeric\(\)](#)

Examples

```
library(tibble)

convert_na_to_blanks(c("a", "b", NA, "d", NA))

df <- tibble(
  a = structure(c("a", "b", NA, "c"), label = "A"),
  b = structure(c(1, NA, 21, 9), label = "B"),
  c = structure(c(TRUE, FALSE, TRUE, TRUE), label = "C"),
  d = structure(c(NA, NA, "s", "q"), label = "D")
)
print(df)
convert_na_to_blanks(df)
```

count_vals

*Count Number of Observations Where a Variable Equals a Value***Description**

Count number of observations where a variable equals a value.

Usage

```
count_vals(var, val)
```

Arguments

var	A vector
val	A value

See Also

Utilities for Filtering Observations: [filter_extreme\(\)](#), [filter_joined\(\)](#), [filter_relative\(\)](#), [max_cond\(\)](#), [min_cond\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(admiral)
data <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      1,      "PR",
  "1",      2,      "CR",
  "1",      3,      "NE",
  "1",      4,      "CR",
  "1",      5,      "NE",
  "2",      1,      "CR",
  "2",      2,      "PR",
  "2",      3,      "CR",
  "3",      1,      "CR",
  "4",      1,      "CR",
  "4",      2,      "NE",
  "4",      3,      "NE",
  "4",      4,      "CR",
  "4",      5,      "PR"
)

# add variable providing the number of NEs for each subject
group_by(data, USUBJID) %>%
  mutate(nr_nes = count_vals(var = AVALC, val = "NE"))
```

create_period_dataset *Create a Reference Dataset for Subperiods, Periods, or Phases*

Description

The function creates a reference dataset for subperiods, periods, or phases from the ADSL dataset. The reference dataset can be used to derive subperiod, period, or phase variables like ASPER, ASPRSDT, ASPREDT, APERIOD, APERSDT, APEREDT, TRTA, APHASEN, PHSDTM, PHEDTM, ... in OCCDS and BDS datasets.

Usage

```
create_period_dataset(
  dataset,
  new_vars,
  subject_keys = get_admiral_option("subject_keys")
)
```

Arguments

dataset	ADSL dataset The variables specified by new_vars and subject_keys are expected. For each element of new_vars at least one variable of the form of the right hand side value must be available in the dataset.
new_vars	New variables A named list of variables like exprs(PHSDT = PHwSDT, PHEDT = PHwEDT, APHASE = APHASEw) is expected. The left hand side of the elements defines a variable of the output dataset, the right hand side defines the source variables from the ADSL dataset in CDISC notation. If the lower case letter "w" is used it refers to a phase variable, if the lower case letters "xx" are used it refers to a period variable, and if both "xx" and "w" are used it refers to a subperiod variable. Only one type must be used, e.g., all right hand side values must refer to period variables. It is not allowed to mix for example period and subperiod variables. If period <i>and</i> subperiod variables are required, separate reference datasets must be created.
subject_keys	Variables to uniquely identify a subject A list of expressions where the expressions are symbols as returned by exprs() is expected.

Details

For each subject and each subperiod/period/phase where at least one of the source variable is not NA an observation is added to the output dataset.

Depending on the type of the source variable (subperiod, period, or phase) the variable ASPER, APERIOD, or APHASEN is added and set to the number of the subperiod, period, or phase.

The variables specified for `new_vars` (left hand side) are added to the output dataset and set to the value of the source variable (right hand side).

Value

A period reference dataset (see "Details" section)

See Also

[derive_vars_period\(\)](#)

Creating auxiliary datasets: [consolidate_metadata\(\)](#), [create_query_data\(\)](#), [create_single_dose_dataset\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

# Create reference dataset for periods
adsl <- tribble(
  ~USUBJID, ~AP01SDT, ~AP01EDT, ~AP02SDT, ~AP02EDT, ~TRT01A, ~TRT02A,
  "1", "2021-01-04", "2021-02-06", "2021-02-07", "2021-03-07", "A", "B",
  "2", "2021-02-02", "2021-03-02", "2021-03-03", "2021-04-01", "B", "A",
) %>%
  mutate(
    across(matches("AP\\d\\d[ES]DT"), ymd)
  ) %>%
  mutate(
    STUDYID = "xyz"
  )

create_period_dataset(
  adsl,
  new_vars = exprs(APERSDT = APxxSDT, APEREDT = APxxEDT, TRTA = TRTxxA)
)

# Create reference dataset for phases
adsl <- tribble(
  ~USUBJID, ~PH1SDT, ~PH1EDT, ~PH2SDT, ~PH2EDT, ~APHASE1, ~APHASE2,
  "1", "2021-01-04", "2021-02-06", "2021-02-07", "2021-03-07", "TREATMENT", "FUP",
  "2", "2021-02-02", "2021-03-02", NA, NA, "TREATMENT", NA
) %>%
  mutate(
    across(matches("PH\\d[ES]DT"), ymd)
  ) %>%
  mutate(
    STUDYID = "xyz"
  )

create_period_dataset(
  adsl,
  new_vars = exprs(PHSDT = PHwSDT, PHEDT = PHwEDT, APHASE = APHASEw)
```

```

)

# Create reference datasets for subperiods
adsl <- tribble(
  ~USUBJID, ~P01S1SDT, ~P01S1EDT, ~P01S2SDT, ~P01S2EDT, ~P02S1SDT, ~P02S1EDT,
  "1", "2021-01-04", "2021-01-19", "2021-01-20", "2021-02-06", "2021-02-07", "2021-03-07",
  "2", "2021-02-02", "2021-03-02", NA, NA, "2021-03-03", "2021-04-01"
) %>%
  mutate(
    across(matches("P\\d\\dS\\d[ES]DT"), ymd)
  ) %>%
  mutate(
    STUDYID = "xyz"
  )

create_period_dataset(
  adsl,
  new_vars = exprs(ASPRSDT = PxxSwSDT, ASPREDT = PxxSwEDT)
)

```

create_query_data	<i>Creates a queries dataset as input dataset to the dataset_queries argument in derive_vars_query()</i>
-------------------	--

Description

Creates a queries dataset as input dataset to the dataset_queries argument in the derive_vars_query() function as defined in the [Queries Dataset Documentation](#).

Usage

```
create_query_data(queries, version = NULL, get_terms_fun = NULL)
```

Arguments

queries	List of queries A list of query() objects is expected.
version	Dictionary version The dictionary version used for coding the terms should be specified. If any of the queries is a basket (SMQ, SDG,) or a customized query including a basket, the parameter needs to be specified. <i>Permitted Values:</i> A character string (the expected format is company-specific)
get_terms_fun	Function which returns the terms For each query specified for the queries parameter referring to a basket (i.e., those where the definition field is set to a basket_select() object or a list which contains at least one basket_select() object) the specified function is called to retrieve the terms defining the query. This function is not provided by admiral as it is company specific, i.e., it has to be implemented at company level.

The function must return a dataset with all the terms defining the basket. The output dataset must contain the following variables.

- **TERM_LEVEL**: the variable to be used for defining a term of the basket, e.g., AEDECOD
- **TERM_NAME**: the name of the term if the variable **TERM_LEVEL** is referring to is character
- **TERM_ID**: the numeric id of the term if the variable **TERM_LEVEL** is referring to is numeric
- **QUERY_NAME**: the name of the basket. The values must be the same for all observations.

The function must provide the following parameters

- **basket_select**: A `basket_select()` object.
- **version**: The dictionary version. The value specified for the `version` in the `create_query_data()` call is passed to this parameter.
- **keep_id**: If set to `TRUE`, the output dataset must contain the `QUERY_ID` variable. The variable must be set to the numeric id of the basket.
- **temp_env**: A temporary environment is passed to this parameter. It can be used to store data which is used for all baskets in the `create_query_data()` call. For example if SMQs need to be read from a database all SMQs can be read and stored in the environment when the first SMQ is handled. For the other SMQs the terms can be retrieved from the environment instead of accessing the database again.

Details

For each `query()` object listed in the `queries` argument, the terms belonging to the query (`TERM_LEVEL`, `TERM_NAME`, `TERM_ID`) are determined with respect to the `definition` field of the query: if the `definition` field of the `query()` object is

- a `basket_select()` object, the terms are read from the basket database by calling the function specified for the `get_terms_fun` parameter.
- a data frame, the terms stored in the data frame are used.
- a list of data frames and `basket_select()` objects, all terms from the data frames and all terms read from the basket database referenced by the `basket_select()` objects are collated.

The following variables (as described in [Queries Dataset Documentation](#)) are created:

- **VAR_PREFIX**: Prefix of the variables to be created by `derive_vars_query()` as specified by the `prefix` element.
- **QUERY_NAME**: Name of the query as specified by the `name` element.
- **QUERY_ID**: Id of the query as specified by the `id` element. If the `id` element is not specified for a query, the variable is set to `NA`. If the `id` element is not specified for any query, the variable is not created.
- **QUERY_SCOPE**: scope of the query as specified by the `scope` element of the `basket_select()` object. For queries not defined by a `basket_select()` object, the variable is set to `NA`. If none of the queries is defined by a `basket_select()` object, the variable is not created.

- QUERY_SCOPE_NUM: numeric scope of the query. It is set to 1 if the scope is broad. Otherwise it is set to 2. If the add_scope_num element equals FALSE, the variable is set to NA. If the add_scope_num element equals FALSE for all baskets or none of the queries is an basket , the variable is not created.
- TERM_LEVEL: Name of the variable used to identify the terms.
- TERM_NAME: Value of the term variable if it is a character variable.
- TERM_ID: Value of the term variable if it is a numeric variable.
- VERSION: Set to the value of the version argument. If it is not specified, the variable is not created.

Value

A dataset to be used as input dataset to the dataset_queries argument in derive_vars_query()

See Also

[derive_vars_query\(\)](#), [query\(\)](#), [basket_select\(\)](#), [Queries Dataset Documentation](#)

Creating auxiliary datasets: [consolidate_metadata\(\)](#), [create_period_dataset\(\)](#), [create_single_dose_dataset\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
library(admiral)

# creating a query dataset for a customized query
cqterms <- tribble(
  ~TERM_NAME, ~TERM_ID,
  "APPLICATION SITE ERYTHEMA", 10003041L,
  "APPLICATION SITE PRURITUS", 10003053L
) %>%
  mutate(TERM_LEVEL = "AEDECOD")

cq <- query(
  prefix = "CQ01",
  name = "Application Site Issues",
  definition = cqterms
)

create_query_data(queries = list(cq))

# create a query dataset for SMQs
pregsmq <- query(
  prefix = "SMQ02",
  id = auto,
  definition = basket_select(
    name = "Pregnancy and neonatal topics (SMQ)",
    scope = "NARROW",
    type = "smq"
  )
)
```

```

)
)

bilismq <- query(
  prefix = "SMQ04",
  definition = basket_select(
    id = 20000121L,
    scope = "BROAD",
    type = "smq"
  )
)

# The get_terms function from admiral.test is used for this example.
# In a real application a company-specific function must be used.
create_query_data(
  queries = list(pregsmq, bilismq),
  get_terms_fun = admiral.test::get_terms,
  version = "20.1"
)

# create a query dataset for SDGs
sdg <- query(
  prefix = "SDG01",
  id = auto,
  definition = basket_select(
    name = "5-aminosalicylates for ulcerative colitis",
    scope = NA_character_,
    type = "sdg"
  )
)

# The get_terms function from admiral.test is used for this example.
# In a real application a company-specific function must be used.
create_query_data(
  queries = list(sdg),
  get_terms_fun = admiral.test::get_terms,
  version = "2019-09"
)

# creating a query dataset for a customized query including SMQs
# The get_terms function from admiral.test is used for this example.
# In a real application a company-specific function must be used.
create_query_data(
  queries = list(
    query(
      prefix = "CQ03",
      name = "Special issues of interest",
      definition = list(
        basket_select(
          name = "Pregnancy and neonatal topics (SMQ)",
          scope = "NARROW",
          type = "smq"
        )
      ),
    )
  )
)

```

```

        cqterms
      )
    )
  ),
  get_terms_fun = admiral.test::get_terms,
  version = "20.1"
)

```

```
create_single_dose_dataset
```

Create dataset of single doses

Description

Derives dataset of single dose from aggregate dose information. This may be necessary when e.g. calculating last dose before an adverse event in ADAE or deriving a total dose parameter in ADEX when EXDOSFRQ != ONCE.

Usage

```

create_single_dose_dataset(
  dataset,
  dose_freq = EXDOSFRQ,
  start_date = ASTDT,
  start_datetime = NULL,
  end_date = AENDT,
  end_datetime = NULL,
  lookup_table = dose_freq_lookup,
  lookup_column = CDISC_VALUE,
  nominal_time = NULL,
  keep_source_vars = expr_c(exprs(USUBJID), dose_freq, start_date, start_datetime,
    end_date, end_datetime)
)

```

Arguments

dataset	Input dataset The columns specified by dose_freq, start_date and the end_date parameters are expected.
dose_freq	The dose frequency The aggregate dosing frequency used for multiple doses in a row. Permitted Values: defined by lookup table.
start_date	The start date A date object is expected. This object cannot contain NA values. Refer to derive_vars_dt() to impute and derive a date from a date character vector to a date object.

start_datetime	<p>The start date-time</p> <p>A date-time object is expected. This object cannot contain NA values.</p> <p>Refer to <code>derive_vars_dtm()</code> to impute and derive a date-time from a date character vector to a date object.</p> <p>If the input dataset contains frequencies which refer to DOSE_WINDOW equals "HOUR" or "MINUTE", the parameter must be specified.</p>
end_date	<p>The end date</p> <p>A date or date-time object is expected. This object cannot contain NA values.</p> <p>Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object.</p>
end_datetime	<p>The end date-time</p> <p>A date-time object is expected. This object cannot contain NA values.</p> <p>Refer to <code>derive_vars_dtm()</code> to impute and derive a date-time from a date character vector to a date object.</p> <p>If the input dataset contains frequencies which refer to DOSE_WINDOW equals "HOUR" or "MINUTE", the parameter must be specified.</p>
lookup_table	<p>The dose frequency value lookup table</p> <p>The table used to look up dose_freq values and determine the appropriate multiplier to be used for row generation. If a lookup table other than the default is used, it must have columns DOSE_WINDOW, DOSE_COUNT, and CONVERSION_FACTOR. The default table dose_freq_lookup is described in detail here.</p> <p>Permitted Values for DOSE_WINDOW: "MINUTE", "HOUR", "DAY", "WEEK", "MONTH", "YEAR"</p>
lookup_column	<p>The dose frequency value column in the lookup table</p> <p>The column of lookup_table.</p>
nominal_time	<p>The nominal relative time from first dose (NFRLT)</p> <p>Used for PK analysis, this will be in hours and should be 0 for the first dose. It can be derived as $(VISITDY - 1) * 24$ for example. This will be expanded as the single dose dataset is created. For example an EXDOFRQ of "QD" will result in the nominal_time being incremented by 24 hours for each expanded record.</p> <p>The value can be NULL if not needed.</p>
keep_source_vars	<p>List of variables to be retained from source dataset</p> <p>This parameter can be specified if additional information is required in the output dataset. For example EXTRT for studies with more than one drug.</p>

Details

Each aggregate dose row is split into multiple rows which each represent a single dose. The number of completed dose periods between start_date or start_datetime and end_date or end_datetime is calculated with `compute_duration` and multiplied by DOSE_COUNT. For DOSE_WINDOW values of "WEEK", "MONTH", and "YEAR", CONVERSION_FACTOR is used to convert into days the time object to be added to start_date.

Observations with dose frequency "ONCE" are copied to the output dataset unchanged.

Value

The input dataset with a single dose per row.

See Also

Creating auxiliary datasets: [consolidate_metadata\(\)](#), [create_period_dataset\(\)](#), [create_query_data\(\)](#)

Examples

```
# Example with default lookup

library(lubridate)
library(stringr)
library(tibble)
library(dplyr)

data <- tribble(
  ~USUBJID, ~EXDOSFRQ, ~ASTDT, ~ASTDTM, ~AENDT, ~AENDTM,
  "P01", "Q2D", ymd("2021-01-01"), ymd_hms("2021-01-01 10:30:00"),
  ymd("2021-01-07"), ymd_hms("2021-01-07 11:30:00"),
  "P01", "Q3D", ymd("2021-01-08"), ymd_hms("2021-01-08 12:00:00"),
  ymd("2021-01-14"), ymd_hms("2021-01-14 14:00:00"),
  "P01", "EVERY 2 WEEKS", ymd("2021-01-15"), ymd_hms("2021-01-15 09:57:00"),
  ymd("2021-01-29"), ymd_hms("2021-01-29 10:57:00")
)

create_single_dose_dataset(data)

# Example with custom lookup

custom_lookup <- tribble(
  ~Value, ~DOSE_COUNT, ~DOSE_WINDOW, ~CONVERSION_FACTOR,
  "Q30MIN", (1 / 30), "MINUTE", 1,
  "Q90MIN", (1 / 90), "MINUTE", 1
)

data <- tribble(
  ~USUBJID, ~EXDOSFRQ, ~ASTDT, ~ASTDTM, ~AENDT, ~AENDTM,
  "P01", "Q30MIN", ymd("2021-01-01"), ymd_hms("2021-01-01T06:00:00"),
  ymd("2021-01-01"), ymd_hms("2021-01-01T07:00:00"),
  "P02", "Q90MIN", ymd("2021-01-01"), ymd_hms("2021-01-01T06:00:00"),
  ymd("2021-01-01"), ymd_hms("2021-01-01T09:00:00")
)

create_single_dose_dataset(data,
  lookup_table = custom_lookup,
  lookup_column = Value,
  start_datetime = ASTDTM,
  end_datetime = AENDTM
)

# Example with nominal time
```

```

data <- tribble(
  ~USUBJID, ~EXDOSFRQ, ~NFRLT, ~ASTDT, ~ASTDTM, ~AENDT, ~AENDTM,
  "P01", "BID", 0, ymd("2021-01-01"), ymd_hms("2021-01-01 08:00:00"),
  ymd("2021-01-07"), ymd_hms("2021-01-07 20:00:00"),
  "P01", "BID", 168, ymd("2021-01-08"), ymd_hms("2021-01-08 08:00:00"),
  ymd("2021-01-14"), ymd_hms("2021-01-14 20:00:00"),
  "P01", "BID", 336, ymd("2021-01-15"), ymd_hms("2021-01-15 08:00:00"),
  ymd("2021-01-29"), ymd_hms("2021-01-29 20:00:00")
)

create_single_dose_dataset(data,
  dose_freq = EXDOSFRQ,
  start_date = ASTDT,
  start_datetime = ASTDTM,
  end_date = AENDT,
  end_datetime = AENDTM,
  lookup_table = dose_freq_lookup,
  lookup_column = CDISC_VALUE,
  nominal_time = NFRLT,
  keep_source_vars = exprs(
    USUBJID, EXDOSFRQ, ASTDT, ASTDTM, AENDT, AENDTM, NFRLT
  )
)

# Example - derive a single dose dataset with imputations

# For either single drug administration records, or multiple drug administration
# records covering a range of dates, fill-in of missing treatment end datetime
# `EXENDTC` by substitution with an acceptable alternate, for example date of
# death, date of datacut may be required. This example shows the
# maximum possible number of single dose records to be derived. The example
# requires the date of datacut `DCUTDT` to be specified correctly, or
# if not appropriate to use `DCUTDT` as missing treatment end data and missing
# treatment end datetime could set equal to treatment start date and treatment
# start datetime. ADSL variables `DTHDT` and `DCUTDT` are preferred for
# imputation use.
#
# All available trial treatments are included, allowing multiple different
# last dose variables to be created in for example `use_ad_template("ADAE")`
# if required.

adsl <- tribble(
  ~STUDYID, ~USUBJID, ~DTHDT,
  "01", "1211", ymd("2013-01-14"),
  "01", "1083", ymd("2013-08-02"),
  "01", "1445", ymd("2014-11-01"),
  "01", "1015", NA,
  "01", "1023", NA
)

ex <- tribble(
  ~STUDYID, ~USUBJID, ~EXSEQ, ~EXTRT, ~EXDOSE, ~EXDOSU, ~EXDOSFRQ, ~EXSTDTC, ~EXENDTC,
  "01", "1015", 1, "PLAC", 0, "mg", "QD", "2014-01-02", "2014-01-16",

```

```

"01", "1015", 2, "PLAC", 0, "mg", "QD", "2014-06-17", "2014-06-18",
"01", "1015", 3, "PLAC", 0, "mg", "QD", "2014-06-19", NA_character_,
"01", "1023", 1, "PLAC", 0, "mg", "QD", "2012-08-05", "2012-08-27",
"01", "1023", 2, "PLAC", 0, "mg", "QD", "2012-08-28", "2012-09-01",
"01", "1211", 1, "XANO", 54, "mg", "QD", "2012-11-15", "2012-11-28",
"01", "1211", 2, "XANO", 54, "mg", "QD", "2012-11-29", NA_character_,
"01", "1445", 1, "PLAC", 0, "mg", "QD", "2014-05-11", "2014-05-25",
"01", "1445", 2, "PLAC", 0, "mg", "QD", "2014-05-26", "2014-11-01",
"01", "1083", 1, "PLAC", 0, "mg", "QD", "2013-07-22", "2013-08-01"
)

adsl_death <- adsl %>%
  mutate(
    DTHDTM = convert_date_to_dtm(DTHDT),
    # Remove `DCUT` setup line below if ADSL `DCUTDT` is populated.
    DCUTDT = convert_dtc_to_dt("2015-03-06"), # Example only, enter date.
    DCUTDTM = convert_date_to_dtm(DCUTDT)
  )

# Select valid dose records, non-missing `EXSTDTC` and `EXDOSE`.
ex_mod <- ex %>%
  filter(!is.na(EXSTDTC) & !is.na(EXDOSE)) %>%
  derive_vars_merged(adsl_death, by_vars = exprs(STUDYID, USUBJID)) %>%
  # Example, set up missing `EXDOSFRQ` as QD daily dosing regime.
  # Replace with study dosing regime per trial treatment.
  mutate(EXDOSFRQ = if_else(is.na(EXDOSFRQ), "QD", EXDOSFRQ)) %>%
  # Create EXxxDTM variables and replace missing `EXENDTM`.
  derive_vars_dtm(
    dtc = EXSTDTC,
    new_vars_prefix = "EXST",
    date_imputation = "first",
    time_imputation = "first",
    flag_imputation = "none",
  ) %>%
  derive_vars_dtm_to_dt(exprs(EXSTDTC)) %>%
  derive_vars_dtm(
    dtc = EXENDTC,
    new_vars_prefix = "EXEN",
    # Maximum imputed treatment end date must not be not greater than
    # date of death or after the datacut date.
    max_dates = exprs(DTHDTM, DCUTDTM),
    date_imputation = "last",
    time_imputation = "last",
    flag_imputation = "none",
    highest_imputation = "Y",
  ) %>%
  derive_vars_dtm_to_dt(exprs(EXENDTM)) %>%
  # Select only unique values.
  # Removes duplicated records before final step.
  distinct(
    STUDYID, USUBJID, EXTRT, EXDOSE, EXDOSFRQ, DCUTDT, DTHDT, EXSTDTC,
    EXSTDTC, EXENDT, EXENDTM, EXSTDTC, EXENDTC
  )

```

```

create_single_dose_dataset(
  ex_mod,
  start_date = EXSTDT,
  start_datetime = EXSTD TM,
  end_date = EXENDT,
  end_datetime = EXEND TM,
  keep_source_vars = exprs(
    STUDYID, USUBJID, EXTRT, EXDOSE, EXDOSFRQ,
    DCUTDT, EXSTDT, EXSTD TM, EXENDT, EXEND TM, EXSTD TC, EXEND TC
  )
)

```

date_source

Create a date_source object

Description

Create a date_source object as input for derive_var_extreme_dt() and derive_var_extreme_dtm().

Usage

```
date_source(dataset_name, filter = NULL, date, traceability_vars = NULL)
```

Arguments

dataset_name	The name of the dataset, i.e. a string, used to search for the date.
filter	An unquoted condition for filtering dataset.
date	A variable providing a date. A date or a datetime can be specified. An unquoted symbol is expected.
traceability_vars	A named list returned by exprs() defining the traceability variables, e.g. exprs(LALVDOM = "AE", LALVSEQ = AESEQ, LALVVAR = "AESTDTC"). The values must be a symbol, a character string, a numeric, or NA.

Value

An object of class date_source.

See Also

[derive_var_extreme_dtm\(\)](#), [derive_var_extreme_dt\(\)](#)

Source Objects: [basket_select\(\)](#), [censor_source\(\)](#), [death_event](#), [dthcaus_source\(\)](#), [event_source\(\)](#), [event\(\)](#), [query\(\)](#), [tte_source\(\)](#)

Examples

```
# treatment end date from ADSL
trt_end_date <- date_source(
  dataset_name = "adsl",
  date = TRTEDT
)

# lab date from LB where assessment was taken, i.e. not "NOT DONE"
lb_date <- date_source(
  dataset_name = "lb",
  filter = LBSTAT != "NOT DONE" | is.na(LBSTAT),
  date = LBDT
)

# death date from ADSL including traceability variables
death_date <- date_source(
  dataset_name = "adsl",
  date = DTHDT,
  traceability_vars = exprs(
    LALVDM = "ADSL",
    LALVVR = "DTHDT"
  )
)
```

death_event

Pre-Defined Time-to-Event Source Objects

Description

These pre-defined tte_source objects can be used as input to [derive_param_tte\(\)](#).

Usage

death_event

lastalive_censor

ae_event

ae_ser_event

ae_gr1_event

ae_gr2_event

ae_gr3_event

```
ae_gr4_event
ae_gr5_event
ae_gr35_event
ae_sev_event
ae_wd_event
```

Details

To see the definition of the various objects simply print the object in the R console, e.g. `print(death_event)`. For details of how to use these objects please refer to `derive_param_tte()`.

See Also

`derive_param_tte()`, `tte_source()`, `event_source()`, `censor_source()`

Source Objects: `basket_select()`, `censor_source()`, `date_source()`, `dthcaus_source()`, `event_source()`, `event()`, `query()`, `tte_source()`

Examples

```
# This shows the definition of all pre-defined `tte_source` objects that ship
# with {admiral}
for (obj in list_tte_source_objects())$object) {
  cat(obj, "\n")
  print(get(obj))
  cat("\n")
}
```

default_qtc_paramcd *Get Default Parameter Code for Corrected QT*

Description

Get Default Parameter Code for Corrected QT

Usage

```
default_qtc_paramcd(method)
```

Arguments

method	Method used to QT correction Permitted Values: "Bazett", "Fridericia", "Sagie"
--------	---

Value

"QTCBR" if method is "Bazett", "QTCFR" if it's "Fridericia" or "QTLCR" if it's "Sagie". An error otherwise.

See Also

BDS-Findings Functions for adding Parameters/Records: [derive_expected_records\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_rr\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```
default_qtc_paramcd("Sagie")
```

derivation_slice	<i>Create a derivation_slice Object</i>
------------------	---

Description

Create a `derivation_slice` object as input for `slice_derivation()`.

Usage

```
derivation_slice(filter, args = NULL)
```

Arguments

<code>filter</code>	An unquoted condition for defining the observations of the slice
<code>args</code>	Arguments of the derivation to be used for the slice A <code>params()</code> object is expected.

Value

An object of class `derivation_slice`

See Also

[slice_derivation\(\)](#), [params\(\)](#)

Higher Order Functions: [call_derivation\(\)](#), [restrict_derivation\(\)](#), [slice_derivation\(\)](#)

derive_derived_param *Adds a Parameter Computed from the Analysis Value of Other Parameters*

Description

[Deprecated]

This function is deprecated. Please use `derive_param_computed()` instead.

Usage

```
derive_derived_param(
  dataset,
  by_vars,
  parameters,
  analysis_value,
  set_values_to,
  filter = NULL,
  constant_by_vars = NULL,
  constant_parameters = NULL
)
```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>parameters</code>.</p>
by_vars	<p>Grouping variables</p> <p>For each group defined by <code>by_vars</code> an observation is added to the output dataset. Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p><i>Permitted Values:</i> list of variables</p>
parameters	<p>Required parameter codes</p> <p>It is expected that all parameter codes (<code>PARAMCD</code>) which are required to derive the new parameter are specified for this parameter or the <code>constant_parameters</code> parameter.</p> <p><i>Permitted Values:</i> A character vector of <code>PARAMCD</code> values</p>
analysis_value	<p>Definition of the analysis value</p> <p>An expression defining the analysis value (<code>AVAL</code>) of the new parameter is expected. The analysis values of the parameters specified by <code>parameters</code> can be accessed using <code>AVAL.<parameter code></code>, e.g., <code>AVAL.SYSBP</code>.</p> <p><i>Permitted Values:</i> An unquoted expression</p>

set_values_to	<p>Variables to be set</p> <p>The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter.</p> <p><i>Permitted Values:</i> List of variable-value pairs</p>
filter	<p>Filter condition</p> <p>The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account.</p> <p><i>Permitted Values:</i> a condition</p>
constant_by_vars	<p>By variables for constant parameters</p> <p>The constant parameters (parameters that are measured only once) are merged to the other parameters using the specified variables. (Refer to Example 2)</p> <p><i>Permitted Values:</i> list of variables</p>
constant_parameters	<p>Required constant parameter codes</p> <p>It is expected that all the parameter codes (PARAMCD) which are required to derive the new parameter and are measured only once are specified here. For example if BMI should be derived and height is measured only once while weight is measured at each visit. Height could be specified in the <code>constant_parameters</code> parameter. (Refer to Example 2)</p> <p><i>Permitted Values:</i> A character vector of PARAMCD values</p>

Value

The input dataset with the new parameter added. Note, a variable will only be populated in the new parameter rows if it is specified in `by_vars`.

See Also

Other deprecated: `derive_param_first_event()`, `derive_var_agegr_fda()`, `derive_var_confirmation_flag()`, `derive_var_disposition_status()`, `derive_var_worst_flag()`, `derive_vars_disposition_reason()`, `derive_vars_merged_dtm()`, `derive_vars_merged_dt()`, `filter_confirmation()`, `format_eoxxstt_default()`, `format_reason_default()`, `vars()`

derive_expected_records

Derive Expected Records

Description

Add expected records as new observations for each 'by group' when the dataset contains missing observations.

Usage

```
derive_expected_records(
  dataset,
  dataset_expected_obs,
  by_vars = NULL,
  set_values_to = NULL
)
```

Arguments

dataset	Input dataset A data frame, the columns from dataset_expected_obs and specified by the by_vars parameter are expected.
dataset_expected_obs	Expected observations dataset Data frame with the expected observations, e.g., all the expected combinations of PARAMCD, PARAM, AVISIT, AVISITN, ...
by_vars	Grouping variables For each group defined by by_vars those observations from dataset_expected_obs are added to the output dataset which do not have a corresponding observation in the input dataset.
set_values_to	Variables to be set The specified variables are set to the specified values for the new observations. A list of variable name-value pairs is expected. <ul style="list-style-type: none"> • LHS refers to a variable. • RHS refers to the values to set to the variable. This can be a string, a symbol, a numeric value or NA, e.g., exprs(PARAMCD = "TDOSE", PARCAT1 = "OVERALL"). More general expression are not allowed.

Details

For each group (the variables specified in the by_vars parameter), those records from dataset_expected_obs that are missing in the input dataset are added to the output dataset.

Value

The input dataset with the missed expected observations added for each by_vars. Note, a variable will only be populated in the new parameter rows if it is specified in by_vars or set_values_to.

See Also

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_rr\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```

library(tibble)

adqs <- tribble(
  ~USUBJID, ~PARAMCD, ~AVISITN, ~AVISIT, ~AVAL,
  "1",      "a",      1, "WEEK 1",  10,
  "1",      "b",      1, "WEEK 1",  11,
  "2",      "a",      2, "WEEK 2",  12,
  "2",      "b",      2, "WEEK 2",  14
)

# Example 1. visit variables are parameter independent
parm_visit_ref <- tribble(
  ~AVISITN, ~AVISIT,
  1,        "WEEK 1",
  2,        "WEEK 2"
)

derive_expected_records(
  dataset = adqs,
  dataset_expected_obs = parm_visit_ref,
  by_vars = exprs(USUBJID, PARAMCD),
  set_values_to = exprs(DTYPE = "DERIVED")
)

# Example 2. visit variables are parameter dependent
parm_visit_ref <- tribble(
  ~PARAMCD, ~AVISITN, ~AVISIT,
  "a",      1, "WEEK 1",
  "a",      2, "WEEK 2",
  "b",      1, "WEEK 1"
)

derive_expected_records(
  dataset = adqs,
  dataset_expected_obs = parm_visit_ref,
  by_vars = exprs(USUBJID, PARAMCD),
  set_values_to = exprs(DTYPE = "DERIVED")
)

```

derive_extreme_event *Add the Worst or Best Observation for Each By Group as New Records*

Description

Add the first available record from events for each by group as new records, all variables of the selected observation are kept. It can be used for selecting the extreme observation from a series of user-defined events. This distinguishes `derive_extreme_event()` from `derive_extreme_records()`, where extreme records are derived based on certain order of existing variables.

Usage

```

derive_extreme_event(
  dataset,
  by_vars = NULL,
  events,
  order,
  mode,
  check_type = "warning",
  set_values_to
)

```

Arguments

dataset	Input dataset The variables specified by the order and the by_vars parameter are expected.
by_vars	Grouping variables <i>Default:</i> NULL <i>Permitted Values:</i> list of variables created by exprs()
events	Conditions and new values defining events A list of event() objects is expected. Only observations listed in the events are considered for deriving extreme event. If multiple records meet the filter condition, take the first record sorted by order.
order	Sort order If a particular event from events has more than one observation, within the event and by group, the records are ordered by the specified order. <i>Permitted Values:</i> list of variables or desc(<variable>) function calls created by exprs(), e.g., exprs(ADT, desc(AVAL))
mode	Selection mode (first or last) If a particular event from events has more than one observation, "first"/"last" is to select the first/last record of this type of events sorting by order. <i>Permitted Values:</i> "first", "last"
check_type	Check uniqueness? If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order. <i>Default:</i> "warning" <i>Permitted Values:</i> "none", "warning", "error"
set_values_to	Variables to be set The specified variables are set to the specified values for the new observations. A list of variable name-value pairs is expected. <ul style="list-style-type: none"> • LHS refers to a variable. • RHS refers to the values to set to the variable. This can be a string, a symbol, a numeric value or NA, e.g., exprs(PARAMCD = "TDOSE", PARCAT1 = "OVERALL"). More general expression are not allowed.

Details

1. Construct a dataset based on events: apply the filter condition and `set_values_to` to the input dataset.
2. For each group (with respect to the variables specified for the `by_vars` parameter) the first or last observation (with respect to the order specified for the `order` parameter and the mode specified for the `mode` parameter) is selected.
3. The variables specified by the `set_values_to` parameter are added to the selected observations.
4. The observations are added to input dataset.

Value

The input dataset with the best or worst observation of each by group added as new observations.

See Also

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_expected_records\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_rr\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```
library(tibble)

adqs <- tribble(
  ~USUBJID, ~PARAMCD,      ~AVALC,      ~ADY,
  "1",      "NO SLEEP",    "N",        1,
  "1",      "WAKE UP",     "N",        2,
  "1",      "FALL ASLEEP", "N",        3,
  "2",      "NO SLEEP",    "N",        1,
  "2",      "WAKE UP",     "Y",        2,
  "2",      "WAKE UP",     "Y",        3,
  "2",      "FALL ASLEEP", "N",        4,
  "3",      "NO SLEEP",    NA_character_, 1
)

# Add a new record for each USUBJID storing the the worst sleeping problem.
derive_extreme_event(
  adqs,
  by_vars = exprs(USUBJID),
  events = list(
    event(
      condition = PARAMCD == "NO SLEEP" & AVALC == "Y",
      set_values_to = exprs(AVALC = "No sleep", AVAL = 1)
    ),
    event(
      condition = PARAMCD == "WAKE UP" & AVALC == "Y",
      set_values_to = exprs(AVALC = "Waking up more than three times", AVAL = 2)
    )
  )
)
```

```

),
event(
  condition = PARAMCD == "FALL ASLEEP" & AVALC == "Y",
  set_values_to = exprs(AVALC = "More than 30 mins to fall asleep", AVAL = 3)
),
event(
  condition = all(AVALC == "N"),
  set_values_to = exprs(
    AVALC = "No sleeping problems", AVAL = 4
  )
),
event(
  condition = TRUE,
  set_values_to = exprs(AVALC = "Missing", AVAL = 99)
),
order = exprs(ADY),
mode = "last",
set_values_to = exprs(
  PARAMCD = "WSP",
  PARAM = "Worst Sleeping Problems"
)
)
)

```

derive_extreme_records

Add the First or Last Observation for Each By Group as New Records

Description

Add the first or last observation for each by group as new observations. It can be used for example for adding the maximum or minimum value as a separate visit. All variables of the selected observation are kept. This distinguishes `derive_extreme_records()` from `derive_summary_records()`, where only the by variables are populated for the new records.

Usage

```

derive_extreme_records(
  dataset,
  by_vars = NULL,
  order,
  mode,
  check_type = "warning",
  filter = NULL,
  set_values_to
)

```

Arguments

dataset	Input dataset
by_vars	The variables specified by the order and the by_vars parameter are expected. Grouping variables <i>Default:</i> NULL <i>Permitted Values:</i> list of variables created by <code>exprs()</code>
order	Sort order Within each by group the observations are ordered by the specified order. <i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code> , e.g., <code>exprs(ADT, desc(AVAL))</code>
mode	Selection mode (first or last) If "first" is specified, the first observation of each by group is added to the input dataset. If "last" is specified, the last observation of each by group is added to the input dataset. <i>Permitted Values:</i> "first", "last"
check_type	Check uniqueness? If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order. <i>Default:</i> "warning" <i>Permitted Values:</i> "none", "warning", "error"
filter	Filter for observations to consider Only observations fulfilling the specified condition are taken into account for selecting the first or last observation. If the parameter is not specified, all observations are considered. <i>Default:</i> NULL <i>Permitted Values:</i> a condition
set_values_to	Variables to be set The specified variables are set to the specified values for the new observations. A list of variable name-value pairs is expected. <ul style="list-style-type: none"> • LHS refers to a variable. • RHS refers to the values to set to the variable. This can be a string, a symbol, a numeric value or NA, e.g., <code>exprs(PARAMCD = "TDOSE", PARCAT1 = "OVERALL")</code>. More general expression are not allowed.

Details

1. The input dataset is restricted as specified by the filter parameter.
2. For each group (with respect to the variables specified for the by_vars parameter) the first or last observation (with respect to the order specified for the order parameter and the mode specified for the mode parameter) is selected.
3. The variables specified by the set_values_to parameter are added to the selected observations.
4. The observations are added to input dataset.

Value

The input dataset with the first or last observation of each by group added as new observations.

See Also

BDS-Findings Functions for adding Parameters/Records: `default_qtc_paramcd()`, `derive_expected_records()`, `derive_extreme_event()`, `derive_locf_records()`, `derive_param_bmi()`, `derive_param_bsa()`, `derive_param_computed()`, `derive_param_doseint()`, `derive_param_exist_flag()`, `derive_param_exposure()`, `derive_param_extreme_event()`, `derive_param_framingham()`, `derive_param_map()`, `derive_param_qtc()`, `derive_param_rr()`, `derive_param_wbc_abs()`, `derive_summary_records()`

Examples

```
library(tibble)

adlb <- tribble(
  ~USUBJID, ~AVISITN, ~AVAL, ~LBSEQ,
  "1",      1,        113,    1,
  "1",      2,        113,    2,
  "1",      3,        117,    3,
  "2",      1,        101,    1,
  "2",      2,        101,    2,
  "2",      3,         95,    3
)

# Add a new record for each USUBJID storing the minimum value (first AVAL).
# If multiple records meet the minimum criterion, take the first value by
# AVISITN. Set AVISITN = 97 and DTYPE = MINIMUM for these new records.
derive_extreme_records(
  adlb,
  by_vars = exprs(USUBJID),
  order = exprs(AVAL, AVISITN),
  mode = "first",
  filter = !is.na(AVAL),
  set_values_to = exprs(
    AVISITN = 97,
    DTYPE = "MINIMUM"
  )
)

# Add a new record for each USUBJID storing the maximum value (last AVAL).
# If multiple records meet the maximum criterion, take the first value by
# AVISITN. Set AVISITN = 98 and DTYPE = MAXIMUM for these new records.
derive_extreme_records(
  adlb,
  by_vars = exprs(USUBJID),
  order = exprs(desc(AVAL), AVISITN),
  mode = "first",
  filter = !is.na(AVAL),
  set_values_to = exprs(
    AVISITN = 98,
    DTYPE = "MAXIMUM"
  )
)
```

```

    )
  )

  # Add a new record for each USUBJID storing for the last value.
  # Set AVISITN = 99 and DTYPE = LOV for these new records.
  derive_extreme_records(
    adlb,
    by_vars = exprs(USUBJID),
    order = exprs(AVISITN),
    mode = "last",
    set_values_to = exprs(
      AVISITN = 99,
      DTYPE = "LOV"
    )
  )
)

```

derive_locf_records *Derive LOCF (Last Observation Carried Forward) Records*

Description

Adds LOCF records as new observations for each 'by group' when the dataset does not contain observations for missed visits/time points.

Usage

```

derive_locf_records(
  dataset,
  dataset_expected_obs,
  by_vars,
  analysis_var = AVAL,
  order,
  keep_vars = NULL
)

```

Arguments

dataset	Input dataset The columns specified by the by_vars, analysis_var, order, keep_vars parameters are expected.
dataset_expected_obs	Expected observations dataset Data frame with all the combinations of PARAMCD, PARAM, AVISIT, AVISITN, ... which are expected in the dataset is expected.
by_vars	Grouping variables For each group defined by by_vars those observations from dataset_expected_obs are added to the output dataset which do not have a corresponding observation in the input dataset or for which analysis_var is NA for the corresponding observation in the input dataset.

analysis_var	Analysis variable. <i>Default:</i> AVAL <i>Permitted Values:</i> a variable
order	List of variables for sorting a dataset The dataset is sorted by order before carrying the last observation forward (eg. AVAL) within each by_vars.
keep_vars	Variables that need carrying the last observation forward Keep variables that need carrying the last observation forward other than analysis_var (e.g., PARAMN, VISITNUM). If by default NULL, only variables specified in by_vars and analysis_var will be populated in the newly created records.

Details

For each group (with respect to the variables specified for the by_vars parameter) those observations from dataset_expected_obs are added to the output dataset

- which do not have a corresponding observation in the input dataset or
- for which analysis_var is NA for the corresponding observation in the input dataset.

For the new observations, analysis_var is set to the non-missing analysis_var of the previous observation in the input dataset (when sorted by order) and DTYPE is set to "LOCF".

Value

The input dataset with the new "LOCF" observations added for each by_vars. Note, a variable will only be populated in the new parameter rows if it is specified in by_vars.

Author(s)

G Gayatri

See Also

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_expected_records\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_rr\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```
library(dplyr)
library(tibble)

advs <- tribble(
  ~STUDYID, ~USUBJID, ~PARAMCD, ~PARAMN, ~AVAL, ~AVISITN, ~AVISIT,
  "CDISC01", "01-701-1015", "PULSE", 1, 61, 0, "BASELINE",
  "CDISC01", "01-701-1015", "PULSE", 1, 60, 2, "WEEK 6",
  "CDISC01", "01-701-1015", "DIABP", 2, 51, 0, "BASELINE",
```

```

"CDISC01", "01-701-1015", "DIABP",      2,   50,      2, "WEEK 2",
"CDISC01", "01-701-1015", "DIABP",      2,   51,      4, "WEEK 4",
"CDISC01", "01-701-1015", "DIABP",      2,   50,      6, "WEEK 6",
"CDISC01", "01-701-1015", "SYSBP",      3,  121,      0, "BASELINE",
"CDISC01", "01-701-1015", "SYSBP",      3,  121,      2, "WEEK 2",
"CDISC01", "01-701-1015", "SYSBP",      3,  121,      4, "WEEK 4",
"CDISC01", "01-701-1015", "SYSBP",      3,  121,      6, "WEEK 6",
"CDISC01", "01-701-1028", "PULSE",       1,   65,      0, "BASELINE",
"CDISC01", "01-701-1028", "DIABP",      2,   79,      0, "BASELINE",
"CDISC01", "01-701-1028", "DIABP",      2,   80,      2, "WEEK 2",
"CDISC01", "01-701-1028", "DIABP",      2,   NA,      4, "WEEK 4",
"CDISC01", "01-701-1028", "DIABP",      2,   NA,      6, "WEEK 6",
"CDISC01", "01-701-1028", "SYSBP",      3,  130,      0, "BASELINE",
"CDISC01", "01-701-1028", "SYSBP",      3,  132,      2, "WEEK 2"
)

# A dataset with all the combinations of PARAMCD, PARAM, AVISIT, AVISITN, ... which are expected.
advs_expected_obsv <- tribble(
  ~PARAMCD, ~AVISITN, ~AVISIT,
  "PULSE",      0, "BASELINE",
  "PULSE",      6, "WEEK 6",
  "DIABP",      0, "BASELINE",
  "DIABP",      2, "WEEK 2",
  "DIABP",      4, "WEEK 4",
  "DIABP",      6, "WEEK 6",
  "SYSBP",      0, "BASELINE",
  "SYSBP",      2, "WEEK 2",
  "SYSBP",      4, "WEEK 4",
  "SYSBP",      6, "WEEK 6"
)

derive_locf_records(
  data = advs,
  dataset_expected_obs = advs_expected_obsv,
  by_vars = exprs(STUDYID, USUBJID, PARAMCD),
  order = exprs(AVISITN, AVISIT),
  keep_vars = exprs(PARAMN)
)

```

```
derive_param_bmi
```

```
Adds a Parameter for BMI
```

Description

Adds a record for BMI/Body Mass Index using Weight and Height each by group (e.g., subject and visit) where the source parameters are available.

Usage

```

derive_param_bmi(
  dataset,
  by_vars,
  set_values_to = exprs(PARAMCD = "BMI"),
  weight_code = "WEIGHT",
  height_code = "HEIGHT",
  get_unit_expr,
  filter = NULL
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>weight_code</code> and <code>height_code</code>.</p>
by_vars	<p>Grouping variables</p> <p>For each group defined by <code>by_vars</code> an observation is added to the output dataset. Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p><i>Permitted Values:</i> list of variables</p>
set_values_to	<p>Variables to be set</p> <p>The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter.</p> <p><i>Permitted Values:</i> List of variable-value pairs</p>
weight_code	<p>WEIGHT parameter code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the <code>WEIGHT</code>. It is expected that <code>WEIGHT</code> is measured in kg</p> <p><i>Permitted Values:</i> character value</p>
height_code	<p>HEIGHT parameter code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the <code>HEIGHT</code>. It is expected that <code>HEIGHT</code> is measured in cm</p> <p><i>Permitted Values:</i> character value</p>
get_unit_expr	<p>An expression providing the unit of the parameter</p> <p>The result is used to check the units of the input parameters.</p> <p><i>Permitted Values:</i> A variable of the input dataset or a function call</p>
filter	<p>Filter condition</p> <p>The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account.</p> <p><i>Permitted Values:</i> a condition</p>

Details

The analysis value of the new parameter is derived as

$$BMI = \frac{WEIGHT}{HEIGHT^2}$$

Value

The input dataset with the new parameter added. Note, a variable will only be populated in the new parameter rows if it is specified in `by_vars`.

See Also

BDS-Findings Functions for adding Parameters/Records: `default_qtc_paramcd()`, `derive_expected_records()`, `derive_extreme_event()`, `derive_extreme_records()`, `derive_locf_records()`, `derive_param_bsa()`, `derive_param_computed()`, `derive_param_doseint()`, `derive_param_exist_flag()`, `derive_param_exposure()`, `derive_param_extreme_event()`, `derive_param_framingham()`, `derive_param_map()`, `derive_param_qtc()`, `derive_param_rr()`, `derive_param_wbc_abs()`, `derive_summary_records()`

Examples

```
library(tibble)

advs <- tribble(
  ~USUBJID,    ~PARAMCD, ~PARAM,      ~AVAL, ~AVISIT,
  "01-701-1015", "HEIGHT", "Height (cm)", 147,  "SCREENING",
  "01-701-1015", "WEIGHT", "Weight (kg)", 54.0, "SCREENING",
  "01-701-1015", "WEIGHT", "Weight (kg)", 54.4, "BASELINE",
  "01-701-1015", "WEIGHT", "Weight (kg)", 53.1, "WEEK 2",
  "01-701-1028", "HEIGHT", "Height (cm)", 163,  "SCREENING",
  "01-701-1028", "WEIGHT", "Weight (kg)", 78.5, "SCREENING",
  "01-701-1028", "WEIGHT", "Weight (kg)", 80.3, "BASELINE",
  "01-701-1028", "WEIGHT", "Weight (kg)", 80.7,  "WEEK 2"
)

derive_param_bmi(
  advs,
  by_vars = exprs(USUBJID, AVISIT),
  weight_code = "WEIGHT",
  height_code = "HEIGHT",
  set_values_to = exprs(
    PARAMCD = "BMI",
    PARAM = "Body Mass Index (kg/m^2)"
  ),
  get_unit_expr = extract_unit(PARAM)
)
```

derive_param_bsa	<i>Adds a Parameter for BSA (Body Surface Area) Using the Specified Method</i>
------------------	--

Description

Adds a record for BSA (Body Surface Area) using the specified derivation method for each by group (e.g., subject and visit) where the source parameters are available.

Usage

```
derive_param_bsa(
  dataset,
  by_vars,
  method,
  set_values_to = exprs(PARAMCD = "BSA"),
  height_code = "HEIGHT",
  weight_code = "WEIGHT",
  get_unit_expr,
  filter = NULL
)
```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>HEIGHT</code> and <code>WEIGHT</code>.</p>
by_vars	<p>Grouping variables</p> <p>For each group defined by <code>by_vars</code> an observation is added to the output dataset. Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p><i>Permitted Values:</i> list of variables</p>
method	<p>Derivation method to use. Note that <code>HEIGHT</code> is expected in cm and <code>WEIGHT</code> is expected in kg:</p> <p>Mosteller: $\sqrt{\text{height} * \text{weight} / 3600}$</p> <p>DuBois-DuBois: $0.20247 * (\text{height}/100) ^ 0.725 * \text{weight} ^ 0.425$</p> <p>Haycock: $0.024265 * \text{height} ^ 0.3964 * \text{weight} ^ 0.5378$</p> <p>Gehan-George: $0.0235 * \text{height} ^ 0.42246 * \text{weight} ^ 0.51456$</p> <p>Boyd: $0.0003207 * (\text{height} ^ 0.3) * (1000 * \text{weight}) ^ (0.7285 - (0.0188 * \log_{10}(1000 * \text{weight})))$</p> <p>Fujimoto: $0.008883 * \text{height} ^ 0.663 * \text{weight} ^ 0.444$</p> <p>Takahira: $0.007241 * \text{height} ^ 0.725 * \text{weight} ^ 0.425$</p> <p><i>Permitted Values:</i> character value</p>

set_values_to	<p>Variables to be set</p> <p>The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter.</p> <p><i>Permitted Values:</i> List of variable-value pairs</p>
height_code	<p>HEIGHT parameter code</p> <p>The observations where PARAMCD equals the specified value are considered as the HEIGHT assessments. It is expected that HEIGHT is measured in cm.</p> <p><i>Permitted Values:</i> character value</p>
weight_code	<p>WEIGHT parameter code</p> <p>The observations where PARAMCD equals the specified value are considered as the WEIGHT assessments. It is expected that WEIGHT is measured in kg.</p> <p><i>Permitted Values:</i> character value</p>
get_unit_expr	<p>An expression providing the unit of the parameter</p> <p>The result is used to check the units of the input parameters.</p> <p><i>Permitted Values:</i> A variable of the input dataset or a function call</p>
filter	<p>Filter condition</p> <p>The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account.</p> <p><i>Permitted Values:</i> a condition</p>

Value

The input dataset with the new parameter added. Note, a variable will only be populated in the new parameter rows if it is specified in `by_vars`.

See Also

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_expected_records\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_rr\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```
library(tibble)

advs <- tribble(
  ~USUBJID, ~PARAMCD, ~PARAM, ~AVAL, ~VISIT,
  "01-701-1015", "HEIGHT", "Height (cm)", 170, "BASELINE",
  "01-701-1015", "WEIGHT", "Weight (kg)", 75, "BASELINE",
  "01-701-1015", "WEIGHT", "Weight (kg)", 78, "MONTH 1",
  "01-701-1015", "WEIGHT", "Weight (kg)", 80, "MONTH 2",
  "01-701-1028", "HEIGHT", "Height (cm)", 185, "BASELINE",
  "01-701-1028", "WEIGHT", "Weight (kg)", 90, "BASELINE",
  "01-701-1028", "WEIGHT", "Weight (kg)", 88, "MONTH 1",
```

```

    "01-701-1028", "WEIGHT", "Weight (kg)", 85, "MONTH 2",
  )

  derive_param_bsa(
    advs,
    by_vars = exprs(USUBJID, VISIT),
    method = "Mosteller",
    set_values_to = exprs(
      PARAMCD = "BSA",
      PARAM = "Body Surface Area (m^2)"
    ),
    get_unit_expr = extract_unit(PARAM)
  )

  derive_param_bsa(
    advs,
    by_vars = exprs(USUBJID, VISIT),
    method = "Fujimoto",
    set_values_to = exprs(
      PARAMCD = "BSA",
      PARAM = "Body Surface Area (m^2)"
    ),
    get_unit_expr = extract_unit(PARAM)
  )

```

derive_param_computed *Adds a Parameter Computed from the Analysis Value of Other Parameters*

Description

Adds a parameter computed from the analysis value of other parameters. It is expected that the analysis value of the new parameter is defined by an expression using the analysis values of other parameters. For example mean arterial pressure (MAP) can be derived from systolic (SYSBP) and diastolic blood pressure (DIABP) with the formula

$$MAP = \frac{SYSBP + 2DIABP}{3}$$

Usage

```

derive_param_computed(
  dataset,
  by_vars,
  parameters,
  analysis_value,
  set_values_to,
  filter = NULL,
  constant_by_vars = NULL,
  constant_parameters = NULL
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>parameters</code>.</p>
by_vars	<p>Grouping variables</p> <p>For each group defined by <code>by_vars</code> an observation is added to the output dataset. Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p><i>Permitted Values:</i> list of variables</p>
parameters	<p>Required parameter codes</p> <p>It is expected that all parameter codes (<code>PARAMCD</code>) which are required to derive the new parameter are specified for this parameter or the <code>constant_parameters</code> parameter.</p> <p><i>Permitted Values:</i> A character vector of <code>PARAMCD</code> values</p>
analysis_value	<p>Definition of the analysis value</p> <p>An expression defining the analysis value (<code>AVAL</code>) of the new parameter is expected. The analysis values of the parameters specified by <code>parameters</code> can be accessed using <code>AVAL.<parameter code></code>, e.g., <code>AVAL.SYSBP</code>.</p> <p><i>Permitted Values:</i> An unquoted expression</p>
set_values_to	<p>Variables to be set</p> <p>The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter.</p> <p><i>Permitted Values:</i> List of variable-value pairs</p>
filter	<p>Filter condition</p> <p>The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account.</p> <p><i>Permitted Values:</i> a condition</p>
constant_by_vars	<p>By variables for constant parameters</p> <p>The constant parameters (parameters that are measured only once) are merged to the other parameters using the specified variables. (Refer to Example 2)</p> <p><i>Permitted Values:</i> list of variables</p>
constant_parameters	<p>Required constant parameter codes</p> <p>It is expected that all the parameter codes (<code>PARAMCD</code>) which are required to derive the new parameter and are measured only once are specified here. For example if <code>BMI</code> should be derived and <code>height</code> is measured only once while <code>weight</code> is measured at each visit. <code>height</code> could be specified in the <code>constant_parameters</code> parameter. (Refer to Example 2)</p> <p><i>Permitted Values:</i> A character vector of <code>PARAMCD</code> values</p>

Details

For each group (with respect to the variables specified for the `by_vars` parameter) an observation is added to the output dataset if the filtered input dataset contains exactly one observation for each parameter code specified for parameters.

For the new observations AVAL is set to the value specified by `analysis_value` and the variables specified for `set_values_to` are set to the provided values. The values of the other variables of the input dataset are set to NA.

Value

The input dataset with the new parameter added. Note, a variable will only be populated in the new parameter rows if it is specified in `by_vars`.

See Also

BDS-Findings Functions for adding Parameters/Records: `default_qtc_paramcd()`, `derive_expected_records()`, `derive_extreme_event()`, `derive_extreme_records()`, `derive_locf_records()`, `derive_param_bmi()`, `derive_param_bsa()`, `derive_param_doseint()`, `derive_param_exist_flag()`, `derive_param_exposure()`, `derive_param_extreme_event()`, `derive_param_framingham()`, `derive_param_map()`, `derive_param_qtc()`, `derive_param_rr()`, `derive_param_wbc_abs()`, `derive_summary_records()`

Examples

```
library(tibble)

# Example 1: Derive MAP
advs <- tribble(
  ~USUBJID, ~PARAMCD, ~PARAM, ~AVAL, ~AVALU, ~VISIT,
  "01-701-1015", "DIABP", "Diastolic Blood Pressure (mmHg)", 51, "mmHg", "BASELINE",
  "01-701-1015", "DIABP", "Diastolic Blood Pressure (mmHg)", 50, "mmHg", "WEEK 2",
  "01-701-1015", "SYSBP", "Systolic Blood Pressure (mmHg)", 121, "mmHg", "BASELINE",
  "01-701-1015", "SYSBP", "Systolic Blood Pressure (mmHg)", 121, "mmHg", "WEEK 2",
  "01-701-1028", "DIABP", "Diastolic Blood Pressure (mmHg)", 79, "mmHg", "BASELINE",
  "01-701-1028", "DIABP", "Diastolic Blood Pressure (mmHg)", 80, "mmHg", "WEEK 2",
  "01-701-1028", "SYSBP", "Systolic Blood Pressure (mmHg)", 130, "mmHg", "BASELINE",
  "01-701-1028", "SYSBP", "Systolic Blood Pressure (mmHg)", 132, "mmHg", "WEEK 2"
)

derive_param_computed(
  advs,
  by_vars = exprs(USUBJID, VISIT),
  parameters = c("SYSBP", "DIABP"),
  analysis_value = (AVAL.SYSBP + 2 * AVAL.DIABP) / 3,
  set_values_to = exprs(
    PARAMCD = "MAP",
    PARAM = "Mean Arterial Pressure (mmHg)",
    AVALU = "mmHg"
  )
)

# Example 2: Derive BMI where height is measured only once
```

```

advs <- tribble(
  ~USUBJID, ~PARAMCD, ~PARAM, ~AVAL, ~AVALU, ~VISIT,
  "01-701-1015", "HEIGHT", "Height (cm)", 147, "cm", "SCREENING",
  "01-701-1015", "WEIGHT", "Weight (kg)", 54.0, "kg", "SCREENING",
  "01-701-1015", "WEIGHT", "Weight (kg)", 54.4, "kg", "BASELINE",
  "01-701-1015", "WEIGHT", "Weight (kg)", 53.1, "kg", "WEEK 2",
  "01-701-1028", "HEIGHT", "Height (cm)", 163, "cm", "SCREENING",
  "01-701-1028", "WEIGHT", "Weight (kg)", 78.5, "kg", "SCREENING",
  "01-701-1028", "WEIGHT", "Weight (kg)", 80.3, "kg", "BASELINE",
  "01-701-1028", "WEIGHT", "Weight (kg)", 80.7, "kg", "WEEK 2"
)

derive_param_computed(
  advs,
  by_vars = exprs(USUBJID, VISIT),
  parameters = "WEIGHT",
  analysis_value = AVAL.WEIGHT / (AVAL.HEIGHT / 100)^2,
  set_values_to = exprs(
    PARAMCD = "BMI",
    PARAM = "Body Mass Index (kg/m^2)",
    AVALU = "kg/m^2"
  ),
  constant_parameters = c("HEIGHT"),
  constant_by_vars = exprs(USUBJID)
)

```

derive_param_doseint *Adds a Parameter for Dose Intensity*

Description

Adds a record for the dose intensity for each by group (e.g., subject and visit) where the source parameters are available.

Usage

```

derive_param_doseint(
  dataset,
  by_vars,
  set_values_to = exprs(PARAMCD = "TNDOSINT"),
  tadm_code = "TNDOSE",
  tpadm_code = "TSNDOSE",
  zero_doses = "Inf",
  filter = NULL
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>tadm_code</code> and <code>padm_code</code>.</p>
by_vars	<p>Grouping variables</p> <p>Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p>Permitted Values: list of variables</p>
set_values_to	<p>Variables to be set</p> <p>The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter.</p> <p><i>Permitted Values:</i> List of variable-value pairs</p>
tadm_code	<p>Total Doses Administered parameter code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the total dose administered. The <code>AVAL</code> associated with this <code>PARAMCD</code> will be the numerator of the dose intensity calculation.</p> <p>Permitted Values: character value</p>
tpadm_code	<p>Total Doses Planned parameter code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the total planned dose. The <code>AVAL</code> associated with this <code>PARAMCD</code> will be the denominator of the dose intensity calculation.</p> <p>Permitted Values: character value</p>
zero_doses	<p>Flag indicating logic for handling 0 planned or administered doses for a <code>by_vars</code> group</p> <p>Default: Inf</p> <p>Permitted Values: Inf, 100</p> <p>No record is returned if either the planned (<code>tpadm_code</code>) or administered (<code>tadm_code</code>) <code>AVAL</code> are NA. No record is returned if a record does not exist for both <code>tadm_code</code> and <code>tpadm_code</code> for the specified <code>by_var</code>.</p> <p>If <code>zero_doses = Inf</code>:</p> <ol style="list-style-type: none"> 1. If the planned dose (<code>tpadm_code</code>) is 0 and administered dose (<code>tadm_code</code>) is 0, NaN is returned. 2. If the planned dose (<code>tpadm_code</code>) is 0 and the administered dose (<code>tadm_code</code>) is > 0, Inf is returned. <p>If <code>zero_doses = 100</code>:</p> <ol style="list-style-type: none"> 1. If the planned dose (<code>tpadm_code</code>) is 0 and administered dose (<code>tadm_code</code>) is 0, 0 is returned. 2. If the planned dose (<code>tpadm_code</code>) is 0 and the administered dose (<code>tadm_code</code>) is > 0, 100 is returned.

filter Filter condition
 The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account.
Permitted Values: a condition

Details

The analysis value of the new parameter is derived as Total Dose / Planned Dose * 100

Value

The input dataset with the new parameter rows added. Note, a variable will only be populated in the new parameter rows if it is specified in `by_vars`.

See Also

BDS-Findings Functions for adding Parameters/Records: `default_qtc_paramcd()`, `derive_expected_records()`, `derive_extreme_event()`, `derive_extreme_records()`, `derive_locf_records()`, `derive_param_bmi()`, `derive_param_bsa()`, `derive_param_computed()`, `derive_param_exist_flag()`, `derive_param_exposure()`, `derive_param_extreme_event()`, `derive_param_framingham()`, `derive_param_map()`, `derive_param_qtc()`, `derive_param_rr()`, `derive_param_wbc_abs()`, `derive_summary_records()`

Examples

```
library(tibble)
library(lubridate, warn.conflicts = FALSE)

adex <- tribble(
  ~USUBJID, ~PARAMCD, ~VISIT, ~ANL01FL, ~ASTDT, ~AENDT, ~AVAL,
  "P001", "TNDOSE", "V1", "Y", ymd("2020-01-01"), ymd("2020-01-30"), 59,
  "P001", "TSNDOSE", "V1", "Y", ymd("2020-01-01"), ymd("2020-02-01"), 96,
  "P001", "TNDOSE", "V2", "Y", ymd("2020-02-01"), ymd("2020-03-15"), 88,
  "P001", "TSNDOSE", "V2", "Y", ymd("2020-02-05"), ymd("2020-03-01"), 88,
  "P002", "TNDOSE", "V1", "Y", ymd("2021-01-01"), ymd("2021-01-30"), 0,
  "P002", "TSNDOSE", "V1", "Y", ymd("2021-01-01"), ymd("2021-02-01"), 0,
  "P002", "TNDOSE", "V2", "Y", ymd("2021-02-01"), ymd("2021-03-15"), 52,
  "P002", "TSNDOSE", "V2", "Y", ymd("2021-02-05"), ymd("2021-03-01"), 0
)

derive_param_doseint(
  adex,
  by_vars = exprs(USUBJID, VISIT),
  set_values_to = exprs(PARAMCD = "TNDOSINT"),
  tadm_code = "TNDOSE",
  tpadm_code = "TSNDOSE"
)

derive_param_doseint(
  adex,
  by_vars = exprs(USUBJID, VISIT),
  set_values_to = exprs(PARAMCD = "TDOSINT2"),
```

```

tadm_code = "TNDOSE",
tpadm_code = "TSNDOSE",
zero_doses = "100"
)

```

```
derive_param_exist_flag
```

Add an Existence Flag Parameter

Description

Add a new parameter indicating that a certain event exists in a dataset. AVALC and AVAL indicate if an event occurred or not. For example, the function can derive a parameter indicating if there is measurable disease at baseline.

Usage

```

derive_param_exist_flag(
  dataset = NULL,
  dataset_adsl,
  dataset_add,
  condition,
  true_value = "Y",
  false_value = NA_character_,
  missing_value = NA_character_,
  filter_add = NULL,
  aval_fun = yn_to_numeric,
  subject_keys = get_admiral_option("subject_keys"),
  set_values_to
)

```

Arguments

dataset	Input dataset The variables specified for subject_keys and the PARAMCD variable are expected.
dataset_adsl	ADSL input dataset The variables specified for subject_keys are expected. For each subject (as defined by subject_keys) from the specified dataset (dataset_adsl), the existence flag is calculated and added as a new observation to the input datasets (dataset)
dataset_add	Additional dataset The variables specified by the subject_keys parameter are expected. This dataset is used to check if an event occurred or not. Any observation in the dataset fulfilling the event condition (condition) is considered as an event.

condition	<p>Event condition</p> <p>The condition is evaluated at the additional dataset (<code>dataset_add</code>).</p> <p>For all subjects where it evaluates as TRUE at least once AVALC is set to the true value (<code>true_value</code>) for the new observations.</p> <p>For all subjects where it evaluates as FALSE or NA for all observations AVALC is set to the false value (<code>false_value</code>).</p> <p>For all subjects not present in the additional dataset AVALC is set to the missing value (<code>missing_value</code>).</p>
true_value	<p>True value</p> <p>For all subjects with at least one observations in the additional dataset (<code>dataset_add</code>) fulfilling the event condition (<code>condition</code>), AVALC is set to the specified value (<code>true_value</code>).</p> <p><i>Default:</i> "Y"</p> <p><i>Permitted Value:</i> A character scalar</p>
false_value	<p>False value</p> <p>For all subjects with at least one observations in the additional dataset (<code>dataset_add</code>) but none of them is fulfilling the event condition (<code>condition</code>), AVALC is set to the specified value (<code>false_value</code>).</p> <p><i>Default:</i> NA_character_</p> <p><i>Permitted Value:</i> A character scalar</p>
missing_value	<p>Values used for missing information</p> <p>For all subjects without an observation in the additional dataset (<code>dataset_add</code>), AVALC is set to the specified value (<code>missing_value</code>).</p> <p><i>Default:</i> NA_character_</p> <p><i>Permitted Value:</i> A character scalar</p>
filter_add	<p>Filter for additional data</p> <p>Only observations fulfilling the specified condition are taken into account for flagging. If the parameter is not specified, all observations are considered.</p> <p><i>Permitted Values:</i> a condition</p>
aval_fun	<p>Function to map character analysis value (AVALC) to numeric analysis value (AVAL)</p> <p>The (first) argument of the function must expect a character vector and the function must return a numeric vector.</p> <p><i>Default:</i> <code>yn_to_numeric</code> (see <code>yn_to_numeric()</code> for details)</p>
subject_keys	<p>Variables to uniquely identify a subject</p> <p>A list of symbols created using <code>exprs()</code> is expected.</p>
set_values_to	<p>Variables to set</p> <p>A named list returned by <code>exprs()</code> defining the variables to be set for the new parameter, e.g. <code>exprs(PARAMCD = "MDIS", PARAM = "Measurable Disease at Baseline")</code> is expected. The values must be symbols, character strings, numeric values, or NA.</p>

Details

1. The additional dataset (`dataset_add`) is restricted to the observations matching the `filter_add` condition.
2. For each subject in `dataset_ads1` a new observation is created.
 - The `AVALC` variable is added and set to the true value (`true_value`) if for the subject at least one observation exists in the (restricted) additional dataset where the condition evaluates to `TRUE`.
 - It is set to the false value (`false_value`) if for the subject at least one observation exists and for all observations the condition evaluates to `FALSE` or `NA`.
 - Otherwise, it is set to the missing value (`missing_value`), i.e., for those subject not in `dataset_add`.
3. The `AVAL` variable is added and set to `aval_fun(AVALC)`.
4. The variables specified by the `set_values_to` parameter are added to the new observations.
5. The new observations are added to input dataset.

Value

The input dataset with a new parameter indicating if an event occurred (`AVALC`, `AVAL`, and the variables specified by `subject_keys` and `set_value_to` are populated for the new parameter)

See Also

BDS-Findings Functions for adding Parameters/Records: `default_qtc_paramcd()`, `derive_expected_records()`, `derive_extreme_event()`, `derive_extreme_records()`, `derive_locf_records()`, `derive_param_bmi()`, `derive_param_bsa()`, `derive_param_computed()`, `derive_param_doseint()`, `derive_param_exposure()`, `derive_param_extreme_event()`, `derive_param_framingham()`, `derive_param_map()`, `derive_param_qtc()`, `derive_param_rr()`, `derive_param_wbc_abs()`, `derive_summary_records()`

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

# Derive a new parameter for measurable disease at baseline
adsl <- tribble(
  ~USUBJID,
  "1",
  "2",
  "3"
) %>%
  mutate(STUDYID = "XX1234")

tu <- tribble(
  ~USUBJID, ~VISIT, ~TUSTRESC,
  "1", "SCREENING", "TARGET",
  "1", "WEEK 1", "TARGET",
  "1", "WEEK 5", "TARGET",
```

```

    "1",      "WEEK 9",      "NON-TARGET",
    "2",      "SCREENING",  "NON-TARGET",
    "2",      "SCREENING",  "NON-TARGET"
  ) %>%
  mutate(
    STUDYID = "XX1234",
    TUTESTCD = "TUMIDENT"
  )

derive_param_exist_flag(
  dataset_adsl = adsl,
  dataset_add = tu,
  filter_add = TUTESTCD == "TUMIDENT" & VISIT == "SCREENING",
  condition = TUSTRESC == "TARGET",
  false_value = "N",
  missing_value = "N",
  set_values_to = exprs(
    PARAMCD = "MDIS",
    PARAM = "Measurable Disease at Baseline"
  )
)

```

derive_param_exposure *Add an Aggregated Parameter and Derive the Associated Start and End Dates*

Description

Add a record computed from the aggregated analysis value of another parameter and compute the start (ASTDT(M)) and end date (AENDT(M)) as the minimum and maximum date by by_vars.

Usage

```

derive_param_exposure(
  dataset,
  by_vars,
  input_code,
  analysis_var,
  summary_fun,
  filter = NULL,
  set_values_to = NULL
)

```

Arguments

dataset	Input dataset
---------	---------------

- The variables specified by the by_vars, analysis_var parameters and PARAMCD are expected,
- Either ASTDTM and AENDTM or ASTDT and AENDT are also expected.

by_vars	<p>Grouping variables</p> <p>For each group defined by by_vars an observation is added to the output dataset. Only variables specified in by_vars will be populated in the newly created records.</p> <p><i>Permitted Values:</i> list of variables</p>
input_code	<p>Required parameter code</p> <p>The observations where PARAMCD equals the specified value are considered to compute the summary record.</p> <p><i>Permitted Values:</i> A character of PARAMCD value</p>
analysis_var	Analysis variable.
summary_fun	<p>Function that takes as an input the analysis_var and performs the calculation. This can include built-in functions as well as user defined functions, for example mean or function(x) mean(x, na.rm = TRUE).</p>
filter	<p>Filter condition</p> <p>The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account.</p> <p><i>Permitted Values:</i> a condition</p>
set_values_to	<p>Variable-value pairs</p> <p>Set a list of variables to some specified value for the new observation(s)</p> <ul style="list-style-type: none"> • LHS refer to a variable. It is expected that at least PARAMCD is defined. • RHS refers to the values to set to the variable. This can be a string, a symbol, a numeric value or NA. (e.g. exprs(PARAMCD = "TDOSE", PARCAT1 = "OVERALL")). More general expression are not allowed. <p><i>Permitted Values:</i> List of variable-value pairs</p>

Details

For each group (with respect to the variables specified for the by_vars parameter), an observation is added to the output dataset and the defined values are set to the defined variables

Value

The input dataset with a new record added for each group (with respect to the variables specified for the by_vars parameter). That is, a variable will only be populated in this new record if it is specified in by_vars. For each new record,

- the variable specified analysis_var is computed as defined by summary_fun,
- the variable(s) specified on the LHS of set_values_to are set to their paired value (RHS). In addition, the start and end date are computed as the minimum/maximum dates by by_vars.

If the input datasets contains

- both AxxDTM and AxxDT then all ASTDTM,AENDTM, ASTDT, AENDT are computed
- only AxxDTM then ASTDTM,AENDTM are computed
- only AxxDT then ASTDT,AENDT are computed.

See Also

BDS-Findings Functions for adding Parameters/Records: `default_qtc_paramcd()`, `derive_expected_records()`, `derive_extreme_event()`, `derive_extreme_records()`, `derive_locf_records()`, `derive_param_bmi()`, `derive_param_bsa()`, `derive_param_computed()`, `derive_param_doseint()`, `derive_param_exist_flag()`, `derive_param_extreme_event()`, `derive_param_framingham()`, `derive_param_map()`, `derive_param_qtc()`, `derive_param_rr()`, `derive_param_wbc_abs()`, `derive_summary_records()`

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate, warn.conflicts = FALSE)
library(stringr, warn.conflicts = FALSE)
adex <- tribble(
  ~USUBJID, ~PARAMCD, ~AVAL, ~AVALC, ~VISIT, ~ASTDT, ~AENDT,
  "1015", "DOSE", 80, NA_character_, "BASELINE", ymd("2014-01-02"), ymd("2014-01-16"),
  "1015", "DOSE", 85, NA_character_, "WEEK 2", ymd("2014-01-17"), ymd("2014-06-18"),
  "1015", "DOSE", 82, NA_character_, "WEEK 24", ymd("2014-06-19"), ymd("2014-07-02"),
  "1015", "ADJ", NA, NA_character_, "BASELINE", ymd("2014-01-02"), ymd("2014-01-16"),
  "1015", "ADJ", NA, NA_character_, "WEEK 2", ymd("2014-01-17"), ymd("2014-06-18"),
  "1015", "ADJ", NA, NA_character_, "WEEK 24", ymd("2014-06-19"), ymd("2014-07-02"),
  "1017", "DOSE", 80, NA_character_, "BASELINE", ymd("2014-01-05"), ymd("2014-01-19"),
  "1017", "DOSE", 50, NA_character_, "WEEK 2", ymd("2014-01-20"), ymd("2014-05-10"),
  "1017", "DOSE", 65, NA_character_, "WEEK 24", ymd("2014-05-10"), ymd("2014-07-02"),
  "1017", "ADJ", NA, NA_character_, "BASELINE", ymd("2014-01-05"), ymd("2014-01-19"),
  "1017", "ADJ", NA, "ADVERSE EVENT", "WEEK 2", ymd("2014-01-20"), ymd("2014-05-10"),
  "1017", "ADJ", NA, NA_character_, "WEEK 24", ymd("2014-05-10"), ymd("2014-07-02")
) %>%
mutate(ASTDTM = ymd_hms(paste(ASTDT, "00:00:00")), AENDTM = ymd_hms(paste(AENDT, "00:00:00")))

# Cumulative dose
adex %>%
  derive_param_exposure(
    by_vars = exprs(USUBJID),
    set_values_to = exprs(PARAMCD = "TDOSE", PARCAT1 = "OVERALL"),
    input_code = "DOSE",
    analysis_var = AVAL,
    summary_fun = function(x) sum(x, na.rm = TRUE)
  ) %>%
  select(-ASTDTM, -AENDTM)

# average dose in w2-24
adex %>%
  derive_param_exposure(
    by_vars = exprs(USUBJID),
    filter = VISIT %in% c("WEEK 2", "WEEK 24"),
    set_values_to = exprs(PARAMCD = "AVDW224", PARCAT1 = "WEEK2-24"),
    input_code = "DOSE",
    analysis_var = AVAL,
    summary_fun = function(x) mean(x, na.rm = TRUE)
  ) %>%
  select(-ASTDTM, -AENDTM)
```

```
# Any dose adjustment?
adex %>%
  derive_param_exposure(
    by_vars = exprs(USUBJID),
    set_values_to = exprs(PARAMCD = "TADJ", PARCAT1 = "OVERALL"),
    input_code = "ADJ",
    analysis_var = AVALC,
    summary_fun = function(x) if_else(sum(!is.na(x)) > 0, "Y", NA_character_)
  ) %>%
  select(-ASTDTM, -AENDTM)
```

```
derive_param_extreme_event
```

Add an Extreme Event Parameter

Description

Add a new parameter for the first or last event occurring in a dataset. The variable given in `new_var` indicates if an event occurred or not. For example, the function can derive a parameter for the first disease progression.

Usage

```
derive_param_extreme_event(
  dataset = NULL,
  dataset_adsl,
  dataset_source,
  filter_source,
  order = NULL,
  new_var = NULL,
  true_value = "Y",
  false_value = "N",
  mode = "first",
  subject_keys = get_admiral_option("subject_keys"),
  set_values_to,
  check_type = "warning"
)
```

Arguments

<code>dataset</code>	Input dataset The PARAMCD variable is expected.
<code>dataset_adsl</code>	ADSL input dataset The variables specified for <code>subject_keys</code> are expected. For each observation of the specified dataset a new observation is added to the input dataset.

dataset_source	Source dataset All observations in the specified dataset fulfilling the condition specified by filter_source are considered as an event. The variables specified by the subject_keys and order parameter (if applicable) are expected.
filter_source	Source filter All observations in dataset_source fulfilling the specified condition are considered as an event. For subjects with at least one event new_var is set to true_value. For all other subjects new_var is set to false_value.
order	Order variable List of symbols for sorting the source dataset (dataset_source). <i>Permitted Values:</i> list of variables or desc(<variable>) function calls created by exprs(), e.g., exprs(ADT, desc(AVAL)).
new_var	New variable The name of the variable which will indicate whether an event happened or not.
true_value	True value For all subjects with at least one observation in the source dataset (dataset_source) fulfilling the event condition (filter_source), new_var is set to the specified value true_value.
false_value	False value For all other subjects in dataset_adsl without an event, new_var is set to the specified value false_value.
mode	Selection mode (first or last) If "first" is specified, the first observation of each subject is selected. If "last" is specified, the last observation of each subject is selected. <i>Permitted Values:</i> "first", "last"
subject_keys	Variables to uniquely identify a subject A list of symbols created using exprs() is expected.
set_values_to	Variables to set A named list returned by exprs() defining the variables to be set for the new parameter, e.g. exprs(PARAMCD = "PD", PARAM = "Disease Progression") is expected. The values must be symbols, character strings, numeric values, or NA. Note, if you require a date or datetime variable to be populated, this needs to be defined here.
check_type	Check uniqueness? If "warning" or "error" is specified, a message is issued if the observations of the input dataset restricted to the source parameter (source_param) are not unique with respect to the subject keys (subject_key parameter) and order variables (order parameter). <i>Permitted Values:</i> "none", "warning", "error"

Details

1. The source dataset (dataset_source) is restricted to observations fulfilling filter_source.
2. For each subject (with respect to the variables specified for the subject_keys parameter) either the first or last observation from the restricted source dataset is selected. This is depending on mode, (with respect to order, if applicable) where the event condition (filter_source parameter) is fulfilled.
3. For each observation in dataset_adsl a new observation is created. For subjects with event new_var is set to true_value. For all other subjects new_var is set to false_value. For subjects with event all variables from dataset_source are kept. For subjects without event all variables which are in both dataset_adsl and dataset_source are kept.
4. The variables specified by the set_values_to parameter are added to the new observations.
5. The new observations are added to input dataset.

Value

The input dataset with a new parameter indicating if and when an event occurred

See Also

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_expected_records\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_rr\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

# Derive a new parameter for the first disease progression (PD)
adsl <- tribble(
  ~USUBJID, ~DTHDT,
  "1",      ymd("2022-05-13"),
  "2",      ymd(""),
  "3",      ymd("")
) %>%
  mutate(STUDYID = "XX1234")

adrs <- tribble(
  ~USUBJID, ~ADTC,      ~AVALC,
  "1",      "2020-01-02", "PR",
  "1",      "2020-02-01", "CR",
  "1",      "2020-03-01", "CR",
  "1",      "2020-04-01", "SD",
  "2",      "2021-06-15", "SD",
  "2",      "2021-07-16", "PD",
  "2",      "2021-09-14", "PD"
```

```

) %>%
  mutate(
    STUDYID = "XX1234",
    ADT = ymd(ADTC),
    PARAMCD = "OVR",
    PARAM = "Overall Response",
    ANLØ1FL = "Y"
  ) %>%
  select(-ADTC)

derive_param_extreme_event(
  adrs,
  dataset_adsl = adsl,
  dataset_source = adrs,
  filter_source = PARAMCD == "OVR" & AVALC == "PD",
  order = exprs(ADT),
  new_var = AVALC,
  true_value = "Y",
  false_value = "N",
  mode = "first",
  set_values_to = exprs(
    PARAMCD = "PD",
    PARAM = "Disease Progression",
    ANLØ1FL = "Y",
    ADT = ADT
  )
)

# derive parameter indicating death
derive_param_extreme_event(
  dataset_adsl = adsl,
  dataset_source = adsl,
  filter_source = !is.na(DTHDT),
  new_var = AVALC,
  true_value = "Y",
  false_value = "N",
  mode = "first",
  set_values_to = exprs(
    PARAMCD = "DEATH",
    PARAM = "Death",
    ANLØ1FL = "Y",
    ADT = DTHDT
  )
)

```

 derive_param_first_event

Add a First Event Parameter

Description**[Deprecated]**

This function is *deprecated*, please use `derive_param_extreme_event()` instead with the `order` argument instead of the `date_var` argument.

Usage

```
derive_param_first_event(
  dataset,
  dataset_adsl,
  dataset_source,
  filter_source,
  date_var,
  subject_keys = exprs(STUDYID, USUBJID),
  set_values_to,
  check_type = "warning"
)
```

Arguments

<code>dataset</code>	Input dataset The PARAMCD variable is expected.
<code>dataset_adsl</code>	ADSL input dataset The variables specified for <code>subject_keys</code> are expected. For each observation of the specified dataset a new observation is added to the input dataset.
<code>dataset_source</code>	Source dataset All observations in the specified dataset fulfilling the condition specified by <code>filter_source</code> are considered as event. The variables specified by the <code>subject_keys</code> and <code>date_var</code> parameter are expected.
<code>filter_source</code>	Source filter All observations in <code>dataset_source</code> fulfilling the specified condition are considered as event. For subjects with at least one event AVALC is set to "Y", AVAL to 1, and ADT to the first date where the condition is fulfilled. For all other subjects AVALC is set to "N", AVAL to 0, and ADT to NA.
<code>date_var</code>	Date variable Date variable in the source dataset (<code>dataset_source</code>). The variable is used to sort the source dataset. ADT is set to the specified variable for events.
<code>subject_keys</code>	Variables to uniquely identify a subject A list of symbols created using <code>exprs()</code> is expected.
<code>set_values_to</code>	Variables to set A named list returned by <code>exprs()</code> defining the variables to be set for the new parameter, e.g. <code>exprs(PARAMCD = "PD", PARAM = "Disease Progression")</code> is expected. The values must be symbols, character strings, numeric values, or NA.

check_type Check uniqueness?
 If "warning" or "error" is specified, a message is issued if the observations of the input dataset restricted to the source parameter (source_param) are not unique with respect to the subject keys (subject_key parameter) and ADT.
Permitted Values: "none", "warning", "error"

Details

1. The input dataset is restricted to observations fulfilling filter_source.
2. For each subject (with respect to the variables specified for the subject_keys parameter) the first observation (with respect to date_var) where the event condition (filter_source parameter) is fulfilled is selected.
3. For each observation in dataset_ads1 a new observation is created. For subjects with event AVALC is set to "Y", AVAL to 1, and ADT to the first date where the event condition is fulfilled. For all other subjects AVALC is set to "N", AVAL to 0, and ADT to NA. For subjects with event all variables from dataset_source are kept. For subjects without event all variables which are in both dataset_ads1 and dataset_source are kept.
4. The variables specified by the set_values_to parameter are added to the new observations.
5. The new observations are added to input dataset.

Value

The input dataset with a new parameter indicating if and when an event occurred

See Also

Other deprecated: [derive_derived_param\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

derive_param_framingham

Adds a Parameter for Framingham Heart Study Cardiovascular Disease 10-Year Risk Score

Description

Adds a record for framingham score (FCVD101) for each by group (e.g., subject and visit) where the source parameters are available.

Usage

```

derive_param_framingham(
  dataset,
  by_vars,
  set_values_to = exprs(PARAMCD = "FCVD101"),
  sysbp_code = "SYSBP",
  chol_code = "CHOL",
  cholhdl_code = "CHOLHDL",
  age = AGE,
  sex = SEX,
  smokefl = SMOKEFL,
  diabetfl = DIABETFL,
  trthypfl = TRTHYPFL,
  get_unit_expr,
  filter = NULL
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>sysbp_code</code>, <code>chol_code</code> and <code>hdl_code</code>.</p>
by_vars	<p>Grouping variables</p> <p>For each group defined by <code>by_vars</code> an observation is added to the output dataset. Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p><i>Permitted Values:</i> list of variables</p>
set_values_to	<p>Variables to be set</p> <p>The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter.</p> <p><i>Permitted Values:</i> List of variable-value pairs</p>
sysbp_code	<p>Systolic blood pressure parameter code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the systolic blood pressure assessments.</p> <p><i>Permitted Values:</i> character value</p>
chol_code	<p>Total serum cholesterol code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the total cholesterol assessments. This must be measured in mg/dL.</p> <p><i>Permitted Values:</i> character value</p>
cholhdl_code	<p>HDL serum cholesterol code</p>

	The observations where PARAMCD equals the specified value are considered as the HDL cholesterol assessments. This must be measured in mg/dL. <i>Permitted Values:</i> character value
age	Subject age A variable containing the subject's age. <i>Permitted Values:</i> A numeric variable name that refers to a subject age column of the input dataset
sex	Subject sex A variable containing the subject's sex. <i>Permitted Values:</i> A character variable name that refers to a subject sex column of the input dataset
smokefl	Smoking status flag A flag indicating smoking status. <i>Permitted Values:</i> A character variable name that refers to a smoking status column of the input dataset.
diabetfl	Diabetic flag A flag indicating diabetic status. <i>Permitted Values:</i> A character variable name that refers to a diabetic status column of the input dataset
trthypfl	Treated with hypertension medication flag A flag indicating if a subject was treated with hypertension medication. <i>Permitted Values:</i> A character variable name that refers to a column that indicates whether a subject is treated for high blood pressure
get_unit_expr	An expression providing the unit of the parameter The result is used to check the units of the input parameters. <i>Permitted Values:</i> A variable of the input dataset or a function call
filter	Filter condition The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account. <i>Permitted Values:</i> a condition

Details

The values of age, sex, smokefl, diabetfl and trthypfl will be added to the by_vars list. The predicted probability of having cardiovascular disease (CVD) within 10-years according to Framingham formula. See AHA Journal article General Cardiovascular Risk Profile for Use in Primary Care for reference.

For Women:

Factor	Amount
Age	2.32888
Total Chol	1.20904
HDL Chol	-0.70833
Sys BP	2.76157

Sys BP + Hypertension Meds	2.82263
Smoker	0.52873
Non-Smoker	0
Diabetic	0.69154
Not Diabetic	0
Average Risk	26.1931
Risk Period	0.95012

For Men:

Factor	Amount
Age	3.06117
Total Chol	1.12370
HDL Chol	-0.93263
Sys BP	1.93303
Sys BP + Hypertension Meds	2.99881
Smoker	.65451
Non-Smoker	0
Diabetic	0.57367
Not Diabetic	0
Average Risk	23.9802
Risk Period	0.88936

The equation for calculating risk:

$$RiskFactors = (\log(Age)*AgeFactor) + (\log(TotalChol)*TotalCholFactor) + (\log(CholHDL)*CholHDLFactor)$$

$$Risk = 100 * (1 - RiskPeriodFactor^{RiskFactors})$$

Value

The input dataset with the new parameter added

See Also

[compute_framingham\(\)](#)

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_expected_records\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_rr\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```

library(tibble)

adcvrisk <- tribble(
  ~USUBJID, ~PARAMCD, ~PARAM, ~AVAL, ~AVALU,
  ~VISIT, ~AGE, ~SEX, ~SMOKEFL, ~DIABETFL, ~TRTHYPFL,
  "01-701-1015", "SYSBP", "Systolic Blood Pressure (mmHg)", 121,
  "mmHg", "BASELINE", 44, "F", "N", "N", "N",
  "01-701-1015", "SYSBP", "Systolic Blood Pressure (mmHg)", 115,
  "mmHg", "WEEK 2", 44, "F", "N", "N", "Y",
  "01-701-1015", "CHOL", "Total Cholesterol (mg/dL)", 216.16,
  "mg/dL", "BASELINE", 44, "F", "N", "N", "N",
  "01-701-1015", "CHOL", "Total Cholesterol (mg/dL)", 210.78,
  "mg/dL", "WEEK 2", 44, "F", "N", "N", "Y",
  "01-701-1015", "CHOLHDL", "Cholesterol/HDL-Cholesterol (mg/dL)", 54.91,
  "mg/dL", "BASELINE", 44, "F", "N", "N", "N",
  "01-701-1015", "CHOLHDL", "Cholesterol/HDL-Cholesterol (mg/dL)", 26.72,
  "mg/dL", "WEEK 2", 44, "F", "N", "N", "Y",
  "01-701-1028", "SYSBP", "Systolic Blood Pressure (mmHg)", 119,
  "mmHg", "BASELINE", 55, "M", "Y", "Y", "Y",
  "01-701-1028", "SYSBP", "Systolic Blood Pressure (mmHg)", 101,
  "mmHg", "WEEK 2", 55, "M", "Y", "Y", "Y",
  "01-701-1028", "CHOL", "Total Cholesterol (mg/dL)", 292.01,
  "mg/dL", "BASELINE", 55, "M", "Y", "Y", "Y",
  "01-701-1028", "CHOL", "Total Cholesterol (mg/dL)", 246.73,
  "mg/dL", "WEEK 2", 55, "M", "Y", "Y", "Y",
  "01-701-1028", "CHOLHDL", "Cholesterol/HDL-Cholesterol (mg/dL)", 65.55,
  "mg/dL", "BASELINE", 55, "M", "Y", "Y", "Y",
  "01-701-1028", "CHOLHDL", "Cholesterol/HDL-Cholesterol (mg/dL)", 44.62,
  "mg/dL", "WEEK 2", 55, "M", "Y", "Y", "Y"
)

adcvrisk %>%
  derive_param_framingham(
    by_vars = exprs(USUBJID, VISIT),
    set_values_to = exprs(
      PARAMCD = "FCVD101",
      PARAM = "FCVD1-Framingham CVD 10-Year Risk Score (%)"
    ),
    get_unit_expr = AVALU
  )

derive_param_framingham(
  adcvrisk,
  by_vars = exprs(USUBJID, VISIT),
  set_values_to = exprs(
    PARAMCD = "FCVD101",
    PARAM = "FCVD1-Framingham CVD 10-Year Risk Score (%)"
  ),
  get_unit_expr = extract_unit(PARAM)
)

```

derive_param_map	<i>Adds a Parameter for Mean Arterial Pressure</i>
------------------	--

Description

Adds a record for mean arterial pressure (MAP) for each by group (e.g., subject and visit) where the source parameters are available.

Usage

```
derive_param_map(
  dataset,
  by_vars,
  set_values_to = exprs(PARAMCD = "MAP"),
  sysbp_code = "SYSBP",
  diabp_code = "DIABP",
  hr_code = NULL,
  get_unit_expr,
  filter = NULL
)
```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>sysbp_code</code>, <code>diabp_code</code> and <code>hr_code</code>.</p>
by_vars	<p>Grouping variables</p> <p>For each group defined by <code>by_vars</code> an observation is added to the output dataset. Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p><i>Permitted Values:</i> list of variables</p>
set_values_to	<p>Variables to be set</p> <p>The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter.</p> <p><i>Permitted Values:</i> List of variable-value pairs</p>
sysbp_code	<p>Systolic blood pressure parameter code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the systolic blood pressure assessments.</p> <p><i>Permitted Values:</i> character value</p>

diabp_code	Diastolic blood pressure parameter code The observations where PARAMCD equals the specified value are considered as the diastolic blood pressure assessments. Permitted Values: character value
hr_code	Heart rate parameter code The observations where PARAMCD equals the specified value are considered as the heart rate assessments. Permitted Values: character value
get_unit_expr	An expression providing the unit of the parameter The result is used to check the units of the input parameters. Permitted Values: A variable of the input dataset or a function call
filter	Filter condition The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account. <i>Permitted Values:</i> a condition

Details

The analysis value of the new parameter is derived as

$$\frac{2DIABP + SYSBP}{3}$$

if it is based on diastolic and systolic blood pressure and

$$DIABP + 0.01e^{4.14 - \frac{40.74}{HR}} (SYSBP - DIABP)$$

if it is based on diastolic, systolic blood pressure, and heart rate.

Value

The input dataset with the new parameter added. Note, a variable will only be populated in the new parameter rows if it is specified in `by_vars`.

See Also

BDS-Findings Functions for adding Parameters/Records: `default_qtc_paramcd()`, `derive_expected_records()`, `derive_extreme_event()`, `derive_extreme_records()`, `derive_locf_records()`, `derive_param_bmi()`, `derive_param_bsa()`, `derive_param_computed()`, `derive_param_doseint()`, `derive_param_exist_flag()`, `derive_param_exposure()`, `derive_param_extreme_event()`, `derive_param_framingham()`, `derive_param_qtc()`, `derive_param_rr()`, `derive_param_wbc_abs()`, `derive_summary_records()`

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)

advs <- tribble(
  ~USUBJID, ~PARAMCD, ~PARAM, ~AVAL, ~VISIT,
```

```

"01-701-1015", "PULSE", "Pulse (beats/min)", 59, "BASELINE",
"01-701-1015", "PULSE", "Pulse (beats/min)", 61, "WEEK 2",
"01-701-1015", "DIABP", "Diastolic Blood Pressure (mmHg)", 51, "BASELINE",
"01-701-1015", "DIABP", "Diastolic Blood Pressure (mmHg)", 50, "WEEK 2",
"01-701-1015", "SYSBP", "Systolic Blood Pressure (mmHg)", 121, "BASELINE",
"01-701-1015", "SYSBP", "Systolic Blood Pressure (mmHg)", 121, "WEEK 2",
"01-701-1028", "PULSE", "Pulse (beats/min)", 62, "BASELINE",
"01-701-1028", "PULSE", "Pulse (beats/min)", 77, "WEEK 2",
"01-701-1028", "DIABP", "Diastolic Blood Pressure (mmHg)", 79, "BASELINE",
"01-701-1028", "DIABP", "Diastolic Blood Pressure (mmHg)", 80, "WEEK 2",
"01-701-1028", "SYSBP", "Systolic Blood Pressure (mmHg)", 130, "BASELINE",
"01-701-1028", "SYSBP", "Systolic Blood Pressure (mmHg)", 132, "WEEK 2"
)

# Derive MAP based on diastolic and systolic blood pressure
advs %>%
  derive_param_map(
    by_vars = exprs(USUBJID, VISIT),
    set_values_to = exprs(
      PARAMCD = "MAP",
      PARAM = "Mean Arterial Pressure (mmHg)"
    ),
    get_unit_expr = extract_unit(PARAM)
  ) %>%
  filter(PARAMCD != "PULSE")

# Derive MAP based on diastolic and systolic blood pressure and heart rate
derive_param_map(
  advs,
  by_vars = exprs(USUBJID, VISIT),
  hr_code = "PULSE",
  set_values_to = exprs(
    PARAMCD = "MAP",
    PARAM = "Mean Arterial Pressure (mmHg)"
  ),
  get_unit_expr = extract_unit(PARAM)
)

```

```
derive_param_qtc
```

Adds a Parameter for Corrected QT (an ECG measurement)

Description

Adds a record for corrected QT using either Bazett's, Fridericia's or Sagie's formula for each by group (e.g., subject and visit) where the source parameters are available.

Usage

```
derive_param_qtc(
  dataset,
```

```

    by_vars,
    method,
    set_values_to = default_qtc_paramcd(method),
    qt_code = "QT",
    rr_code = "RR",
    get_unit_expr,
    filter = NULL
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> and the <code>unit_var</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>qt_code</code> and <code>rr_code</code>.</p>
by_vars	<p>Grouping variables</p> <p>Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p>Permitted Values: list of variables</p>
method	<p>Method used to QT correction</p> <p>Permitted Values: "Bazett", "Fridericia", "Sagie"</p>
set_values_to	<p>Variables to be set</p> <p>The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter.</p> <p><i>Permitted Values:</i> List of variable-value pairs</p>
qt_code	<p>QT parameter code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the QT interval assessments. It is expected that QT is measured in msec.</p> <p>Permitted Values: character value</p>
rr_code	<p>RR parameter code</p> <p>The observations where <code>PARAMCD</code> equals the specified value are considered as the RR interval assessments. It is expected that RR is measured in msec.</p> <p>Permitted Values: character value</p>
get_unit_expr	<p>An expression providing the unit of the parameter</p> <p>The result is used to check the units of the input parameters.</p> <p>Permitted Values: A variable of the input dataset or a function call</p>
filter	<p>Filter condition</p> <p>The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account.</p> <p><i>Permitted Values:</i> a condition</p>

Value

The input dataset with the new parameter added. Note, a variable will only be populated in the new parameter rows if it is specified in `by_vars`.

See Also

[compute_qtc\(\)](#)

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_expected_records\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_rr\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```
library(tibble)

adeg <- tribble(
  ~USUBJID, ~PARAMCD, ~PARAM, ~AVAL, ~AVALU, ~VISIT,
  "01-701-1015", "HR", "Heart Rate (beats/min)", 70.14, "beats/min", "BASELINE",
  "01-701-1015", "QT", "QT Duration (msec)", 370, "msec", "WEEK 2",
  "01-701-1015", "HR", "Heart Rate (beats/min)", 62.66, "beats/min", "WEEK 1",
  "01-701-1015", "RR", "RR Duration (msec)", 710, "msec", "WEEK 2",
  "01-701-1028", "HR", "Heart Rate (beats/min)", 85.45, "beats/min", "BASELINE",
  "01-701-1028", "QT", "QT Duration (msec)", 480, "msec", "WEEK 2",
  "01-701-1028", "QT", "QT Duration (msec)", 350, "msec", "WEEK 3",
  "01-701-1028", "HR", "Heart Rate (beats/min)", 56.54, "beats/min", "WEEK 3",
  "01-701-1028", "RR", "RR Duration (msec)", 842, "msec", "WEEK 2",
)

derive_param_qtc(
  adeg,
  by_vars = exprs(USUBJID, VISIT),
  method = "Bazett",
  set_values_to = exprs(
    PARAMCD = "QTcBR",
    PARAM = "QTcB - Bazett's Correction Formula Rederived (msec)",
    AVALU = "msec"
  ),
  get_unit_expr = AVALU
)

derive_param_qtc(
  adeg,
  by_vars = exprs(USUBJID, VISIT),
  method = "Fridericia",
  set_values_to = exprs(
    PARAMCD = "QTcFR",
    PARAM = "QTcF - Fridericia's Correction Formula Rederived (msec)",
    AVALU = "msec"
  ),
)
```

```

    get_unit_expr = extract_unit(PARAM)
  )

  derive_param_qtc(
    adeg,
    by_vars = exprs(USUBJID, VISIT),
    method = "Sagie",
    set_values_to = exprs(
      PARAMCD = "QTLCR",
      PARAM = "QTlc - Sagie's Correction Formula Rederived (msec)",
      AVALU = "msec"
    ),
    get_unit_expr = extract_unit(PARAM)
  )

```

derive_param_rr	<i>Adds a Parameter for Derived RR (an ECG measurement)</i>
-----------------	---

Description

Adds a record for derived RR based on heart rate for each by group (e.g., subject and visit) where the source parameters are available.

Usage

```

derive_param_rr(
  dataset,
  by_vars,
  set_values_to = exprs(PARAMCD = "RRR"),
  hr_code = "HR",
  get_unit_expr,
  filter = NULL
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> parameter, <code>PARAMCD</code>, and <code>AVAL</code> are expected.</p> <p>The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset after restricting it by the filter condition (<code>filter</code> parameter) and to the parameters specified by <code>hr_code</code>.</p>
by_vars	<p>Grouping variables</p> <p>For each group defined by <code>by_vars</code> an observation is added to the output dataset. Only variables specified in <code>by_vars</code> will be populated in the newly created records.</p> <p><i>Permitted Values:</i> list of variables</p>

set_values_to	Variables to be set The specified variables are set to the specified values for the new observations. For example <code>exprs(PARAMCD = "MAP")</code> defines the parameter code for the new parameter. <i>Permitted Values:</i> List of variable-value pairs
hr_code	HR parameter code The observations where PARAMCD equals the specified value are considered as the heart rate assessments. <i>Permitted Values:</i> character value
get_unit_expr	An expression providing the unit of the parameter The result is used to check the units of the input parameters. <i>Permitted Values:</i> A variable of the input dataset or a function call
filter	Filter condition The specified condition is applied to the input dataset before deriving the new parameter, i.e., only observations fulfilling the condition are taken into account. <i>Permitted Values:</i> a condition

Details

The analysis value of the new parameter is derived as

$$\frac{60000}{HR}$$

Value

The input dataset with the new parameter added. Note, a variable will only be populated in the new parameter rows if it is specified in `by_vars`.

See Also

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_expected_records\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_wbc_abs\(\)](#), [derive_summary_records\(\)](#)

Examples

```
library(tibble)

adeg <- tribble(
  ~USUBJID, ~PARAMCD, ~PARAM, ~AVAL, ~AVALU, ~VISIT,
  "01-701-1015", "HR", "Heart Rate", 70.14, "beats/min", "BASELINE",
  "01-701-1015", "QT", "QT Duration", 370, "msec", "WEEK 2",
  "01-701-1015", "HR", "Heart Rate", 62.66, "beats/min", "WEEK 1",
  "01-701-1015", "RR", "RR Duration", 710, "msec", "WEEK 2",
  "01-701-1028", "HR", "Heart Rate", 85.45, "beats/min", "BASELINE",
  "01-701-1028", "QT", "QT Duration", 480, "msec", "WEEK 2",
```



```

    "01-701-1028", "QT", "QT Duration", 350, "msec", "WEEK 3",
    "01-701-1028", "HR", "Heart Rate", 56.54, "beats/min", "WEEK 3",
    "01-701-1028", "RR", "RR Duration", 842, "msec", "WEEK 2"
  )

  derive_param_rr(
    adeg,
    by_vars = exprs(USUBJID, VISIT),
    set_values_to = exprs(
      PARAMCD = "RRR",
      PARAM = "RR Duration Rederived (msec)",
      AVALU = "msec"
    ),
    get_unit_expr = AVALU
  )

```

derive_param_tte *Derive a Time-to-Event Parameter*

Description

Add a time-to-event parameter to the input dataset.

Usage

```

derive_param_tte(
  dataset = NULL,
  dataset_adsl,
  source_datasets,
  by_vars = NULL,
  start_date = TRTSDT,
  event_conditions,
  censor_conditions,
  create_datetime = FALSE,
  set_values_to,
  subject_keys = get_admiral_option("subject_keys")
)

```

Arguments

dataset	Input dataset The PARAMCD variable is expected.
dataset_adsl	ADSL input dataset The variables specified for start_date, start_imputation_flag, and subject_keys are expected.
source_datasets	Source datasets A named list of datasets is expected. The dataset_name field of tte_source() refers to the dataset provided in the list.

by_vars	<p>By variables</p> <p>If the parameter is specified, separate time to event parameters are derived for each by group.</p> <p>The by variables must be in at least one of the source datasets. Each source dataset must contain either all by variables or none of the by variables.</p> <p>The by variables are not included in the output dataset.</p>
start_date	<p>Time to event origin date</p> <p>The variable STARTDT is set to the specified date. The value is taken from the ADSL dataset.</p> <p>If the event or censoring date is before the origin date, ADT is set to the origin date.</p> <p>If the specified variable is imputed, the corresponding date imputation flag must be specified for start_imputation_flag.</p>
event_conditions	<p>Sources and conditions defining events</p> <p>A list of event_source() objects is expected.</p>
sensor_conditions	<p>Sources and conditions defining censorings</p> <p>A list of sensor_source() objects is expected.</p>
create_datetime	<p>Create datetime variables?</p> <p>If set to TRUE, variables ADTM and STARTDTM are created. Otherwise, variables ADT and STARTDT are created.</p>
set_values_to	<p>Variables to set</p> <p>A named list returned by exprs() defining the variables to be set for the new parameter, e.g. exprs(PARAMCD = "OS", PARAM = "Overall Survival") is expected. The values must be symbols, character strings, numeric values, expressions, or NA.</p>
subject_keys	<p>Variables to uniquely identify a subject</p> <p>A list of symbols created using exprs() is expected.</p>

Details

The following steps are performed to create the observations of the new parameter:

Deriving the events:

1. For each event source dataset the observations as specified by the filter element are selected. Then for each patient the first observation (with respect to date) is selected.
2. The ADT variable is set to the variable specified by the date element. If the date variable is a datetime variable, only the datepart is copied.
3. The CNSR variable is added and set to the censor element.
4. The variables specified by the set_values_to element are added.
5. The selected observations of all event source datasets are combined into a single dataset.

6. For each patient the first observation (with respect to the ADT variable) from the single dataset is selected.

Deriving the censoring observations:

1. For each censoring source dataset the observations as specified by the filter element are selected. Then for each patient the last observation (with respect to date) is selected.
2. The ADT variable is set to the variable specified by the date element. If the date variable is a datetime variable, only the datepart is copied.
3. The CNSR variable is added and set to the censor element.
4. The variables specified by the set_values_to element are added.
5. The selected observations of all censoring source datasets are combined into a single dataset.
6. For each patient the last observation (with respect to the ADT variable) from the single dataset is selected.

For each subject (as defined by the subject_keys parameter) an observation is selected. If an event is available, the event observation is selected. Otherwise the censoring observation is selected.

Finally:

1. The variables specified for start_date and start_imputation_flag are joined from the ADSL dataset. Only subjects in both datasets are kept, i.e., subjects with both an event or censoring and an observation in dataset_adsl.
2. The variables as defined by the set_values_to parameter are added.
3. The ADT/ADTM variable is set to the maximum of ADT/ADTM and STARTDT/STARTDTM (depending on the create_datetime parameter).
4. The new observations are added to the output dataset.

Value

The input dataset with the new parameter added

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)
data("admiral_adsl")

adsl <- admiral_adsl

# derive overall survival parameter
death <- event_source(
  dataset_name = "adsl",
  filter = DTHFL == "Y",
  date = DTHDT,
  set_values_to = exprs(
    EVNTDESC = "DEATH",
    SRCDOM = "ADSL",
```

```

    SRCVAR = "DTHDT"
  )
)

last_alive_dt <- censor_source(
  dataset_name = "adsl",
  date = LSTALVDT,
  set_values_to = exprs(
    EVNTDESC = "LAST DATE KNOWN ALIVE",
    SRCDOM = "ADSL",
    SRCVAR = "LSTALVDT"
  )
)

derive_param_tte(
  dataset_adsl = adsl,
  event_conditions = list(death),
  censor_conditions = list(last_alive_dt),
  source_datasets = list(adsl = adsl),
  set_values_to = exprs(
    PARAMCD = "OS",
    PARAM = "Overall Survival"
  )
) %>%
select(-STUDYID) %>%
filter(row_number() %in% 20:30)

# derive duration of response
# only observations for subjects in dataset_adsl are created
adsl <- tribble(
  ~USUBJID, ~DTHFL, ~DTHDT, ~RSPDT,
  "01", "Y", ymd("2021-06-12"), ymd("2021-03-04"),
  "02", "N", NA, NA,
  "03", "Y", ymd("2021-08-21"), NA,
  "04", "N", NA, ymd("2021-04-14")
) %>%
mutate(STUDYID = "AB42")

adrs <- tribble(
  ~USUBJID, ~AVALC, ~ADT, ~ASEQ,
  "01", "SD", ymd("2021-01-03"), 1,
  "01", "PR", ymd("2021-03-04"), 2,
  "01", "PD", ymd("2021-05-05"), 3,
  "02", "PD", ymd("2021-02-03"), 1,
  "04", "SD", ymd("2021-02-13"), 1,
  "04", "PR", ymd("2021-04-14"), 2,
  "04", "CR", ymd("2021-05-15"), 3
) %>%
mutate(STUDYID = "AB42", PARAMCD = "OVR")

pd <- event_source(
  dataset_name = "adrs",
  filter = AVALC == "PD",

```

```
    date = ADT,
    set_values_to = exprs(
      EVENTDESC = "PD",
      SRCDOM = "ADRS",
      SRCVAR = "ADTM",
      SRCSEQ = ASEQ
    )
  )
)

death <- event_source(
  dataset_name = "adsl",
  filter = DTHFL == "Y",
  date = DTHDT,
  set_values_to = exprs(
    EVENTDESC = "DEATH",
    SRCDOM = "ADSL",
    SRCVAR = "DTHDT"
  )
)

lastvisit <- censor_source(
  dataset_name = "adrs",
  date = ADT,
  censor = 1,
  set_values_to = exprs(
    EVENTDESC = "LAST TUMOR ASSESSMENT",
    SRCDOM = "ADRS",
    SRCVAR = "ADTM",
    SRCSEQ = ASEQ
  )
)

first_response <- censor_source(
  dataset_name = "adsl",
  date = RSPDT,
  censor = 1,
  set_values_to = exprs(
    EVENTDESC = "FIRST RESPONSE",
    SRCDOM = "ADSL",
    SRCVAR = "RSPDT"
  )
)

derive_param_tte(
  dataset_adsl = filter(adsl, !is.na(RSPDT)),
  start_date = RSPDT,
  event_conditions = list(pd, death),
  censor_conditions = list(lastvisit, first_response),
  source_datasets = list(adsl = adsl, adrs = adrs),
  set_values_to = exprs(
    PARAMCD = "DURRSP",
    PARAM = "Duration of Response"
  )
)
```

```

)

# derive time to adverse event for each preferred term
adsl <- tribble(
  ~USUBJID, ~TRTSDT,          ~EOSDT,
  "01",     ymd("2020-12-06"), ymd("2021-03-06"),
  "02",     ymd("2021-01-16"), ymd("2021-02-03")
) %>%
  mutate(STUDYID = "AB42")

ae <- tribble(
  ~USUBJID, ~AESTDTC,          ~AESEQ, ~AEDECOD,
  "01",     "2021-01-03T10:56", 1,      "Flu",
  "01",     "2021-03-04",      2,      "Cough",
  "01",     "2021",            3,      "Flu"
) %>%
  mutate(STUDYID = "AB42")

ae_ext <- derive_vars_dt(
  ae,
  dtc = AESTDTC,
  new_vars_prefix = "AEST",
  highest_imputation = "M",
  flag_imputation = "none"
)

ttae <- event_source(
  dataset_name = "ae",
  date = AESTDT,
  set_values_to = exprs(
    EVNTDESC = "AE",
    SRCDOM = "AE",
    SRCVAR = "AESTDTC",
    SRCSEQ = AESEQ
  )
)

eos <- censor_source(
  dataset_name = "adsl",
  date = EOSDT,
  set_values_to = exprs(
    EVNTDESC = "END OF STUDY",
    SRCDOM = "ADSL",
    SRCVAR = "EOSDT"
  )
)

derive_param_tte(
  dataset_adsl = adsl,
  by_vars = exprs(AEDECOD),
  start_date = TRTSDT,
  event_conditions = list(ttae),
  censor_conditions = list(eos),

```

```

source_datasets = list(adsl = adsl, ae = ae_ext),
set_values_to = exprs(
  PARAMCD = paste0("TTAE", as.numeric(as.factor(AEDECOD))),
  PARAM = paste("Time to First", AEDECOD, "Adverse Event"),
  PARCAT1 = "TTAE",
  PARCAT2 = AEDECOD
)
) %>%
select(USUBJID, STARTDT, PARAMCD, PARAM, ADT, CNSR, SRCSEQ)

```

derive_param_wbc_abs *Add a parameter for lab differentials converted to absolute values*

Description

Add a parameter by converting lab differentials from fraction or percentage to absolute values

Usage

```

derive_param_wbc_abs(
  dataset,
  by_vars,
  set_values_to,
  get_unit_expr,
  wbc_unit = "10^9/L",
  wbc_code = "WBC",
  diff_code,
  diff_type = "fraction"
)

```

Arguments

dataset	Input dataset The variables specified by the <code>by_vars</code> argument, <code>PARAMCD</code> , and <code>AVAL</code> are expected to be present. The variable specified by <code>by_vars</code> and <code>PARAMCD</code> must be a unique key of the input dataset, and to the parameters specified by <code>wbc_code</code> and <code>diff_code</code> .
by_vars	Grouping variables Permitted Values: list of variables
set_values_to	Variables to set A named list returned by <code>exprs()</code> defining the variables to be set for the new parameter, e.g. <code>exprs(PARAMCD = "LYMPH", PARAM = "Lymphocytes Abs (10^9/L)")</code> is expected.
get_unit_expr	An expression providing the unit of the parameter The result is used to check the units of the input parameters. Permitted Values: a variable containing unit from the input dataset, or a function call, for example, <code>get_unit_expr = extract_unit(PARAM)</code> .

wbc_unit	A string containing the required unit of the WBC parameter Default: "10^9/L"
wbc_code	White Blood Cell (WBC) parameter The observations where PARAMCD equals the specified value are considered as the WBC absolute results to use for converting the differentials. Default: "WBC" Permitted Values: character value
diff_code	white blood differential parameter The observations where PARAMCD equals the specified value are considered as the white blood differential lab results in fraction or percentage value to be converted into absolute value.
diff_type	A string specifying the type of differential Permitted Values: "percent", "fraction" Default: fraction

Details

If diff_type is "percent", the analysis value of the new parameter is derived as

$$\frac{WhiteBloodCellCount * PercentageValue}{100}$$

If diff_type is "fraction", the analysis value of the new parameter is derived as

$$WhiteBloodCellCount * FractionValue$$

New records are created for each group of records (grouped by by_vars) if 1) the white blood cell component in absolute value is not already available from the input dataset, and 2) the white blood cell absolute value (identified by wbc_code) and the white blood cell differential (identified by diff_code) are both present.

Value

The input dataset with the new parameter added

See Also

BDS-Findings Functions for adding Parameters/Records: [default_qtc_paramcd\(\)](#), [derive_expected_records\(\)](#), [derive_extreme_event\(\)](#), [derive_extreme_records\(\)](#), [derive_locf_records\(\)](#), [derive_param_bmi\(\)](#), [derive_param_bsa\(\)](#), [derive_param_computed\(\)](#), [derive_param_doseint\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_exposure\(\)](#), [derive_param_extreme_event\(\)](#), [derive_param_framingham\(\)](#), [derive_param_map\(\)](#), [derive_param_qtc\(\)](#), [derive_param_rr\(\)](#), [derive_summary_records\(\)](#)

Examples

```
library(tibble)

test_lb <- tribble(
  ~USUBJID, ~PARAMCD, ~AVAL, ~PARAM, ~VISIT,
  "P01", "WBC", 33, "Leukocyte Count (10^9/L)", "CYCLE 1 DAY 1",
```



```

"P01", "WBC", 38, "Leukocyte Count (10^9/L)", "CYCLE 2 DAY 1",
"P01", "LYMLE", 0.90, "Lymphocytes (fraction of 1)", "CYCLE 1 DAY 1",
"P01", "LYMLE", 0.70, "Lymphocytes (fraction of 1)", "CYCLE 2 DAY 1",
"P01", "ALB", 36, "Albumin (g/dL)", "CYCLE 2 DAY 1",
"P02", "WBC", 33, "Leukocyte Count (10^9/L)", "CYCLE 1 DAY 1",
"P02", "LYMPH", 29, "Lymphocytes Abs (10^9/L)", "CYCLE 1 DAY 1",
"P02", "LYMLE", 0.87, "Lymphocytes (fraction of 1)", "CYCLE 1 DAY 1",
"P03", "LYMLE", 0.89, "Lymphocytes (fraction of 1)", "CYCLE 1 DAY 1"
)

derive_param_wbc_abs(
  dataset = test_lb,
  by_vars = exprs(USUBJID, VISIT),
  set_values_to = exprs(
    PARAMCD = "LYMPH",
    PARAM = "Lymphocytes Abs (10^9/L)",
    DTYPE = "CALCULATION"
  ),
  get_unit_expr = extract_unit(PARAM),
  wbc_code = "WBC",
  diff_code = "LYMLE",
  diff_type = "fraction"
)

```

```
derive_summary_records
```

Add New Records Within By Groups Using Aggregation Functions

Description

It is not uncommon to have an analysis need whereby one needs to derive an analysis value (AVAL) from multiple records. The ADaM basic dataset structure variable DTYPE is available to indicate when a new derived records has been added to a dataset.

Usage

```

derive_summary_records(
  dataset,
  by_vars,
  filter = NULL,
  analysis_var,
  summary_fun,
  set_values_to = NULL
)

```

Arguments

dataset A data frame.

by_vars	Variables to consider for generation of groupwise summary records. Providing the names of variables in <code>exprs()</code> will create a groupwise summary and generate summary records for the specified groups.
filter	Filter condition as logical expression to apply during summary calculation. By default, filtering expressions are computed within <code>by_vars</code> as this will help when an aggregating, lagging, or ranking function is involved. For example, <ul style="list-style-type: none"> • <code>filter = (AVAL > mean(AVAL, na.rm = TRUE))</code> will filter all AVAL values greater than mean of AVAL with in <code>by_vars</code>. • <code>filter = (dplyr::n() > 2)</code> will filter n count of <code>by_vars</code> greater than 2.
analysis_var	Analysis variable.
summary_fun	Function that takes as an input the <code>analysis_var</code> and performs the calculation. This can include built-in functions as well as user defined functions, for example <code>mean</code> or <code>function(x) mean(x, na.rm = TRUE)</code> .
set_values_to	Variables to be set The specified variables are set to the specified values for the new observations. A list of variable name-value pairs is expected. <ul style="list-style-type: none"> • LHS refers to a variable. • RHS refers to the values to set to the variable. This can be a string, a symbol, a numeric value or NA, e.g., <code>exprs(PARAMCD = "TDOSE", PARCAT1 = "OVERALL")</code>. More general expression are not allowed.

Details

When all records have same values within `by_vars` then this function will retain those common values in the newly derived records. Otherwise new value will be set to NA.

Value

A data frame with derived records appended to original dataset.

See Also

`get_summary_records()`

BDS-Findings Functions for adding Parameters/Records: `default_qtc_paramcd()`, `derive_expected_records()`, `derive_extreme_event()`, `derive_extreme_records()`, `derive_locf_records()`, `derive_param_bmi()`, `derive_param_bsa()`, `derive_param_computed()`, `derive_param_doseint()`, `derive_param_exist_flag()`, `derive_param_exposure()`, `derive_param_extreme_event()`, `derive_param_framingham()`, `derive_param_map()`, `derive_param_qtc()`, `derive_param_rr()`, `derive_param_wbc_abs()`

Examples

```
library(tibble)
library(dplyr, warn.conflicts = TRUE)

adeg <- tribble(
  ~USUBJID, ~EGSEQ, ~PARAM, ~AVISIT, ~EGDTC, ~AVAL, ~TRTA,
```

```

"XYZ-1001", 1, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:50", 385, "",
"XYZ-1001", 2, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:52", 399, "",
"XYZ-1001", 3, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:56", 396, "",
"XYZ-1001", 4, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:45", 384, "Placebo",
"XYZ-1001", 5, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:48", 393, "Placebo",
"XYZ-1001", 6, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:51", 388, "Placebo",
"XYZ-1001", 7, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:45", 385, "Placebo",
"XYZ-1001", 8, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:48", 394, "Placebo",
"XYZ-1001", 9, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:51", 402, "Placebo",
"XYZ-1002", 1, "QTcF Int. (msec)", "Baseline", "2016-02-22T07:58", 399, "",
"XYZ-1002", 2, "QTcF Int. (msec)", "Baseline", "2016-02-22T07:58", 410, "",
"XYZ-1002", 3, "QTcF Int. (msec)", "Baseline", "2016-02-22T08:01", 392, "",
"XYZ-1002", 4, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:50", 401, "Active 20mg",
"XYZ-1002", 5, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:53", 407, "Active 20mg",
"XYZ-1002", 6, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:56", 400, "Active 20mg",
"XYZ-1002", 7, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:50", 412, "Active 20mg",
"XYZ-1002", 8, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:53", 414, "Active 20mg",
"XYZ-1002", 9, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:56", 402, "Active 20mg",
)

# Summarize the average of the triplicate ECG interval values (AVAL)
derive_summary_records(
  adeg,
  by_vars = exprs(USUBJID, PARAM, AVISIT),
  analysis_var = AVAL,
  summary_fun = function(x) mean(x, na.rm = TRUE),
  set_values_to = exprs(DTYPE = "AVERAGE")
)

advs <- tribble(
  ~USUBJID, ~VSSEQ, ~PARAM, ~AVAL, ~VSSTRESU, ~VISIT, ~VSDTC,
  "XYZ-001-001", 1164, "Weight", 99, "kg", "Screening", "2018-03-19",
  "XYZ-001-001", 1165, "Weight", 101, "kg", "Run-In", "2018-03-26",
  "XYZ-001-001", 1166, "Weight", 100, "kg", "Baseline", "2018-04-16",
  "XYZ-001-001", 1167, "Weight", 94, "kg", "Week 24", "2018-09-30",
  "XYZ-001-001", 1168, "Weight", 92, "kg", "Week 48", "2019-03-17",
  "XYZ-001-001", 1169, "Weight", 95, "kg", "Week 52", "2019-04-14",
)

# Set new values to any variable. Here, `DTYPE = MAXIMUM` refers to `max()` records
# and `DTYPE = AVERAGE` refers to `mean()` records.
derive_summary_records(
  advs,
  by_vars = exprs(USUBJID, PARAM),
  analysis_var = AVAL,
  summary_fun = max,
  set_values_to = exprs(DTYPE = "MAXIMUM")
) %>%
derive_summary_records(
  by_vars = exprs(USUBJID, PARAM),
  analysis_var = AVAL,
  summary_fun = mean,
  set_values_to = exprs(DTYPE = "AVERAGE")
)

```

```

)

# Sample ADEG dataset with triplicate record for only AVISIT = 'Baseline'
adeg <- tribble(
  ~USUBJID, ~EGSEQ, ~PARAM, ~AVISIT, ~EGDTC, ~AVAL, ~TRTA,
  "XYZ-1001", 1, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:50", 385, "",
  "XYZ-1001", 2, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:52", 399, "",
  "XYZ-1001", 3, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:56", 396, "",
  "XYZ-1001", 4, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:48", 393, "Placebo",
  "XYZ-1001", 5, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:51", 388, "Placebo",
  "XYZ-1001", 6, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:48", 394, "Placebo",
  "XYZ-1001", 7, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:51", 402, "Placebo",
  "XYZ-1002", 1, "QTcF Int. (msec)", "Baseline", "2016-02-22T07:58", 399, "",
  "XYZ-1002", 2, "QTcF Int. (msec)", "Baseline", "2016-02-22T07:58", 410, "",
  "XYZ-1002", 3, "QTcF Int. (msec)", "Baseline", "2016-02-22T08:01", 392, "",
  "XYZ-1002", 4, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:53", 407, "Active 20mg",
  "XYZ-1002", 5, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:56", 400, "Active 20mg",
  "XYZ-1002", 6, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:53", 414, "Active 20mg",
  "XYZ-1002", 7, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:56", 402, "Active 20mg",
)

# Compute the average of AVAL only if there are more than 2 records within the
# by group
derive_summary_records(
  adeg,
  by_vars = exprs(USUBJID, PARAM, AVISIT),
  filter = n() > 2,
  analysis_var = AVAL,
  summary_fun = function(x) mean(x, na.rm = TRUE),
  set_values_to = exprs(DTYPE = "AVERAGE")
)

```

derive_vars_aage

Derive Analysis Age

Description

Derives analysis age (AAGE) and analysis age unit (AAGEU)

Usage

```

derive_vars_aage(
  dataset,
  start_date = BRTHDT,
  end_date = RANDDT,
  unit = "years"
)

```

Arguments

dataset	Input dataset The columns specified by the start_date and the end_date parameter are expected.
start_date	The start date A date or date-time object is expected. Refer to derive_vars_dt() to impute and derive a date from a date character vector to a date object. Default: BRTHDT
end_date	The end date A date or date-time object is expected. Refer to derive_vars_dt() to impute and derive a date from a date character vector to a date object. Default: RANDDT
unit	Unit The age is derived in the specified unit Default: 'years' Permitted Values: 'years', 'months', 'weeks', 'days', 'hours', 'minutes', 'seconds'

Details

The age is derived as the integer part of the duration from start to end date in the specified unit. When 'years' or 'months' are specified in the out_unit parameter, because of the underlying lubridate::time_length() function that is used here, results are calculated based on the actual calendar length of months or years rather than assuming equal days every month (30.4375 days) or every year (365.25 days).

Value

The input dataset with AGE and AAGEU added

See Also

[derive_vars_duration\(\)](#)

ADSL Functions that returns variable appended to dataset: [derive_var_age_years\(\)](#), [derive_var_dthcaus\(\)](#), [derive_var_extreme_dtm\(\)](#), [derive_var_extreme_dt\(\)](#), [derive_vars_period\(\)](#)

Examples

```
library(tibble)
library(lubridate)

data <- tribble(
  ~BRTHDT, ~RANDDT,
  ymd("1984-09-06"), ymd("2020-02-24")
)
```

```
derive_vars_aage(data)
```

```
derive_vars_atc      Derive ATC Class Variables
```

Description

Add Anatomical Therapeutic Chemical class variables from FACM to ADCM

Usage

```
derive_vars_atc(
  dataset,
  dataset_facm,
  by_vars = exprs(USUBJID, CMREFID = FAREFID),
  value_var = FASTRESC
)
```

Arguments

dataset	Input dataset The variables specified by the by_vars parameter are required
dataset_facm	FACM dataset The variables specified by the by_vars and value_var parameters, FAGRPID and FATESTCD are required
by_vars	Keys used to merge dataset_facm with dataset <i>Permitted Values:</i> list of variables
value_var	The variable of dataset_facm containing the values of the transposed variables Default: FASTRESC

Value

The input dataset with ATC variables added

See Also

OCCDS Functions: [derive_var_trtemfl\(\)](#), [derive_vars_query\(\)](#), [get_terms_from_db\(\)](#)

Examples

```
library(tibble)

cm <- tribble(
  ~USUBJID, ~CMGRPID, ~CMREFID, ~CMDECOD,
  "BP40257-1001", "14", "1192056", "PARACETAMOL",
  "BP40257-1001", "18", "2007001", "SOLUMEDROL",
```

```

    "BP40257-1002", "19", "2791596", "SPIRONOLACTONE"
  )
  facm <- tribble(
    ~USUBJID, ~FAGRPID, ~FAREFID, ~FATESTCD, ~FASTRESC,
    "BP40257-1001", "1", "1192056", "CMATC1CD", "N",
    "BP40257-1001", "1", "1192056", "CMATC2CD", "N02",
    "BP40257-1001", "1", "1192056", "CMATC3CD", "N02B",
    "BP40257-1001", "1", "1192056", "CMATC4CD", "N02BE",
    "BP40257-1001", "1", "2007001", "CMATC1CD", "D",
    "BP40257-1001", "1", "2007001", "CMATC2CD", "D10",
    "BP40257-1001", "1", "2007001", "CMATC3CD", "D10A",
    "BP40257-1001", "1", "2007001", "CMATC4CD", "D10AA",
    "BP40257-1001", "2", "2007001", "CMATC1CD", "D",
    "BP40257-1001", "2", "2007001", "CMATC2CD", "D07",
    "BP40257-1001", "2", "2007001", "CMATC3CD", "D07A",
    "BP40257-1001", "2", "2007001", "CMATC4CD", "D07AA",
    "BP40257-1001", "3", "2007001", "CMATC1CD", "H",
    "BP40257-1001", "3", "2007001", "CMATC2CD", "H02",
    "BP40257-1001", "3", "2007001", "CMATC3CD", "H02A",
    "BP40257-1001", "3", "2007001", "CMATC4CD", "H02AB",
    "BP40257-1002", "1", "2791596", "CMATC1CD", "C",
    "BP40257-1002", "1", "2791596", "CMATC2CD", "C03",
    "BP40257-1002", "1", "2791596", "CMATC3CD", "C03D",
    "BP40257-1002", "1", "2791596", "CMATC4CD", "C03DA"
  )

  derive_vars_atc(cm, facm)

```

```
derive_vars_disposition_reason
```

Derive a Disposition Reason at a Specific Timepoint

Description

[Deprecated]

This function is *deprecated*. Please use `derive_vars_merged()` and specify the `filter_add` argument to derive the respective variables.

Derive a disposition reason from the the relevant records in the disposition domain.

Usage

```

derive_vars_disposition_reason(
  dataset,
  dataset_ds,
  new_var,
  reason_var,
  new_var_spe = NULL,
  reason_var_spe = NULL,

```

```

format_new_vars = format_reason_default,
filter_ds,
subject_keys = get_admiral_option("subject_keys")
)

```

Arguments

dataset	Input dataset
dataset_ds	Dataset containing the disposition information (e.g. ds) The dataset must contain: <ul style="list-style-type: none"> • STUDYID, USUBJID, • The variable(s) specified in the reason_var (and reason_var_spe, if required) • The variables used in filter_ds.
new_var	Name of the disposition reason variable A variable name is expected (e.g. DCSREAS).
reason_var	The variable used to derive the disposition reason A variable name is expected (e.g. DSDECOD).
new_var_spe	Name of the disposition reason detail variable A variable name is expected (e.g. DCSREASP). If new_var_spe is specified, it is expected that reason_var_spe is also specified, otherwise an error is issued. Default: NULL
reason_var_spe	The variable used to derive the disposition reason detail A variable name is expected (e.g. DSTERM). If new_var_spe is specified, it is expected that reason_var_spe is also specified, otherwise an error is issued. Default: NULL
format_new_vars	The function used to derive the reason(s) This function is used to derive the disposition reason(s) and must follow the below conventions <ul style="list-style-type: none"> • If only the main reason for discontinuation needs to be derived (i.e. new_var_spe is NULL), the function must have at least one character vector argument, e.g. <code>format_reason <- function(reason)</code> and new_var will be derived as <code>new_var = format_reason(reason_var)</code>. Typically, the content of the function would return reason_var or NA depending on the value (e.g. <code>if_else (reason != "COMPLETED" & !is.na(reason), reason, NA_character_)</code>). <code>DCSREAS = format_reason(DSDECOD)</code> returns <code>DCSREAS = DSDECOD</code> when <code>DSDECOD</code> is not 'COMPLETED' nor NA, NA otherwise. • If both the main reason and the details needs to be derived (new_var_spe is specified) the function must have two character vectors argument, e.g. <code>format_reason2 <- function(reason, reason_spe)</code> and new_var will be derived as <code>new_var = format_reason(reason_var)</code>, new_var_spe will be derived as <code>new_var_spe = format_reason(reason_var, reason_var_spe)</code>. Typically, the content of the function would return reason_var_spe or NA depending on the reason_var value (e.g. <code>if_else (reason == "OTHER",</code>

	reason_spe, NA_character_)). DCSREASP = format_reason(DSDECOD, DSTERM) returns DCSREASP = DSTERM when DSDECOD is equal to 'OTHER'.
	Default: format_reason_default, see format_reason_default() for details.
filter_ds	Filter condition for the disposition data. Filter used to select the relevant disposition data. It is expected that the filter restricts dataset_ds such that there is at most one observation per patient. An error is issued otherwise. Permitted Values: logical expression.
subject_keys	Variables to uniquely identify a subject A list of expressions where the expressions are symbols as returned by <code>exprs()</code> is expected.

Details

This functions returns the main reason for discontinuation (e.g. DCSREAS or DCTREAS). The reason for discontinuation is derived based on reason_var (e.g. DSDECOD) and format_new_vars. If new_var_spe is not NULL, then the function will also return the details associated with the reason for discontinuation (e.g. DCSREASP). The details associated with the reason for discontinuation are derived based on reason_var_spe (e.g. DSTERM), reason_var and format_new_vars.

Value

the input dataset with the disposition reason(s) (new_var and if required new_var_spe) added.

See Also

[format_reason_default\(\)](#)

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

derive_vars_dt

Derive/Impute a Date from a Date Character Vector

Description

Derive a date ('--DT') from a date character vector ('--DTC'). The date can be imputed (see `date_imputation` argument) and the date imputation flag ('--DTF') can be added.

Usage

```

derive_vars_dt(
  dataset,
  new_vars_prefix,
  dtc,
  highest_imputation = "n",
  date_imputation = "first",
  flag_imputation = "auto",
  min_dates = NULL,
  max_dates = NULL,
  preserve = FALSE
)

```

Arguments

dataset	Input dataset. The date character vector (dtc) must be present.
new_vars_prefix	Prefix used for the output variable(s). A character scalar is expected. For the date variable "DT" is appended to the specified prefix and for the date imputation flag "DTF". I.e., for new_vars_prefix = "AST" the variables ASTDT and ASTDTF are created.
dtc	The '--DTC' date to impute A character date is expected in a format like yyyy-mm-dd or yyyy-mm-ddThh:mm:ss. Trailing components can be omitted and - is a valid "missing" value for any component.
highest_imputation	Highest imputation level The highest_imputation argument controls which components of the DTC value are imputed if they are missing. All components up to the specified level are imputed. If a component at a higher level than the highest imputation level is missing, NA_character_ is returned. For example, for highest_imputation = "D" "2020" results in NA_character_ because the month is missing. If "n" is specified no imputation is performed, i.e., if any component is missing, NA_character_ is returned. If "Y" is specified, date_imputation should be "first" or "last" and min_dates or max_dates should be specified respectively. Otherwise, NA_character_ is returned if the year component is missing. <i>Default:</i> "n" <i>Permitted Values:</i> "Y" (year, highest level), "M" (month), "D" (day), "n" (none, lowest level)
date_imputation	The value to impute the day/month when a datepart is missing. A character value is expected, either as a

- format with month and day specified as "mm-dd": e.g. "06-15" for the 15th of June (The year can not be specified; for imputing the year "first" or "last" together with min_dates or max_dates argument can be used (see examples).),
- or as a keyword: "first", "mid", "last" to impute to the first/mid/last day/month.

The argument is ignored if highest_imputation is less then "D".

Default: "first"

flag_imputation

Whether the date imputation flag must also be derived.

If "auto" is specified, the date imputation flag is derived if the date_imputation argument is not null.

Default: "auto"

Permitted Values: "auto", "date" or "none"

min_dates

Minimum dates

A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example

```
impute_dtc_dtm(
  "2020-11",
  min_dates = list(
    ymd_hms("2020-12-06T12:12:12"),
    ymd_hms("2020-11-11T11:11:11")
  ),
  highest_imputation = "M"
)
```

returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).

max_dates

Maximum dates

A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.

preserve

Preserve day if month is missing and day is present

For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "MID").

Permitted Values: TRUE, FALSE

Default: FALSE

Details

In admiral we don't allow users to pick any single part of the date/time to impute, we only enable to impute up to a highest level, i.e. you couldn't choose to say impute months, but not days.

The presence of a '--DTF' variable is checked and if it already exists in the input dataset, a warning is issued and '--DTF' will be overwritten.

Value

The input dataset with the date '--DT' (and the date imputation flag '--DTF' if requested) added.

See Also

Date/Time Derivation Functions that returns variable appended to dataset: [derive_var_trtdurd\(\)](#), [derive_vars_dtm_to_dt\(\)](#), [derive_vars_dtm_to_tm\(\)](#), [derive_vars_dtm\(\)](#), [derive_vars_duration\(\)](#), [derive_vars_dy\(\)](#)

Examples

```
library(tibble)
library(lubridate)

mhdt <- tribble(
  ~MHSTDTC,
  "2019-07-18T15:25:40",
  "2019-07-18T15:25",
  "2019-07-18",
  "2019-02",
  "2019",
  "2019---07",
  ""
)

# Create ASTDT and ASTDTF
# No imputation for partial date
derive_vars_dt(
  mhdt,
  new_vars_prefix = "AST",
  dtc = MHSTDTC
)

# Create ASTDT and ASTDTF
# Impute partial dates to first day/month
derive_vars_dt(
  mhdt,
  new_vars_prefix = "AST",
  dtc = MHSTDTC,
  highest_imputation = "M"
)

# Impute partial dates to 6th of April
derive_vars_dt(
```

```

    mhdt,
    new_vars_prefix = "AST",
    dtc = MHSTDTC,
    highest_imputation = "M",
    date_imputation = "04-06"
  )

# Create AENDT and AENDTF
# Impute partial dates to last day/month
derive_vars_dt(
  mhdt,
  new_vars_prefix = "AEN",
  dtc = MHSTDTC,
  highest_imputation = "M",
  date_imputation = "last"
)

# Create BIRTHDT
# Impute partial dates to 15th of June. No DTF
derive_vars_dt(
  mhdt,
  new_vars_prefix = "BIRTH",
  dtc = MHSTDTC,
  highest_imputation = "M",
  date_imputation = "mid",
  flag_imputation = "none"
)

# Impute AE start date to the first date and ensure that the imputed date
# is not before the treatment start date
adae <- tribble(
  ~AESTDTC, ~TRTSDTM,
  "2020-12", ymd_hms("2020-12-06T12:12:12"),
  "2020-11", ymd_hms("2020-12-06T12:12:12")
)

derive_vars_dt(
  adae,
  dtc = AESTDTC,
  new_vars_prefix = "AST",
  highest_imputation = "M",
  min_dates = exprs(TRTSDTM)
)

# A user imputing dates as middle month/day, i.e. date_imputation = "mid" can
# use preserve argument to "preserve" partial dates. For example, "2019--07",
# will be displayed as "2019-06-07" rather than 2019-06-15 with preserve = TRUE

derive_vars_dt(
  mhdt,
  new_vars_prefix = "AST",
  dtc = MHSTDTC,
  highest_imputation = "M",

```

```

    date_imputation = "mid",
    preserve = TRUE
  )

```

 derive_vars_dtm

Derive/Impute a Datetime from a Date Character Vector

Description

Derive a datetime object ('--DTM') from a date character vector ('--DTC'). The date and time can be imputed (see `date_imputation/time_imputation` arguments) and the date/time imputation flag ('--DTF', '--TMF') can be added.

Usage

```

derive_vars_dtm(
  dataset,
  new_vars_prefix,
  dtc,
  highest_imputation = "h",
  date_imputation = "first",
  time_imputation = "first",
  flag_imputation = "auto",
  min_dates = NULL,
  max_dates = NULL,
  preserve = FALSE,
  ignore_seconds_flag = FALSE
)

```

Arguments

<code>dataset</code>	Input dataset The date character vector (<code>dtc</code>) must be present.
<code>new_vars_prefix</code>	Prefix used for the output variable(s). A character scalar is expected. For the date variable "DT" is appended to the specified prefix, for the date imputation flag "DTF", and for the time imputation flag "TMF". I.e., for <code>new_vars_prefix = "AST"</code> the variables ASTDT, ASTDTF, and ASTTMF are created.
<code>dtc</code>	The '--DTC' date to impute A character date is expected in a format like <code>yyyy-mm-dd</code> or <code>yyyy-mm-ddThh:mm:ss</code> . Trailing components can be omitted and <code>-</code> is a valid "missing" value for any component.
<code>highest_imputation</code>	Highest imputation level

The `highest_imputation` argument controls which components of the DTC value are imputed if they are missing. All components up to the specified level are imputed.

If a component at a higher level than the highest imputation level is missing, `NA_character_` is returned. For example, for `highest_imputation = "D"` `"2020"` results in `NA_character_` because the month is missing.

If `"n"` is specified, no imputation is performed, i.e., if any component is missing, `NA_character_` is returned.

If `"Y"` is specified, `date_imputation` should be `"first"` or `"last"` and `min_dates` or `max_dates` should be specified respectively. Otherwise, `NA_character_` is returned if the year component is missing.

Default: `"h"`

Permitted Values: `"Y"` (year, highest level), `"M"` (month), `"D"` (day), `"h"` (hour), `"m"` (minute), `"s"` (second), `"n"` (none, lowest level)

`date_imputation`

The value to impute the day/month when a datepart is missing.

A character value is expected, either as a

- format with month and day specified as `"mm-dd"`: e.g. `"06-15"` for the 15th of June (The year can not be specified; for imputing the year `"first"` or `"last"` together with `min_dates` or `max_dates` argument can be used (see examples).),
- or as a keyword: `"first"`, `"mid"`, `"last"` to impute to the first/mid/last day/month.

The argument is ignored if `highest_imputation` is less than `"D"`.

Default: `"first"`.

`time_imputation`

The value to impute the time when a timepart is missing.

A character value is expected, either as a

- format with hour, min and sec specified as `"hh:mm:ss"`: e.g. `"00:00:00"` for the start of the day,
- or as a keyword: `"first"`, `"last"` to impute to the start/end of a day.

The argument is ignored if `highest_imputation = "n"`.

Default: `"first"`.

`flag_imputation`

Whether the date/time imputation flag(s) must also be derived.

If `"auto"` is specified, the date imputation flag is derived if the `date_imputation` argument is not null and the time imputation flag is derived if the `time_imputation` argument is not null

Default: `"auto"`

Permitted Values: `"auto"`, `"date"`, `"time"`, `"both"`, or `"none"`

`min_dates`

Minimum dates

A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the

dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example

```
impute_dtc_dtm(
  "2020-11",
  min_dates = list(
    ymd_hms("2020-12-06T12:12:12"),
    ymd_hms("2020-11-11T11:11:11")
  ),
  highest_imputation = "M"
)
```

returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).

For date variables (not datetime) in the list the time is imputed to "00:00:00". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.

max_dates	<p>Maximum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.</p> <p>For date variables (not datetime) in the list the time is imputed to "23:59:59". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.</p>
preserve	<p>Preserve day if month is missing and day is present</p> <p>For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "mid").</p> <p>Permitted Values: TRUE, FALSE</p> <p><i>Default:</i> FALSE</p>
ignore_seconds_flag	<p>ADaM IG states that given SDTM ('--DTC') variable, if only hours and minutes are ever collected, and seconds are imputed in ('--DTM') as 00, then it is not necessary to set ('--TMF') to 'S'. A user can set this to TRUE so the 'S' Flag is dropped from ('--TMF').</p> <p>A logical value</p> <p><i>Default:</i> FALSE</p>

Details

In admiral we don't allow users to pick any single part of the date/time to impute, we only enable to impute up to a highest level, i.e. you couldn't choose to say impute months, but not days.

The presence of a '--DTF' variable is checked and the variable is not derived if it already exists in the input dataset. However, if '--TMF' already exists in the input dataset, a warning is issued and '--TMF' will be overwritten.

Value

The input dataset with the datetime '--DTM' (and the date/time imputation flag '--DTF', '--TMF') added.

See Also

Date/Time Derivation Functions that returns variable appended to dataset: [derive_var_trtdurd\(\)](#), [derive_vars_dtm_to_dt\(\)](#), [derive_vars_dtm_to_tm\(\)](#), [derive_vars_dt\(\)](#), [derive_vars_duration\(\)](#), [derive_vars_dy\(\)](#)

Examples

```
library(tibble)
library(lubridate)

mhdt <- tribble(
  ~MHSTDTC,
  "2019-07-18T15:25:40",
  "2019-07-18T15:25",
  "2019-07-18",
  "2019-02",
  "2019",
  "2019---07",
  ""
)

derive_vars_dtm(
  mhdt,
  new_vars_prefix = "AST",
  dtc = MHSTDTC,
  highest_imputation = "M"
)

# Impute AE end date to the last date and ensure that the imputed date is not
# after the death or data cut off date
adae <- tribble(
  ~AEENDTC, ~DTHDT, ~DCUTDT,
  "2020-12", ymd("2020-12-06"), ymd("2020-12-24"),
  "2020-11", ymd("2020-12-06"), ymd("2020-12-24")
)

derive_vars_dtm(
  adae,
  dtc = AEENDTC,
  new_vars_prefix = "AEN",
  highest_imputation = "M",
  date_imputation = "last",
  time_imputation = "last",
  max_dates = exprs(DTHDT, DCUTDT)
)

# Seconds has been removed from the input dataset. Function now uses
```

```

# ignore_seconds_flag to remove the 'S' from the --TMF variable.
mhdt <- tribble(
  ~MHSTDTC,
  "2019-07-18T15:25",
  "2019-07-18T15:25",
  "2019-07-18",
  "2019-02",
  "2019",
  "2019---07",
  ""
)

derive_vars_dtm(
  mhdt,
  new_vars_prefix = "AST",
  dtc = MHSTDTC,
  highest_imputation = "M",
  ignore_seconds_flag = TRUE
)

# A user imputing dates as middle month/day, i.e. date_imputation = "MID" can
# use preserve argument to "preserve" partial dates. For example, "2019--07",
# will be displayed as "2019-06-07" rather than 2019-06-15 with preserve = TRUE

derive_vars_dtm(
  mhdt,
  new_vars_prefix = "AST",
  dtc = MHSTDTC,
  highest_imputation = "M",
  date_imputation = "mid",
  preserve = TRUE
)

```

derive_vars_dtm_to_dt *Derive Date Variables from Datetime Variables*

Description

This function creates date(s) as output from datetime variable(s)

Usage

```
derive_vars_dtm_to_dt(dataset, source_vars)
```

Arguments

dataset	Input dataset
source_vars	A list of datetime variables created using <code>exprs()</code> from which dates are to be extracted

Value

A data frame containing the input dataset with the corresponding date (--DT) variable(s) of all datetime variables (--DTM) specified in source_vars.

See Also

Date/Time Derivation Functions that returns variable appended to dataset: [derive_var_trtdurd\(\)](#), [derive_vars_dtm_to_tm\(\)](#), [derive_vars_dtm\(\)](#), [derive_vars_dt\(\)](#), [derive_vars_duration\(\)](#), [derive_vars_dy\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

adcm <- tribble(
  ~USUBJID, ~TRTSDTM,          ~ASTDTM,          ~AENDTM,
  "PAT01",  "2012-02-25 23:00:00", "2012-02-28 19:00:00", "2012-02-25 23:00:00",
  "PAT01",  NA,                "2012-02-28 19:00:00", NA,
  "PAT01",  "2017-02-25 23:00:00", "2013-02-25 19:00:00", "2014-02-25 19:00:00",
  "PAT01",  "2017-02-25 16:00:00", "2017-02-25 14:00:00", "2017-03-25 23:00:00",
  "PAT01",  "2017-02-25 16:00:00", "2017-02-25 14:00:00", "2017-04-29 14:00:00",
) %>%
mutate(
  TRTSDTM = as_datetime(TRTSDTM),
  ASTDTM  = as_datetime(ASTDTM),
  AENDTM  = as_datetime(AENDTM)
)

adcm %>%
  derive_vars_dtm_to_dt(exprs(TRTSDTM, ASTDTM, AENDTM)) %>%
  select(USUBJID, starts_with("TRT"), starts_with("AST"), starts_with("AEN"))
```

derive_vars_dtm_to_tm *Derive Time Variables from Datetime Variables*

Description

This function creates time variable(s) as output from datetime variable(s)

Usage

```
derive_vars_dtm_to_tm(dataset, source_vars)
```

Arguments

dataset	Input dataset
source_vars	A list of datetime variables created using <code>exprs()</code> from which time is to be extracted

Details

The names of the newly added variables are automatically set by replacing the --DTM suffix of the source_vars with --TM. The --TM variables are created using the hms package.

Value

A data frame containing the input dataset with the corresponding time (--TM) variable(s) of all datetime variables (--DTM) specified in source_vars with the correct name.

See Also

Date/Time Derivation Functions that returns variable appended to dataset: [derive_var_trtdurd\(\)](#), [derive_vars_dtm_to_dt\(\)](#), [derive_vars_dtm\(\)](#), [derive_vars_dt\(\)](#), [derive_vars_duration\(\)](#), [derive_vars_dy\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

adcm <- tribble(
  ~USUBJID, ~TRTSDTM, ~ASTDTM, ~AENDTM,
  "PAT01", "2012-02-25 23:41:10", "2012-02-28 19:03:00", "2013-02-25 23:32:16",
  "PAT01", "", "2012-02-28 19:00:00", "",
  "PAT01", "2017-02-25 23:00:02", "2013-02-25 19:00:15", "2014-02-25 19:00:56",
  "PAT01", "2017-02-25 16:00:00", "2017-02-25 14:25:00", "2017-03-25 23:00:00",
  "PAT01", "2017-02-25 16:05:17", "2017-02-25 14:20:00", "2018-04-29 14:06:45",
) %>%
  mutate(
    TRTSDTM = as_datetime(TRTSDTM),
    ASTDTM = as_datetime(ASTDTM),
    AENDTM = as_datetime(AENDTM)
  )

adcm %>%
  derive_vars_dtm_to_tm(exprs(TRTSDTM)) %>%
  select(USUBJID, starts_with("TRT"), everything())

adcm %>%
  derive_vars_dtm_to_tm(exprs(TRTSDTM, ASTDTM, AENDTM)) %>%
  select(USUBJID, starts_with("TRT"), starts_with("AS"), starts_with("AE"))
```

derive_vars_duration *Derive Duration*

Description

Derives duration between two dates, specified by the variables present in input dataset e.g., duration of adverse events, relative day, age, ...

Usage

```

derive_vars_duration(
  dataset,
  new_var,
  new_var_unit = NULL,
  start_date,
  end_date,
  in_unit = "days",
  out_unit = "days",
  floor_in = TRUE,
  add_one = TRUE,
  trunc_out = FALSE
)

```

Arguments

dataset	Input dataset The variables specified by the start_date and the end_date parameter are expected.
new_var	Name of variable to create
new_var_unit	Name of the unit variable If the parameter is not specified, no variable for the unit is created.
start_date	The start date A date or date-time variable is expected. This variable must be present in specified input dataset. Refer to derive_vars_dt() to impute and derive a date from a date character vector to a date object.
end_date	The end date A date or date-time variable is expected. This variable must be present in specified input dataset. Refer to derive_vars_dt() to impute and derive a date from a date character vector to a date object.
in_unit	Input unit See floor_in and add_one parameter for details. Default: 'days' Permitted Values: 'years', 'months', 'days', 'hours', 'minutes', 'min', 'seconds', 'sec'
out_unit	Output unit The duration is derived in the specified unit Default: 'days' Permitted Values: 'years', 'months', 'days', 'hours', 'minutes', 'min', 'seconds', 'sec'
floor_in	Round down input dates? The input dates are round down with respect to the input unit, e.g., if the input unit is 'days', the time of the input dates is ignored.

	Default: ‘TRUE‘
	Permitted Values: TRUE, FALSE
add_one	Add one input unit? If the duration is non-negative, one input unit is added. I.e., the duration can not be zero. Default: TRUE Permitted Values: TRUE, FALSE
trunc_out	Return integer part The fractional part of the duration (in output unit) is removed, i.e., the integer part is returned. Default: FALSE Permitted Values: TRUE, FALSE

Details

The duration is derived as time from start to end date in the specified output unit. If the end date is before the start date, the duration is negative. The start and end date variable must be present in the specified input dataset.

Value

The input dataset with the duration and unit variable added

See Also

[compute_duration\(\)](#)

Date/Time Derivation Functions that returns variable appended to dataset: [derive_var_trtdurd\(\)](#), [derive_vars_dtm_to_dt\(\)](#), [derive_vars_dtm_to_tm\(\)](#), [derive_vars_dtm\(\)](#), [derive_vars_dt\(\)](#), [derive_vars_dy\(\)](#)

Examples

```
library(lubridate)
library(tibble)

# Derive age in years
data <- tribble(
  ~USUBJID, ~BRTHDT, ~RANDDT,
  "P01", ymd("1984-09-06"), ymd("2020-02-24"),
  "P02", ymd("1985-01-01"), NA,
  "P03", NA, ymd("2021-03-10"),
  "P04", NA, NA
)

derive_vars_duration(data,
  new_var = AAGE,
  new_var_unit = AAGEU,
  start_date = BRTHDT,
  end_date = RANDDT,
  out_unit = "years",
```

```

    add_one = FALSE,
    trunc_out = TRUE
  )

  # Derive adverse event duration in days
  data <- tribble(
    ~USUBJID, ~ASTDT, ~AENDT,
    "P01", ymd("2021-03-05"), ymd("2021-03-02"),
    "P02", ymd("2019-09-18"), ymd("2019-09-18"),
    "P03", ymd("1985-01-01"), NA,
    "P04", NA, NA
  )

  derive_vars_duration(data,
    new_var = ADURN,
    new_var_unit = ADURU,
    start_date = ASTDT,
    end_date = AENDT,
    out_unit = "days"
  )

  # Derive adverse event duration in minutes
  data <- tribble(
    ~USUBJID, ~ADTM, ~TRTSDTM,
    "P01", ymd_hms("2019-08-09T04:30:56"), ymd_hms("2019-08-09T05:00:00"),
    "P02", ymd_hms("2019-11-11T10:30:00"), ymd_hms("2019-11-11T11:30:00"),
    "P03", ymd_hms("2019-11-11T00:00:00"), ymd_hms("2019-11-11T04:00:00"),
    "P04", NA, ymd_hms("2019-11-11T12:34:56"),
  )

  derive_vars_duration(data,
    new_var = ADURN,
    new_var_unit = ADURU,
    start_date = ADTM,
    end_date = TRTSDTM,
    in_unit = "minutes",
    out_unit = "minutes",
    add_one = FALSE
  )

  # Derive adverse event start time since last dose in hours
  data <- tribble(
    ~USUBJID, ~ASTDTM, ~LDOSEDTM,
    "P01", ymd_hms("2019-08-09T04:30:56"), ymd_hms("2019-08-08T10:05:00"),
    "P02", ymd_hms("2019-11-11T23:59:59"), ymd_hms("2019-10-11T11:37:00"),
    "P03", ymd_hms("2019-11-11T00:00:00"), ymd_hms("2019-11-10T23:59:59"),
    "P04", ymd_hms("2019-11-11T12:34:56"), NA,
    "P05", NA, ymd_hms("2019-09-28T12:34:56")
  )

  derive_vars_duration(
    data,
    new_var = LDRELTM,
    new_var_unit = LDRELTMU,

```

```

start_date = LDOSEDTM,
end_date = ASTDTM,
in_unit = "hours",
out_unit = "hours",
add_one = FALSE
)

```

derive_vars_dy

Derive Relative Day Variables

Description

Adds relative day variables (--DY) to the dataset, e.g., ASTDY and AENDY.

Usage

```
derive_vars_dy(dataset, reference_date, source_vars)
```

Arguments

dataset	Input dataset The columns specified by the reference_date and the source_vars parameter are expected.
reference_date	The start date column, e.g., date of first treatment A date or date-time object column is expected. Refer to derive_vars_dt() to impute and derive a date from a date character vector to a date object.
source_vars	A list of datetime or date variables created using exprs() from which dates are to be extracted. This can either be a list of date(time) variables or named --DY variables and corresponding -DT(M) variables e.g. exprs(TRTSDTM, ASTDTM, AENDT) or exprs(TRTSDT, ASTDTM, AENDT, DEATHDY = DTHDT). If the source variable does not end in -DT(M), a name for the resulting --DY variable must be provided.

Details

The relative day is derived as number of days from the reference date to the end date. If it is nonnegative, one is added. I.e., the relative day of the reference date is 1. Unless a name is explicitly specified, the name of the resulting relative day variable is generated from the source variable name by replacing DT (or DTM as appropriate) with DY.

Value

The input dataset with --DY corresponding to the --DTM or --DT source variable(s) added

See Also

Date/Time Derivation Functions that returns variable appended to dataset: [derive_var_trtdurd\(\)](#), [derive_vars_dtm_to_dt\(\)](#), [derive_vars_dtm_to_tm\(\)](#), [derive_vars_dtm\(\)](#), [derive_vars_dt\(\)](#), [derive_vars_duration\(\)](#)

Examples

```
library(tibble)
library(lubridate)
library(dplyr, warn.conflicts = FALSE)

datain <- tribble(
  ~TRTSDTM, ~ASTDTM, ~AENDT,
  "2014-01-17T23:59:59", "2014-01-18T13:09:09", "2014-01-20"
) %>%
  mutate(
    TRTSDTM = as_datetime(TRTSDTM),
    ASTDTM = as_datetime(ASTDTM),
    AENDT = ymd(AENDT)
  )

derive_vars_dy(
  datain,
  reference_date = TRTSDTM,
  source_vars = exprs(TRTSDTM, ASTDTM, AENDT)
)

# specifying name of new variables
datain <- tribble(
  ~TRTSDT, ~DTHDT,
  "2014-01-17", "2014-02-01"
) %>%
  mutate(
    TRTSDT = ymd(TRTSDT),
    DTHDT = ymd(DTHDT)
  )

derive_vars_dy(
  datain,
  reference_date = TRTSDT,
  source_vars = exprs(TRTSDT, DEATHDY = DTHDT)
)
```

derive_vars_joined	<i>Add Variables from an Additional Dataset Based on Conditions from Both Datasets</i>
--------------------	--

Description

The function adds variables from an additional dataset to the input dataset. The selection of the observations from the additional dataset can depend on variables from both datasets. For example, add the lowest value (nadir) before the current observation.

Usage

```
derive_vars_joined(
  dataset,
  dataset_add,
  by_vars = NULL,
  order = NULL,
  new_vars = NULL,
  join_vars = NULL,
  filter_add = NULL,
  filter_join = NULL,
  mode = NULL,
  check_type = "warning"
)
```

Arguments

dataset	Input dataset The variables specified by <code>by_vars</code> are expected.
dataset_add	Additional dataset The variables specified by the <code>by_vars</code> , the <code>new_vars</code> , the <code>join_vars</code> , and the <code>order</code> argument are expected.
by_vars	Grouping variables The two datasets are joined by the specified variables. Variables from the additional dataset can be renamed by naming the element, i.e., <code>by_vars = exprs(<name in input dataset>)</code> <i>Permitted Values:</i> list of variables created by <code>exprs()</code>
order	Sort order If the argument is set to a non-null value, for each observation of the input dataset the first or last observation from the joined dataset is selected with respect to the specified order. The specified variables are expected in the additional dataset (<code>dataset_add</code>). If a variable is available in both <code>dataset</code> and <code>dataset_add</code> , the one from <code>dataset_add</code> is used for the sorting. <i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code> , e.g., <code>exprs(ADT, desc(AVAL))</code> or <code>NULL</code>
new_vars	Variables to add The specified variables from the additional dataset are added to the output dataset. Variables can be renamed by naming the element, i.e., <code>new_vars = exprs(<new name> = <old name>)</code> . For example <code>new_vars = exprs(var1, var2)</code> adds variables <code>var1</code> and <code>var2</code> from <code>dataset_add</code> to the input dataset. And <code>new_vars = exprs(var1, new_var2 = old_var2)</code> takes <code>var1</code> and <code>old_var2</code> from <code>dataset_add</code> and adds them to the input dataset renaming <code>old_var2</code> to <code>new_var2</code> .

	<p>If the argument is not specified or set to NULL, all variables from the additional dataset (<code>dataset_add</code>) are added.</p> <p><i>Permitted Values:</i> list of variables created by <code>exprs()</code></p>
<code>join_vars</code>	<p>Variables to use from additional dataset</p> <p>Any extra variables required from the additional dataset for <code>filter_join</code> should be specified for this argument. Variables specified for <code>new_vars</code> do not need to be repeated for <code>join_vars</code>. If a specified variable exists in both the input dataset and the additional dataset, the suffix ".join" is added to the variable from the additional dataset.</p> <p>The variables are not included in the output dataset.</p> <p><i>Permitted Values:</i> list of variables created by <code>exprs()</code></p>
<code>filter_add</code>	<p>Filter for additional dataset (<code>dataset_add</code>)</p> <p>Only observations from <code>dataset_add</code> fulfilling the specified condition are joined to the input dataset. If the argument is not specified, all observations are joined.</p> <p><i>Permitted Values:</i> a condition</p>
<code>filter_join</code>	<p>Filter for the joined dataset</p> <p>The specified condition is applied to the joined dataset. Therefore variables from both datasets <code>dataset</code> and <code>dataset_add</code> can be used.</p> <p><i>Permitted Values:</i> a condition</p>
<code>mode</code>	<p>Selection mode</p> <p>Determines if the first or last observation is selected. If the <code>order</code> argument is specified, <code>mode</code> must be non-null.</p> <p>If the <code>order</code> argument is not specified, the <code>mode</code> argument is ignored.</p> <p><i>Permitted Values:</i> "first", "last", NULL</p>
<code>check_type</code>	<p>Check uniqueness?</p> <p>If "warning" or "error" is specified, the specified message is issued if the observations of the (restricted) joined dataset are not unique with respect to the by variables and the order.</p> <p>This argument is ignored if <code>order</code> is not specified. In this case an error is issued independent of <code>check_type</code> if the restricted joined dataset contains more than one observation for any of the observations of the input dataset.</p> <p><i>Permitted Values:</i> "none", "warning", "error"</p>

Details

1. The records from the additional dataset (`dataset_add`) are restricted to those matching the `filter_add` condition.
2. The input dataset and the (restricted) additional dataset are left joined by the grouping variables (`by_vars`). If no grouping variables are specified, a full join is performed.
3. The joined dataset is restricted by the `filter_join` condition.
4. If `order` is specified, for each observation of the input dataset the first or last observation (depending on `mode`) is selected.

- The variables specified for `new_vars` are renamed (if requested) and merged to the input dataset. I.e., the output dataset contains all observations from the input dataset. For observations without a matching observation in the joined dataset the new variables are set to NA. Observations in the additional dataset which have no matching observation in the input dataset are ignored.

Value

The output dataset contains all observations and variables of the input dataset and additionally the variables specified for `new_vars` from the additional dataset (`dataset_add`).

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(tibble)
library(lubridate)
library(dplyr, warn.conflicts = FALSE)
library(tidyr)

# Add AVISIT (based on time windows), AWLO, and AWHI
adbds <- tribble(
  ~USUBJID, ~ADY,
  "1",      -33,
  "1",      -2,
  "1",       3,
  "1",     24,
  "2",      NA,
)

windows <- tribble(
  ~AVISIT, ~AWLO, ~AWHI,
  "BASELINE", -30, 1,
  "WEEK 1",    2,  7,
  "WEEK 2",    8, 15,
  "WEEK 3",   16, 22,
  "WEEK 4",   23, 30
)

derive_vars_joined(
  adbds,
  dataset_add = windows,
  filter_join = AWLO <= ADY & ADY <= AWHI
)
```

```

# derive the nadir after baseline and before the current observation
adbds <- tribble(
  ~USUBJID, ~ADY, ~AVAL,
  "1",      -7,   10,
  "1",       1,   12,
  "1",       8,   11,
  "1",      15,    9,
  "1",      20,   14,
  "1",      24,   12,
  "2",      13,    8
)

derive_vars_joined(
  adbds,
  dataset_add = adbds,
  by_vars = exprs(USUBJID),
  order = exprs(AVAL),
  new_vars = exprs(NADIR = AVAL),
  join_vars = exprs(ADY),
  filter_add = ADY > 0,
  filter_join = ADY.join < ADY,
  mode = "first",
  check_type = "none"
)

# add highest hemoglobin value within two weeks before AE,
# take earliest if more than one
adae <- tribble(
  ~USUBJID, ~ASTDY,
  "1",       3,
  "1",      22,
  "2",       2
)

adlb <- tribble(
  ~USUBJID, ~PARAMCD, ~ADY, ~AVAL,
  "1",     "HGB",     1,   8.5,
  "1",     "HGB",     3,   7.9,
  "1",     "HGB",     5,   8.9,
  "1",     "HGB",     8,   8.0,
  "1",     "HGB",     9,   8.0,
  "1",     "HGB",    16,   7.4,
  "1",     "HGB",    24,   8.1,
  "1",     "ALB",     1,   42,
)

derive_vars_joined(
  adae,
  dataset_add = adlb,
  by_vars = exprs(USUBJID),
  order = exprs(AVAL, desc(ADY)),
  new_vars = exprs(HGB_MAX = AVAL, HGB_DY = ADY),
  filter_add = PARAMCD == "HGB",

```

```

    filter_join = ASTDY - 14 <= ADY & ADY <= ASTDY,
    mode = "last"
  )

# Add APERIOD, APERIODC based on ADSL
adsl <- tribble(
  ~USUBJID, ~AP01SDT, ~AP01EDT, ~AP02SDT, ~AP02EDT,
  "1",      "2021-01-04", "2021-02-06", "2021-02-07", "2021-03-07",
  "2",      "2021-02-02", "2021-03-02", "2021-03-03", "2021-04-01"
) %>%
  mutate(across(ends_with("DT"), ymd)) %>%
  mutate(STUDYID = "xyz")

period_ref <- create_period_dataset(
  adsl,
  new_vars = exprs(APERSDT = APxxSDT, APEREDT = APxxEDT)
)

period_ref

adae <- tribble(
  ~USUBJID, ~ASTDT,
  "1",      "2021-01-01",
  "1",      "2021-01-05",
  "1",      "2021-02-05",
  "1",      "2021-03-05",
  "1",      "2021-04-05",
  "2",      "2021-02-15",
) %>%
  mutate(
    ASTDT = ymd(ASTDT),
    STUDYID = "xyz"
  )

derive_vars_joined(
  adae,
  dataset_add = period_ref,
  by_vars = exprs(STUDYID, USUBJID),
  join_vars = exprs(APERSDT, APEREDT),
  filter_join = APERSDT <= ASTDT & ASTDT <= APEREDT
)

```

derive_vars_last_dose *Derive Last Dose*

Description

Add EX source variables from last dose to the input dataset.

Usage

```

derive_vars_last_dose(
  dataset,
  dataset_ex,
  filter_ex = NULL,
  by_vars = exprs(STUDYID, USUBJID),
  dose_id = exprs(),
  dose_date,
  analysis_date,
  single_dose_condition = EXDOSFRQ == "ONCE",
  new_vars = NULL,
  traceability_vars = NULL
)

```

Arguments

dataset	Input dataset. The variables specified by the <code>by_vars</code> and <code>analysis_date</code> parameters are expected.
dataset_ex	Input EX dataset. The variables specified by the <code>by_vars</code> , <code>dose_date</code> , <code>new_vars</code> parameters, and source variables from <code>traceability_vars</code> parameter are expected.
filter_ex	Filtering condition applied to EX dataset. For example, it can be used to filter for valid dose. Defaults to <code>NULL</code> .
by_vars	Variables to join by (created by <code>rlang::exprs()</code>).
dose_id	Variables to identify unique dose (created by <code>rlang::exprs()</code>). Defaults to empty <code>exprs()</code> .
dose_date	The EX dose date variable. A date or date-time object is expected.
analysis_date	The analysis date variable. A date or date-time object is expected.
single_dose_condition	The condition for checking if <code>dataset_ex</code> is single dose. An error is issued if the condition is not true. Defaults to <code>(EXDOSFRQ == "ONCE")</code> .
new_vars	Variables to keep from <code>dataset_ex</code> , with the option to rename. Can either be variables created by <code>rlang::exprs()</code> (e.g. <code>exprs(VISIT)</code>), or named list returned by <code>exprs()</code> (e.g. <code>exprs(LSTEXVIS = VISIT)</code>). If set to <code>NULL</code> , then all variables from <code>dataset_ex</code> are kept without renaming. Defaults to <code>NULL</code> .
traceability_vars	A named list returned by <code>exprs()</code> listing the traceability variables, e.g. <code>exprs(LDOSEDOM = "EX", LDOSESEQ = EXSEQ)</code> . The left-hand side (names of the list elements) gives the names of the traceability variables in the returned dataset. The right-hand side (values of the list elements) gives the values of the traceability variables in the returned dataset. These can be either strings or symbols referring to existing variables.

Details

When doing date comparison to identify last dose, date-time imputations are done as follows:

- dose_date: time is imputed to 00:00:00 if the variable is a date variable
- analysis_date: time is imputed to 23:59:59 if the variable is a date variable

The last dose records are identified as follows:

1. The dataset_ex is filtered using filter_ex, if provided. This is useful for, for example, filtering for valid dose only.
2. The datasets dataset and dataset_ex are joined using by_vars.
3. The last dose is identified: the last dose is the EX record with maximum date where dose_date is lower to or equal to analysis_date, subject to both date values are non-NA. The last dose is identified per by_vars. If multiple EX records exist for the same dose_date, then either dose_id needs to be supplied (e.g. dose_id = exprs(EXSEQ)) to identify unique records, or an error is issued. When dose_id is supplied, the last EX record from the same dose_date sorted by dose_id will be used to identify last dose.
4. The EX source variables (as specified in new_vars) from last dose are appended to the dataset and returned to the user.

This function only works correctly for EX dataset with a structure of single dose per row. If your study EX dataset has multiple doses per row, use `create_single_dose_dataset()` to transform the EX dataset into single dose per row structure before calling `derive_vars_last_dose()`.

If variables (other than those specified in `by_vars`) exist in both `dataset` and `dataset_ex`, then join cannot be performed properly and an error is issued. To resolve the error, use `new_vars` to either keep variables unique to `dataset_ex`, or use this option to rename variables from `dataset_ex` (e.g. `new_vars = exprs(LSTEXVIS = VISIT)`).

Value

Input dataset with EX source variables from last dose added.

See Also

`derive_var_last_dose_amt()`, `derive_var_last_dose_date()`, `derive_var_last_dose_grp()`, `create_single_dose_dataset()`

General Derivation Functions for all ADaMs that returns variable appended to dataset: `derive_var_extreme_flag()`, `derive_var_joined_exist_flag()`, `derive_var_last_dose_amt()`, `derive_var_last_dose_date()`, `derive_var_last_dose_grp()`, `derive_var_merged_cat()`, `derive_var_merged_character()`, `derive_var_merged_exist_flag()`, `derive_var_merged_summary()`, `derive_var_obs_number()`, `derive_var_relative_flag()`, `derive_vars_joined()`, `derive_vars_merged_lookup()`, `derive_vars_merged()`, `derive_vars_transposed()`, `get_summary_records()`

Examples

```
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data(admiral_ae)
data(ex_single)

# create datetime variables in input datasets
ex_single <- derive_vars_dtm(
```



```

    head(ex_single, 100),
    dtc = EXENDTC,
    new_vars_prefix = "EXEN",
    flag_imputation = "none"
  )

adae <- admiral_ae %>%
  head(100) %>%
  derive_vars_dtm(
    dtc = AESTDTC,
    new_vars_prefix = "AST",
    highest_imputation = "M"
  )

# add last dose vars
adae %>%
  derive_vars_last_dose(
    dataset_ex = ex_single,
    filter_ex = (EXDOSE > 0 | (EXDOSE == 0 & grepl("PLACEBO", EXTRT))) &
      !is.na(EXENDTM),
    new_vars = exprs(EXDOSE, EXTRT, EXSEQ, EXENDTC, VISIT),
    dose_date = EXENDTM,
    analysis_date = ASTDTM
  ) %>%
  select(STUDYID, USUBJID, AESEQ, AESTDTC, EXDOSE, EXTRT, EXENDTC, EXSEQ, VISIT)

# or with traceability variables
adae %>%
  derive_vars_last_dose(
    dataset_ex = ex_single,
    filter_ex = (EXDOSE > 0 | (EXDOSE == 0 & grepl("PLACEBO", EXTRT))) &
      !is.na(EXENDTM),
    new_vars = exprs(EXDOSE, EXTRT, EXSEQ, EXENDTC, VISIT),
    dose_date = EXENDTM,
    analysis_date = ASTDTM,
    traceability_vars = exprs(LDOSEDOM = "EX", LDOSESEQ = EXSEQ, LDOSEVAR = "EXENDTC")
  ) %>%
  select(STUDYID, USUBJID, AESEQ, AESTDTC, EXDOSE, EXTRT, EXENDTC, LDOSEDOM, LDOSESEQ, LDOSEVAR)

```

derive_vars_merged *Add New Variable(s) to the Input Dataset Based on Variables from Another Dataset*

Description

Add new variable(s) to the input dataset based on variables from another dataset. The observations to merge can be selected by a condition (`filter_add` argument) and/or selecting the first or last observation for each by group (`order` and `mode` argument).

Usage

```

derive_vars_merged(
  dataset,
  dataset_add,
  by_vars,
  order = NULL,
  new_vars = NULL,
  mode = NULL,
  filter_add = NULL,
  match_flag = NULL,
  check_type = "warning",
  duplicate_msg = NULL
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> argument are expected.</p>
dataset_add	<p>Additional dataset</p> <p>The variables specified by the <code>by_vars</code>, the <code>new_vars</code>, and the <code>order</code> argument are expected.</p>
by_vars	<p>Grouping variables</p> <p>The input dataset and the selected observations from the additional dataset are merged by the specified by variables. The by variables must be a unique key of the selected observations. Variables from the additional dataset can be renamed by naming the element, i.e., <code>by_vars = exprs(<name in input dataset> = <name in additional dataset>)</code> similar to the dplyr joins.</p> <p><i>Permitted Values:</i> list of variables created by <code>exprs()</code></p>
order	<p>Sort order</p> <p>If the argument is set to a non-null value, for each by group the first or last observation from the additional dataset is selected with respect to the specified order.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code>, e.g., <code>exprs(ADT, desc(AVAL))</code> or NULL</p>
new_vars	<p>Variables to add</p> <p>The specified variables from the additional dataset are added to the output dataset. Variables can be renamed by naming the element, i.e., <code>new_vars = exprs(<new name> = <old name>)</code>. For example <code>new_vars = exprs(var1, var2)</code> adds variables <code>var1</code> and <code>var2</code> from <code>dataset_add</code> to the input dataset.</p> <p>And <code>new_vars = exprs(var1, new_var2 = old_var2)</code> takes <code>var1</code> and <code>old_var2</code> from <code>dataset_add</code> and adds them to the input dataset renaming <code>old_var2</code> to <code>new_var2</code>.</p> <p>If the argument is not specified or set to NULL, all variables from the additional dataset (<code>dataset_add</code>) are added.</p>

	<i>Default:</i> NULL
	<i>Permitted Values:</i> list of variables created by <code>exprs()</code>
mode	<p>Selection mode</p> <p>Determines if the first or last observation is selected. If the <code>order</code> argument is specified, <code>mode</code> must be non-null.</p> <p>If the <code>order</code> argument is not specified, the <code>mode</code> argument is ignored.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> "first", "last", NULL</p>
filter_add	<p>Filter for additional dataset (<code>dataset_add</code>)</p> <p>Only observations fulfilling the specified condition are taken into account for merging. If the argument is not specified, all observations are considered.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> a condition</p>
match_flag	<p>Match flag</p> <p>If the argument is specified (e.g., <code>match_flag = FLAG</code>), the specified variable (e.g., <code>FLAG</code>) is added to the input dataset. This variable will be <code>TRUE</code> for all selected records from <code>dataset_add</code> which are merged into the input dataset, and <code>NA</code> otherwise.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> Variable name</p>
check_type	<p>Check uniqueness?</p> <p>If "warning" or "error" is specified, the specified message is issued if the observations of the (restricted) additional dataset are not unique with respect to the by variables and the order.</p> <p><i>Default:</i> "warning"</p> <p><i>Permitted Values:</i> "none", "warning", "error"</p>
duplicate_msg	<p>Message of unique check</p> <p>If the uniqueness check fails, the specified message is displayed.</p> <p><i>Default:</i></p> <pre>paste("Dataset `dataset_add` contains duplicate records with respect to", enumerate(vars2chr(by_vars)))</pre>

Details

1. The records from the additional dataset (`dataset_add`) are restricted to those matching the `filter_add` condition.
2. If `order` is specified, for each by group the first or last observation (depending on `mode`) is selected.
3. The variables specified for `new_vars` are renamed (if requested) and merged to the input dataset using `left_join()`. I.e., the output dataset contains all observations from the input dataset. For observations without a matching observation in the additional dataset the new variables are set to `NA`. Observations in the additional dataset which have no matching observation in the input dataset are ignored.

Value

The output dataset contains all observations and variables of the input dataset and additionally the variables specified for `new_vars` from the additional dataset (`dataset_add`).

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(admiral.test)
library(dplyr, warn.conflicts = FALSE)
data("admiral_vs")
data("admiral_dm")

# Merging all dm variables to vs
derive_vars_merged(
  admiral_vs,
  dataset_add = select(admiral_dm, -DOMAIN),
  by_vars = exprs(STUDYID, USUBJID)
) %>%
  select(STUDYID, USUBJID, VSTESTCD, VISIT, VSTPT, VSSTRESN, AGE, AGEU)

# Merge last weight to adsl
data("admiral_adsl")
derive_vars_merged(
  admiral_adsl,
  dataset_add = admiral_vs,
  by_vars = exprs(STUDYID, USUBJID),
  order = exprs(VSDTC),
  mode = "last",
  new_vars = exprs(LASTWGT = VSSTRESN, LASTWGTU = VSSTRESU),
  filter_add = VSTESTCD == "WEIGHT",
  match_flag = vsdatafl
) %>%
  select(STUDYID, USUBJID, AGE, AGEU, LASTWGT, LASTWGTU, vsdatafl)

# Derive treatment start datetime (TRTSDTM)
data(admiral_ex)

## Impute exposure start date to first date/time
ex_ext <- derive_vars_dtm(
  admiral_ex,
  dtc = EXSTDTC,
  new_vars_prefix = "EXST",
  highest_imputation = "M",
```

```
)

## Add first exposure datetime and imputation flags to adsl
derive_vars_merged(
  select(admiral_dm, STUDYID, USUBJID),
  dataset_add = ex_ext,
  by_vars = exprs(STUDYID, USUBJID),
  new_vars = exprs(TRTSDTM = EXSTDTC, TRTSDTF = EXSTDTC, TRTSTMF = EXSTTMF),
  order = exprs(EXSTDTC),
  mode = "first"
)

# Derive treatment start datetime (TRTSDTM)
data(admiral_ex)

## Impute exposure start date to first date/time
ex_ext <- derive_vars_dtm(
  admiral_ex,
  dtc = EXSTDTC,
  new_vars_prefix = "EXST",
  highest_imputation = "M",
)

## Add first exposure datetime and imputation flags to adsl
derive_vars_merged(
  select(admiral_dm, STUDYID, USUBJID),
  dataset_add = ex_ext,
  filter_add = !is.na(EXSTDTC),
  by_vars = exprs(STUDYID, USUBJID),
  new_vars = exprs(TRTSDTM = EXSTDTC, TRTSDTF = EXSTDTC, TRTSTMF = EXSTTMF),
  order = exprs(EXSTDTC),
  mode = "first"
)

# Derive treatment end datetime (TRTEDTM)
## Impute exposure end datetime to last time, no date imputation
ex_ext <- derive_vars_dtm(
  admiral_ex,
  dtc = EXENDTC,
  new_vars_prefix = "EXEN",
  time_imputation = "last",
)

## Add last exposure datetime and imputation flag to adsl
derive_vars_merged(
  select(admiral_dm, STUDYID, USUBJID),
  dataset_add = ex_ext,
  filter_add = !is.na(EXENDTC),
  by_vars = exprs(STUDYID, USUBJID),
  new_vars = exprs(TRTEDTM = EXENDTC, TRTETMF = EXENTMF),
  order = exprs(EXENDTC),
  mode = "last"
)
```

derive_vars_merged_dt *Merge a (Imputed) Date Variable*

Description

[Deprecated]

This function is *deprecated*, please use `derive_vars_dt()` and `derive_vars_merged()` instead.

Merge a imputed date variable and date imputation flag from a dataset to the input dataset. The observations to merge can be selected by a condition and/or selecting the first or last observation for each by group.

Usage

```
derive_vars_merged_dt(
  dataset,
  dataset_add,
  by_vars,
  order = NULL,
  new_vars_prefix,
  filter_add = NULL,
  mode = NULL,
  dtc,
  date_imputation = NULL,
  flag_imputation = "auto",
  min_dates = NULL,
  max_dates = NULL,
  preserve = FALSE,
  check_type = "warning",
  duplicate_msg = NULL
)
```

Arguments

dataset	Input dataset The variables specified by the <code>by_vars</code> argument are expected.
dataset_add	Additional dataset The variables specified by the <code>by_vars</code> , the <code>dtc</code> , and the <code>order</code> argument are expected.
by_vars	Grouping variables The input dataset and the selected observations from the additional dataset are merged by the specified by variables. The by variables must be a unique key of the selected observations. Variables from the additional dataset can be renamed by naming the element, i.e., <code>by_vars = exprs(<name in input dataset> = <name in additional dataset>)</code> similar to the dplyr joins. <i>Permitted Values:</i> list of variables created by <code>exprs()</code>

order	<p>Sort order</p> <p>If the argument is set to a non-null value, for each by group the first or last observation from the additional dataset is selected with respect to the specified order. The imputed date variable can be specified as well (see examples below). Please note that NA is considered as the last value. I.e., if a order variable is NA and mode = "last", this observation is chosen while for mode = "first" the observation is chosen only if there are no observations where the variable is not 'NA'.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> list of variables or desc(<variable>) function calls created by exprs(), e.g., exprs(ADT, desc(AVAL)) or NULL</p>
new_vars_prefix	<p>Prefix used for the output variable(s).</p> <p>A character scalar is expected. For the date variable "DT" is appended to the specified prefix and for the date imputation flag "DTF". I.e., for new_vars_prefix = "AST" the variables ASTDT and ASTDTF are created.</p>
filter_add	<p>Filter for additional dataset (dataset_add)</p> <p>Only observations fulfilling the specified condition are taken into account for merging. If the argument is not specified, all observations are considered.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> a condition</p>
mode	<p>Selection mode</p> <p>Determines if the first or last observation is selected. If the order argument is specified, mode must be non-null.</p> <p>If the order argument is not specified, the mode argument is ignored.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> "first", "last", NULL</p>
dtc	<p>The '--DTC' date to impute</p> <p>A character date is expected in a format like yyyy-mm-dd or yyyy-mm-ddThh:mm:ss. Trailing components can be omitted and - is a valid "missing" value for any component.</p>
date_imputation	<p>The value to impute the day/month when a datepart is missing.</p> <p>A character value is expected, either as a</p> <ul style="list-style-type: none"> • format with month and day specified as "mm-dd": e.g. "06-15" for the 15th of June (The year can not be specified; for imputing the year "first" or "last" together with min_dates or max_dates argument can be used (see examples).), • or as a keyword: "first", "mid", "last" to impute to the first/mid/last day/month. <p>The argument is ignored if highest_imputation is less than "D".</p> <p><i>Default:</i> "first"</p>
flag_imputation	<p>Whether the date imputation flag must also be derived.</p>

	<p>If "auto" is specified, the date imputation flag is derived if the date_imputation argument is not null.</p> <p><i>Default:</i> "auto"</p> <p><i>Permitted Values:</i> "auto", "date" or "none"</p>
min_dates	<p>Minimum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example</p> <pre>impute_dtc_dtm("2020-11", min_dates = list(ymd_hms("2020-12-06T12:12:12"), ymd_hms("2020-11-11T11:11:11")), highest_imputation = "M")</pre> <p>returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).</p>
max_dates	<p>Maximum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.</p>
preserve	<p>Preserve day if month is missing and day is present</p> <p>For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "MID").</p> <p><i>Permitted Values:</i> TRUE, FALSE</p> <p><i>Default:</i> FALSE</p>
check_type	<p>Check uniqueness?</p> <p>If "warning" or "error" is specified, the specified message is issued if the observations of the (restricted) additional dataset are not unique with respect to the by variables and the order.</p> <p><i>Default:</i> "warning"</p> <p><i>Permitted Values:</i> "none", "warning", "error"</p>
duplicate_msg	<p>Message of unique check</p> <p>If the uniqueness check fails, the specified message is displayed.</p> <p><i>Default:</i></p> <pre>paste("Dataset `dataset_add` contains duplicate records with respect to", enumerate(vars2chr(by_vars)))</pre>

Details

1. The additional dataset is restricted to the observations matching the `filter_add` condition.
2. The date variable and if requested, the date imputation flag is added to the additional dataset.
3. If `order` is specified, for each by group the first or last observation (depending on mode) is selected.
4. The date and flag variables are merged to the input dataset.

Value

The output dataset contains all observations and variables of the input dataset and additionally the variable `<new_vars_prefix>DT` and optionally the variable `<new_vars_prefix>DTF` derived from the additional dataset (`dataset_add`).

See Also

Other deprecated: `derive_derived_param()`, `derive_param_first_event()`, `derive_var_agegr_fda()`, `derive_var_confirmation_flag()`, `derive_var_disposition_status()`, `derive_var_worst_flag()`, `derive_vars_disposition_reason()`, `derive_vars_merged_dtm()`, `filter_confirmation()`, `format_eoxxstt_default()`, `format_reason_default()`, `vars()`

derive_vars_merged_dtm

Merge a (Imputed) Datetime Variable

Description**[Deprecated]**

This function is *deprecated*, please use `derive_vars_dtm()` and `derive_vars_merged()` instead.

Merge a imputed datetime variable, date imputation flag, and time imputation flag from a dataset to the input dataset. The observations to merge can be selected by a condition and/or selecting the first or last observation for each by group.

Usage

```
derive_vars_merged_dtm(
  dataset,
  dataset_add,
  by_vars,
  order = NULL,
  new_vars_prefix,
  filter_add = NULL,
  mode = NULL,
  dtc,
  date_imputation = NULL,
  time_imputation = "00:00:00",
```

```

flag_imputation = "auto",
min_dates = NULL,
max_dates = NULL,
preserve = FALSE,
check_type = "warning",
duplicate_msg = NULL
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> argument are expected.</p>
dataset_add	<p>Additional dataset</p> <p>The variables specified by the <code>by_vars</code>, the <code>dtc</code>, and the <code>order</code> argument are expected.</p>
by_vars	<p>Grouping variables</p> <p>The input dataset and the selected observations from the additional dataset are merged by the specified by variables. The by variables must be a unique key of the selected observations. Variables from the additional dataset can be renamed by naming the element, i.e., <code>by_vars = exprs(<name in input dataset> = <name in additional dataset>)</code> similar to the <code>dplyr</code> joins.</p> <p><i>Permitted Values:</i> list of variables created by <code>exprs()</code></p>
order	<p>Sort order</p> <p>If the argument is set to a non-null value, for each by group the first or last observation from the additional dataset is selected with respect to the specified order. The imputed datetime variable can be specified as well (see examples below).</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code>, e.g., <code>exprs(ADT, desc(AVAL))</code> or NULL</p>
new_vars_prefix	<p>Prefix used for the output variable(s).</p> <p>A character scalar is expected. For the date variable "DT" is appended to the specified prefix, for the date imputation flag "DTF", and for the time imputation flag "TMF". I.e., for <code>new_vars_prefix = "AST"</code> the variables <code>ASTDT</code>, <code>ASTDTF</code>, and <code>ASTTMF</code> are created.</p>
filter_add	<p>Filter for additional dataset (<code>dataset_add</code>)</p> <p>Only observations fulfilling the specified condition are taken into account for merging. If the argument is not specified, all observations are considered.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> a condition</p>
mode	<p>Selection mode</p> <p>Determines if the first or last observation is selected. If the <code>order</code> argument is specified, <code>mode</code> must be non-null.</p> <p>If the <code>order</code> argument is not specified, the <code>mode</code> argument is ignored.</p>

	<i>Default:</i> NULL
	<i>Permitted Values:</i> "first", "last", NULL
dtc	<p>The '--DTC' date to impute</p> <p>A character date is expected in a format like yyyy-mm-dd or yyyy-mm-ddThh:mm:ss. Trailing components can be omitted and - is a valid "missing" value for any component.</p>
date_imputation	<p>The value to impute the day/month when a datepart is missing.</p> <p>A character value is expected, either as a</p> <ul style="list-style-type: none"> • format with month and day specified as "mm-dd": e.g. "06-15" for the 15th of June (The year can not be specified; for imputing the year "first" or "last" together with min_dates or max_dates argument can be used (see examples).), • or as a keyword: "first", "mid", "last" to impute to the first/mid/last day/month. <p>The argument is ignored if highest_imputation is less than "D".</p> <p><i>Default:</i> "first".</p>
time_imputation	<p>The value to impute the time when a timepart is missing.</p> <p>A character value is expected, either as a</p> <ul style="list-style-type: none"> • format with hour, min and sec specified as "hh:mm:ss": e.g. "00:00:00" for the start of the day, • or as a keyword: "first", "last" to impute to the start/end of a day. <p>The argument is ignored if highest_imputation = "n".</p> <p><i>Default:</i> "first".</p>
flag_imputation	<p>Whether the date/time imputation flag(s) must also be derived.</p> <p>If "auto" is specified, the date imputation flag is derived if the date_imputation argument is not null and the time imputation flag is derived if the time_imputation argument is not null</p> <p><i>Default:</i> "auto"</p> <p><i>Permitted Values:</i> "auto", "date", "time", "both", or "none"</p>
min_dates	<p>Minimum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example</p> <pre>impute_dtc_dtm("2020-11", min_dates = list(ymd_hms("2020-12-06T12:12:12"), ymd_hms("2020-11-11T11:11:11")))</pre>

```

    ),
    highest_imputation = "M"
  )

```

returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).

For date variables (not datetime) in the list the time is imputed to "00:00:00". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.

max_dates

Maximum dates

A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.

For date variables (not datetime) in the list the time is imputed to "23:59:59". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.

preserve

Preserve day if month is missing and day is present

For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "mid").

Permitted Values: TRUE, FALSE

Default: FALSE

check_type

Check uniqueness?

If "warning" or "error" is specified, the specified message is issued if the observations of the (restricted) additional dataset are not unique with respect to the by variables and the order.

Default: "warning"

Permitted Values: "none", "warning", "error"

duplicate_msg

Message of unique check

If the uniqueness check fails, the specified message is displayed.

Default:

```

paste("Dataset `dataset_add` contains duplicate records with respect to",
      enumerate(vars2chr(by_vars)))

```

Details

1. The additional dataset is restricted to the observations matching the filter_add condition.
2. The datetime variable and if requested, the date imputation flag and time imputation flag is added to the additional dataset.
3. If order is specified, for each by group the first or last observation (depending on mode) is selected.
4. The date and flag variables are merged to the input dataset.

Value

The output dataset contains all observations and variables of the input dataset and additionally the variable <new_vars_prefix>DT and optionally the variables <new_vars_prefix>DTF and <new_vars_prefix>TMF derived from the additional dataset (dataset_add).

See Also

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

derive_vars_merged_lookup

Merge Lookup Table with Source Dataset

Description

Merge user-defined lookup table with the input dataset. Optionally print a list of records from the input dataset that do not have corresponding mapping from the lookup table.

Usage

```
derive_vars_merged_lookup(
  dataset,
  dataset_add,
  by_vars,
  order = NULL,
  new_vars = NULL,
  mode = NULL,
  filter_add = NULL,
  check_type = "warning",
  duplicate_msg = NULL,
  print_not_mapped = TRUE
)
```

Arguments

dataset	Input dataset The variables specified by the by_vars argument are expected.
dataset_add	Lookup table The variables specified by the by_vars argument are expected.
by_vars	Grouping variables The input dataset and the selected observations from the additional dataset are merged by the specified by variables. The by variables must be a unique key of the selected observations. Variables from the additional dataset can be renamed

by naming the element, i.e., `by_vars = exprs(<name in input dataset> = <name in additional dataset>)` similar to the `dplyr` joins.

Permitted Values: list of variables created by `exprs()`

order	<p>Sort order</p> <p>If the argument is set to a non-null value, for each by group the first or last observation from the additional dataset is selected with respect to the specified order.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code>, e.g., <code>exprs(ADT, desc(AVAL))</code> or NULL</p>
new_vars	<p>Variables to add</p> <p>The specified variables from the additional dataset are added to the output dataset. Variables can be renamed by naming the element, i.e., <code>new_vars = exprs(<new name> = <old name>)</code>. For example <code>new_vars = exprs(var1, var2)</code> adds variables <code>var1</code> and <code>var2</code> from <code>dataset_add</code> to the input dataset.</p> <p>And <code>new_vars = exprs(var1, new_var2 = old_var2)</code> takes <code>var1</code> and <code>old_var2</code> from <code>dataset_add</code> and adds them to the input dataset renaming <code>old_var2</code> to <code>new_var2</code>.</p> <p>If the argument is not specified or set to NULL, all variables from the additional dataset (<code>dataset_add</code>) are added.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> list of variables created by <code>exprs()</code></p>
mode	<p>Selection mode</p> <p>Determines if the first or last observation is selected. If the <code>order</code> argument is specified, <code>mode</code> must be non-null.</p> <p>If the <code>order</code> argument is not specified, the <code>mode</code> argument is ignored.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> "first", "last", NULL</p>
filter_add	<p>Filter for additional dataset (<code>dataset_add</code>)</p> <p>Only observations fulfilling the specified condition are taken into account for merging. If the argument is not specified, all observations are considered.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> a condition</p>
check_type	<p>Check uniqueness?</p> <p>If "warning" or "error" is specified, the specified message is issued if the observations of the (restricted) additional dataset are not unique with respect to the by variables and the order.</p> <p><i>Default:</i> "warning"</p> <p><i>Permitted Values:</i> "none", "warning", "error"</p>
duplicate_msg	<p>Message of unique check</p> <p>If the uniqueness check fails, the specified message is displayed.</p> <p><i>Default:</i></p>

```
paste("Dataset `dataset_add` contains duplicate records with respect to",
      enumerate(vars2chr(by_vars)))
```

```
print_not_mapped
```

Print a list of unique `by_vars` values that do not have corresponding records from the lookup table?

Default: TRUE

Permitted Values: TRUE, FALSE

Value

The output dataset contains all observations and variables of the input dataset, and add the variables specified in `new_vars` from the lookup table specified in `dataset_add`. Optionally prints a list of unique `by_vars` values that do not have corresponding records from the lookup table (by specifying `print_not_mapped = TRUE`).

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(admiral.test)
library(tibble)
library(dplyr, warn.conflicts = FALSE)
data("admiral_vs")
param_lookup <- tribble(
  ~VSTESTCD, ~VSTEST, ~PARAMCD, ~PARAM,
  "SYSBP", "Systolic Blood Pressure", "SYSBP", "Systolic Blood Pressure (mmHg)",
  "WEIGHT", "Weight", "WEIGHT", "Weight (kg)",
  "HEIGHT", "Height", "HEIGHT", "Height (cm)",
  "TEMP", "Temperature", "TEMP", "Temperature (C)",
  "MAP", "Mean Arterial Pressure", "MAP", "Mean Arterial Pressure (mmHg)",
  "BMI", "Body Mass Index", "BMI", "Body Mass Index(kg/m^2)",
  "BSA", "Body Surface Area", "BSA", "Body Surface Area(m^2)"
)
derive_vars_merged_lookup(
  dataset = admiral_vs,
  dataset_add = param_lookup,
  by_vars = exprs(VSTESTCD),
  new_vars = exprs(PARAMCD),
  print_not_mapped = TRUE
)
```

derive_vars_period *Add Subperiod, Period, or Phase Variables to ADSL*

Description

The function adds subperiod, period, or phase variables like P01S1SDT, P01S2SDT, AP01SDTM, AP02SDTM, TRT01A, TRT02A, PH1SDT, PH2SDT, ... to the input dataset. The values of the variables are defined by a period reference dataset which has one observations per patient and subperiod, period, or phase.

Usage

```
derive_vars_period(
  dataset,
  dataset_ref,
  new_vars,
  subject_keys = get_admiral_option("subject_keys")
)
```

Arguments

dataset	ADSL dataset The variables specified by subject_keys are expected.
dataset_ref	Period reference dataset The variables specified by new_vars and subject_keys are expected. If subperiod variables are requested, APERIOD and ASPER are expected. If period variables are requested, APERIOD is expected. If phase variables are requested, APHASEN is expected.
new_vars	New variables A named list of variables like exprs(PHwSDT = PHSDT, PHwEDT = PHEDT, APHASEw = APHASE) is expected. The left hand side of the elements defines a set of variables (in CDISC notation) to be added to the output dataset. The right hand side defines the source variable from the period reference dataset. If the lower case letter "w" is used it refers to a phase variable, if the lower case letters "xx" are used it refers to a period variable, and if both "xx" and "w" are used it refers to a subperiod variable. Only one type must be used, e.g., all left hand side values must refer to period variables. It is not allowed to mix for example period and subperiod variables. If period <i>and</i> subperiod variables are required, separate calls must be used.
subject_keys	Variables to uniquely identify a subject A list of expressions where the expressions are symbols as returned by exprs() is expected.

Details

For each subperiod/period/phase in the period reference dataset and each element in `new_vars` a variable (LHS value of `new_vars`) is added to the output dataset and set to the value of the source variable (RHS value of `new_vars`).

Value

The input dataset with subperiod/period/phase variables added (see "Details" section)

See Also

[create_period_dataset\(\)](#)

ADSL Functions that returns variable appended to dataset: [derive_var_age_years\(\)](#), [derive_var_dthcaus\(\)](#), [derive_var_extreme_dtm\(\)](#), [derive_var_extreme_dt\(\)](#), [derive_vars_aage\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

adsl <- tibble(STUDYID = "xyz", USUBJID = c("1", "2"))

# Add period variables to ADSL
period_ref <- tribble(
  ~USUBJID, ~APERIOD, ~APERSDT, ~APEREDT,
  "1",      1, "2021-01-04", "2021-02-06",
  "1",      2, "2021-02-07", "2021-03-07",
  "2",      1, "2021-02-02", "2021-03-02",
  "2",      2, "2021-03-03", "2021-04-01"
) %>%
  mutate(
    STUDYID = "xyz",
    APERIOD = as.integer(APERIOD),
    across(matches("APER[ES]DT"), ymd)
  )

derive_vars_period(
  adsl,
  dataset_ref = period_ref,
  new_vars = exprs(APxxSDT = APERSDT, APxxEDT = APEREDT)
) %>%
  select(STUDYID, USUBJID, AP01SDT, AP01EDT, AP02SDT, AP02EDT)

# Add phase variables to ADSL
phase_ref <- tribble(
  ~USUBJID, ~APHASEN, ~PHSDT, ~PHEDT, ~APHASE,
  "1",      1, "2021-01-04", "2021-02-06", "TREATMENT",
  "1",      2, "2021-02-07", "2021-03-07", "FUP",
  "2",      1, "2021-02-02", "2021-03-02", "TREATMENT"
) %>%
```

```

mutate(
  STUDYID = "xyz",
  APHASEN = as.integer(APHASEN),
  across(matches("PH[ES]DT"), ymd)
)

derive_vars_period(
  adsl,
  dataset_ref = phase_ref,
  new_vars = exprs(PHwSDT = PHSdT, PHwEDT = PHEDT, APHASEw = APHASE)
) %>%
  select(STUDYID, USUBJID, PH1SDT, PH1EDT, PH2SDT, PH2EDT, APHASE1, APHASE2)

# Add subperiod variables to ADSL
subperiod_ref <- tribble(
  ~USUBJID, ~APERIOD, ~ASPER, ~ASPRSDT, ~ASPREDT,
  "1",      1,      1, "2021-01-04", "2021-01-19",
  "1",      1,      2, "2021-01-20", "2021-02-06",
  "1",      2,      1, "2021-02-07", "2021-03-07",
  "2",      1,      1, "2021-02-02", "2021-03-02",
  "2",      2,      1, "2021-03-03", "2021-04-01"
) %>%
  mutate(
    STUDYID = "xyz",
    APERIOD = as.integer(APERIOD),
    ASPER = as.integer(ASPER),
    across(matches("ASPR[ES]DT"), ymd)
  )

derive_vars_period(
  adsl,
  dataset_ref = subperiod_ref,
  new_vars = exprs(PxxSwSDT = ASPRSdT, PxxSwEDT = ASPREDT)
) %>%
  select(STUDYID, USUBJID, P01S1SDT, P01S1EDT, P01S2SDT, P01S2EDT, P02S1SDT, P02S1EDT)

```

derive_vars_query *Derive Query Variables*

Description

Derive Query Variables

Usage

```
derive_vars_query(dataset, dataset_queries)
```

Arguments

dataset Input dataset.

dataset_queries A dataset containing required columns VAR_PREFIX, QUERY_NAME, TERM_LEVEL, TERM_NAME, TERM_ID, and optional columns QUERY_ID, QUERY_SCOPE, QUERY_SCOPE_NUM. The content of the dataset will be verified by [assert_valid_queries\(\)](#). [create_query_data\(\)](#) can be used to create the dataset.

Details

This function can be used to derive CDISC variables such as SMQzzNAM, SMQzzCD, SMQzzSC, SMQzzSCN, and CQzzNAM in ADAE and ADMH, and variables such as SDGzzNAM, SDGzzCD, and SDGzzSC in ADCM. An example usage of this function can be found in the [OCCDS vignette](#).

A query dataset is expected as an input to this function. See the [Queries Dataset Documentation vignette](#) for descriptions, or call `data("queries")` for an example of a query dataset.

For each unique element in VAR_PREFIX, the corresponding "NAM" variable will be created. For each unique VAR_PREFIX, if QUERY_ID is not "" or NA, then the corresponding "CD" variable is created; similarly, if QUERY_SCOPE is not "" or NA, then the corresponding "SC" variable will be created; if QUERY_SCOPE_NUM is not "" or NA, then the corresponding "SCN" variable will be created.

For each record in dataset, the "NAM" variable takes the value of QUERY_NAME if the value of TERM_NAME or TERM_ID in dataset_queries matches the value of the respective TERM_LEVEL in dataset. Note that TERM_NAME in dataset_queries dataset may be NA only when TERM_ID is non-NA and vice versa. The "CD", "SC", and "SCN" variables are derived accordingly based on QUERY_ID, QUERY_SCOPE, and QUERY_SCOPE_NUM respectively, whenever not missing.

Value

The input dataset with query variables derived.

See Also

[create_query_data\(\)](#) [assert_valid_queries\(\)](#)

OCCDS Functions: [derive_var_trtemfl\(\)](#), [derive_vars_atc\(\)](#), [get_terms_from_db\(\)](#)

Examples

```
library(tibble)
data("queries")
adae <- tribble(
  ~USUBJID, ~ASTDTM, ~AETERM, ~AESEQ, ~AEDECOD, ~AELLT, ~AELLTCD,
  "01", "2020-06-02 23:59:59", "ALANINE AMINOTRANSFERASE ABNORMAL",
  3, "Alanine aminotransferase abnormal", NA_character_, NA_integer_,
  "02", "2020-06-05 23:59:59", "BASEDOW'S DISEASE",
  5, "Basedow's disease", NA_character_, 1L,
  "03", "2020-06-07 23:59:59", "SOME TERM",
  2, "Some query", "Some term", NA_integer_,
  "05", "2020-06-09 23:59:59", "ALVEOLAR PROTEINOSIS",
```

```

    7, "Alveolar proteinosis", NA_character_, NA_integer_
  )
  derive_vars_query(adae, queries)

```

derive_vars_transposed

Derive Variables by Transposing and Merging a Second Dataset

Description

Adds variables from a vertical dataset after transposing it into a wide one.

Usage

```

derive_vars_transposed(
  dataset,
  dataset_merge,
  by_vars,
  key_var,
  value_var,
  filter = NULL
)

```

Arguments

dataset	Input dataset The variables specified by the <code>by_vars</code> parameter are required
dataset_merge	Dataset to transpose and merge The variables specified by the <code>by_vars</code> , <code>key_var</code> and <code>value_var</code> parameters are expected
by_vars	Keys used to merge <code>dataset_merge</code> with <code>dataset</code>
key_var	The variable of <code>dataset_merge</code> containing the names of the transposed variables
value_var	The variable of <code>dataset_merge</code> containing the values of the transposed variables
filter	Expression used to restrict the records of <code>dataset_merge</code> prior to transposing

Details

After filtering `dataset_merge` based upon the condition provided in `filter`, this dataset is transposed and subsequently merged onto `dataset` using `by_vars` as keys.

Value

The input dataset with transposed variables from `dataset_merge` added

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)

cm <- tribble(
  ~USUBJID, ~CMGRPID, ~CMREFID, ~CMDECOD,
  "BP40257-1001", "14", "1192056", "PARACETAMOL",
  "BP40257-1001", "18", "2007001", "SOLUMEDROL",
  "BP40257-1002", "19", "2791596", "SPIRONOLACTONE"
)

facm <- tribble(
  ~USUBJID, ~FAGRPID, ~FAREFID, ~FATESTCD, ~FASTRESC,
  "BP40257-1001", "1", "1192056", "CMATC1CD", "N",
  "BP40257-1001", "1", "1192056", "CMATC2CD", "N02",
  "BP40257-1001", "1", "1192056", "CMATC3CD", "N02B",
  "BP40257-1001", "1", "1192056", "CMATC4CD", "N02BE",
  "BP40257-1001", "1", "2007001", "CMATC1CD", "D",
  "BP40257-1001", "1", "2007001", "CMATC2CD", "D10",
  "BP40257-1001", "1", "2007001", "CMATC3CD", "D10A",
  "BP40257-1001", "1", "2007001", "CMATC4CD", "D10AA",
  "BP40257-1001", "2", "2007001", "CMATC1CD", "D",
  "BP40257-1001", "2", "2007001", "CMATC2CD", "D07",
  "BP40257-1001", "2", "2007001", "CMATC3CD", "D07A",
  "BP40257-1001", "2", "2007001", "CMATC4CD", "D07AA",
  "BP40257-1001", "3", "2007001", "CMATC1CD", "H",
  "BP40257-1001", "3", "2007001", "CMATC2CD", "H02",
  "BP40257-1001", "3", "2007001", "CMATC3CD", "H02A",
  "BP40257-1001", "3", "2007001", "CMATC4CD", "H02AB",
  "BP40257-1002", "1", "2791596", "CMATC1CD", "C",
  "BP40257-1002", "1", "2791596", "CMATC2CD", "C03",
  "BP40257-1002", "1", "2791596", "CMATC3CD", "C03D",
  "BP40257-1002", "1", "2791596", "CMATC4CD", "C03DA"
)

cm %>%
  derive_vars_transposed(
    facm,
    by_vars = exprs(USUBJID, CMREFID = FAREFID),
    key_var = FATESTCD,
    value_var = FASTRESC
  ) %>%
  select(USUBJID, CMDECOD, starts_with("CMATC"))
```

derive_var_agegr_fda *Derive Age Groups*

Description

[Deprecated]

This function is *deprecated*, please create a user defined function instead.

[Deprecated]

This function is *deprecated*, please create a user defined function instead.

Usage

```
derive_var_agegr_fda(dataset, age_var, age_unit = NULL, new_var)
```

```
derive_var_agegr_ema(dataset, age_var, age_unit = NULL, new_var)
```

Arguments

dataset	Input dataset
age_var	AGE variable
age_unit	AGE unit variable
new_var	New variable to create inside dataset

See Also

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

derive_var_age_years *Derive Age in Years*

Description

Converts the given age variable (age_var) to the unit 'years' from the current units given in the age_var+U variable or age_unit argument and stores in a new variable (new_var).

Usage

```
derive_var_age_years(dataset, age_var, age_unit = NULL, new_var)
```

Arguments

dataset	Input dataset. The column specified by the age_var argument is expected.
age_var	Age variable. A numeric object is expected.
age_unit	Age unit. The age_unit argument is only expected when there is NOT a variable age_var+U in dataset. This gives the unit of the age_var variable and is used to convert AGE to 'years' so that grouping can occur. Default: NULL Permitted Values: 'years', 'months', 'weeks', 'days', 'hours', 'minutes', 'seconds'
new_var	New age variable to be created in years. The returned values are doubles and NOT integers. '

Details

This function is used to convert an age variable into the unit 'years' which can then be used to create age groups. The resulting column contains the equivalent years as a double. Note, underlying computations assume an equal number of days in each year (365.25).

Value

The input dataset (dataset) with new_var variable added in years.

Author(s)

Michael Thorpe

See Also

[derive_vars_duration\(\)](#)

ADSL Functions that returns variable appended to dataset: [derive_var_dthcaus\(\)](#), [derive_var_extreme_dtm\(\)](#), [derive_var_extreme_dt\(\)](#), [derive_vars_aage\(\)](#), [derive_vars_period\(\)](#)

Examples

```

library(tibble)

# Derive age with age units specified
data <- tribble(
  ~AGE, ~AGEU,
  27, "days",
  24, "months",
  3, "years",
  4, "weeks",
  1, "years"
)

derive_var_age_years(data, AGE, new_var = AAGE)

# Derive age without age units variable specified
data <- tribble(
  ~AGE,
  12,
  24,
  36,
  48
)
derive_var_age_years(data, AGE, age_unit = "months", new_var = AAGE)

```

```
derive_var_analysis_ratio
```

Derive Ratio Variable

Description

Derives a ratio variable for a BDS dataset based on user specified variables.

Usage

```
derive_var_analysis_ratio(dataset, numer_var, denom_var, new_var = NULL)
```

Arguments

dataset	Input dataset
numer_var	Variable containing numeric values to be used in the numerator of the ratio calculation.
denom_var	Variable containing numeric values to be used in the denominator of the ratio calculation.
new_var	A user-defined variable that will be appended to the dataset. The default behavior will take the denominator variable and prefix it with R2 and append to the dataset. Using this argument will override this default behavior. Default is NULL.

Details

A user wishing to calculate a Ratio to Baseline, AVAL / BASE will have returned a new variable R2BASE that will be appended to the input dataset. Ratio to Analysis Range Lower Limit AVAL / ANRLO will return a new variable R2ANRLO, and Ratio to Analysis Range Upper Limit AVAL / ANRHI will return a new variable R2ANRLO. Please note how the denominator variable has the prefix R2----. A user can override the default returned variables by using the `new_var` argument. Also, values of 0 in the denominator will return NA in the derivation.

Reference CDISC ADaM Implementation Guide Version 1.1 Section 3.3.4 Analysis Parameter Variables for BDS Datasets

Value

The input dataset with a ratio variable appended

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_anrind\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_basetype\(\)](#), [derive_var_base\(\)](#), [derive_var_chg\(\)](#), [derive_var_ontrtfl\(\)](#), [derive_var_pchg\(\)](#), [derive_var_shift\(\)](#)

Examples

```
library(tibble)

data <- tribble(
  ~USUBJID, ~PARAMCD, ~SEQ, ~AVAL, ~BASE, ~ANRLO, ~ANRHI,
  "P01", "ALT", 1, 27, 27, 6, 34,
  "P01", "ALT", 2, 41, 27, 6, 34,
  "P01", "ALT", 3, 17, 27, 6, 34,
  "P02", "ALB", 1, 38, 38, 33, 49,
  "P02", "ALB", 2, 39, 38, 33, 49,
  "P02", "ALB", 3, 37, 38, 33, 49
)

# Returns "R2" prefixed variables
data %>%
  derive_var_analysis_ratio( numer_var = AVAL, denom_var = BASE ) %>%
  derive_var_analysis_ratio( numer_var = AVAL, denom_var = ANRLO ) %>%
  derive_var_analysis_ratio( numer_var = AVAL, denom_var = ANRHI )

# Returns user-defined variables
data %>%
  derive_var_analysis_ratio( numer_var = AVAL, denom_var = BASE, new_var = R01BASE ) %>%
  derive_var_analysis_ratio( numer_var = AVAL, denom_var = ANRLO, new_var = R01ANRLO ) %>%
  derive_var_analysis_ratio( numer_var = AVAL, denom_var = ANRHI, new_var = R01ANRHI )
```

derive_var_anrind	<i>Derive Reference Range Indicator</i>
-------------------	---

Description

Derive Reference Range Indicator

Usage

```
derive_var_anrind(dataset)
```

Arguments

dataset	The input dataset
---------	-------------------

Details

ANRIND is set to

- "NORMAL" if AVAL is greater or equal ANRLO and less than or equal ANRHI; or if AVAL is greater than or equal ANRLO and ANRHI is missing; or if AVAL is less than or equal ANRHI and ANRLO is missing
- "LOW" if AVAL is less than ANRLO and either A1LO is missing or AVAL is greater than or equal A1LO
- "HIGH" if AVAL is greater than ANRHI and either A1HI is missing or AVAL is less than or equal A1HI
- "LOW LOW" if AVAL is less than A1LO
- "HIGH HIGH" if AVAL is greater than A1HI

Value

The input dataset with additional column ANRIND

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_basetype\(\)](#), [derive_var_base\(\)](#), [derive_var_chg\(\)](#), [derive_var_ontrtfl\(\)](#), [derive_var_pchg\(\)](#), [derive_var_shift\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data(admiral_vs)

ref_ranges <- tribble(
```

```

~PARAMCD, ~ANRLO, ~ANRHI, ~A1LO, ~A1HI,
"DIABP",    60,    80,    40,    90,
"PULSE",    60,    100,   40,    110
)

admiral_vs %>%
  mutate(
    PARAMCD = VSTESTCD,
    AVAL = VSSTRESN
  ) %>%
  filter(PARAMCD %in% c("PULSE", "DIABP")) %>%
  derive_vars_merged(ref_ranges, by_vars = exprs(PARAMCD)) %>%
  derive_var_anrind() %>%
  select(USUBJID, PARAMCD, AVAL, ANRLO:ANRIND)

```

derive_var_atoxgr *Derive Lab High toxicity Grade 0 - 4 and Low Toxicity Grades 0 - (-4)*

Description

Derives character lab grade based on high and low severity/toxicity grade(s).

Usage

```

derive_var_atoxgr(
  dataset,
  lotox_description_var = ATOXDSCCL,
  hitox_description_var = ATOXDSCCH
)

```

Arguments

dataset	Input data set The columns ATOXGRL, ATOXGRH and specified by lotox_description_var, and hitox_description_var parameters are expected.
lotox_description_var	Variable containing the toxicity grade description for low values, eg. "Anemia"
hitox_description_var	Variable containing the toxicity grade description for low values, eg. "Hemoglobin Increased".

Details

Created variable ATOXGR will contain values "-4", "-3", "-2", "-1" for low values and "1", "2", "3", "4" for high values, and will contain "0" if value is gradable and does not satisfy any of the criteria for high or low values. ATOXGR is set to missing if information not available to give a grade.

Function applies the following rules:

- High and low missing - overall missing
- Low grade not missing and > 0 - overall holds low grade
- High grade not missing and > 0 - overall holds high grade
- (Only high direction OR low direction is NORMAL) and high grade normal - overall NORMAL
- (Only low direction OR high direction is NORMAL) and low grade normal - overall NORMAL
- otherwise set to missing

Value

The input data set with the character variable added

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_anrind\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_basetype\(\)](#), [derive_var_base\(\)](#), [derive_var_chg\(\)](#), [derive_var_ontrtfl\(\)](#), [derive_var_pchg\(\)](#), [derive_var_shift\(\)](#)

Examples

```
library(tibble)

adlb <- tribble(
  ~ATOXDACL,      ~ATOXDACH,      ~ATOXDACH,      ~ATOXDACH,
  "Hypoglycemia", "Hyperglycemia", NA_character_, "0",
  "Hypoglycemia", "Hyperglycemia", "0",      "1",
  "Hypoglycemia", "Hyperglycemia", "0",      "0",
  NA_character_,  "INR Increased", NA_character_, "0",
  "Hypophosphatemia", NA_character_, "1",      NA_character_
)

derive_var_atoxgr(adlb)
```

derive_var_atoxgr_dir *Derive Lab Toxicity Grade 0 - 4*

Description

Derives a character lab grade based on severity/toxicity criteria.

Usage

```

derive_var_atoxgr_dir(
  dataset,
  new_var,
  tox_description_var,
  meta_criteria,
  criteria_direction,
  get_unit_expr
)

```

Arguments

dataset	Input data set The columns specified by tox_description_var parameter is expected.
new_var	Name of the character grade variable to create, for example, ATOXGRH or ATOXGRL.
tox_description_var	Variable containing the description of the grading criteria. For example: "Anemia" or "INR Increased".
meta_criteria	Metadata data set holding the criteria (normally a case statement) Permitted Values: atoxgr_criteria_ctcv4, atoxgr_criteria_ctcv5 admiral metadata data set atoxgr_criteria_ctcv4 implements Common Terminology Criteria for Adverse Events (CTCAE) v4.0 admiral metadata data set atoxgr_criteria_ctcv5 implements Common Terminology Criteria for Adverse Events (CTCAE) v5.0 The metadata should have the following variables: <ul style="list-style-type: none"> • TERM: variable to hold the term describing the criteria applied to a particular lab test, eg. "Anemia" or "INR Increased". Note: the variable is case insensitive. • DIRECTION: variable to hold the direction of the abnormality of a particular lab test value. "L" is for LOW values, "H" is for HIGH values. Note: the variable is case insensitive. • SI_UNIT_CHECK: variable to hold unit of particular lab test. Used to check against input data if criteria is based on absolute values. • VAR_CHECK: variable to hold comma separated list of variables used in criteria. Used to check against input data that variables exist. • GRADE_CRITERIA_CODE: variable to hold code that creates grade based on defined criteria.
criteria_direction	Direction (L= Low, H = High) of toxicity grade. Permitted Values: "L", "H"
get_unit_expr	An expression providing the unit of the parameter The result is used to check the units of the input parameters. Compared with SI_UNIT_CHECK in metadata (see meta_criteria parameter). Permitted Values: A variable containing unit from the input dataset, or a function call, for example, get_unit_expr = extract_unit(PARAM).

Details

new_var is derived with values NA, "0", "1", "2", "3", "4", where "4" is the most severe grade

- "4" is where the lab value satisfies the criteria for grade 4.
- "3" is where the lab value satisfies the criteria for grade 3.
- "2" is where the lab value satisfies the criteria for grade 2.
- "1" is where the lab value satisfies the criteria for grade 1.
- "0" is where a grade can be derived and is not grade "1", "2", "3" or "4".
- NA is where a grade cannot be derived.

Value

The input dataset with the character variable added

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_anrind\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_basetype\(\)](#), [derive_var_base\(\)](#), [derive_var_chg\(\)](#), [derive_var_ontrftl\(\)](#), [derive_var_pchg\(\)](#), [derive_var_shift\(\)](#)

Examples

```
library(tibble)

data <- tribble(
  ~ATOXDSCS,           ~AVAL, ~ANRLO, ~ANRHI, ~PARAM,
  "Hypoglycemia",     119,   4,      7,      "Glucose (mmol/L)",
  "Hypoglycemia",     120,   4,      7,      "Glucose (mmol/L)",
  "Anemia",           129,  120,    180,    "Hemoglobin (g/L)",
  "White blood cell decreased", 10,   5,      20,    "White blood cell (10^9/L)",
  "White blood cell decreased", 15,   5,      20,    "White blood cell (10^9/L)",
  "Anemia",           140,  120,    180,    "Hemoglobin (g/L)"
)

derive_var_atoxgr_dir(data,
  new_var = ATOXGRL,
  tox_description_var = ATOXDSCS,
  meta_criteria = atoxgr_criteria_ctcv4,
  criteria_direction = "L",
  get_unit_expr = extract_unit(PARAM)
)

data <- tribble(
  ~ATOXDSCS,           ~AVAL, ~ANRLO, ~ANRHI, ~PARAM,
  "Hyperglycemia",     119,   4,      7,      "Glucose (mmol/L)",
  "Hyperglycemia",     120,   4,      7,      "Glucose (mmol/L)",
  "GGT increased",     129,   0,      30,    "Gamma Glutamyl Transferase (U/L)",
  "Lymphocyte count increased", 4,    1,      4,      "Lymphocytes Abs (10^9/L)",
  "Lymphocyte count increased", 2,    1,      4,      "Lymphocytes Abs (10^9/L)",
  "GGT increased",     140,  120,    180,    "Gamma Glutamyl Transferase (U/L)"
)
```

```

)

derive_var_atoxgr_dir(data,
  new_var = ATOXGRH,
  tox_description_var = ATOXDSCH,
  meta_criteria = atoxgr_criteria_ctcv4,
  criteria_direction = "H",
  get_unit_expr = extract_unit(PARAM)
)

```

derive_var_base *Derive Baseline Variables*

Description

Derive baseline variables, e.g. BASE or BNRIND, in a BDS dataset

Usage

```

derive_var_base(
  dataset,
  by_vars,
  source_var = AVAL,
  new_var = BASE,
  filter = ABLFL == "Y"
)

```

Arguments

dataset	The input dataset
by_vars	Grouping variables uniquely identifying a set of records for which to calculate new_var
source_var	The column from which to extract the baseline value, e.g. AVAL
new_var	The name of the newly created baseline column, e.g. BASE
filter	The condition used to filter dataset for baseline records. By default ABLFL == "Y"

Details

For each `by_vars` group, the baseline record is identified by the condition specified in `filter` which defaults to `ABLFL == "Y"`. Subsequently, every value of the `new_var` variable for the `by_vars` group is set to the value of the `source_var` variable of the baseline record. In case there are multiple baseline records within `by_vars` an error is issued.

Value

A new `data.frame` containing all records and variables of the input dataset plus the `new_var` variable

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_anrind\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_basetype\(\)](#), [derive_var_chg\(\)](#), [derive_var_ontrtfl\(\)](#), [derive_var_pchg\(\)](#), [derive_var_shift\(\)](#)

Examples

```
library(tibble)

dataset <- tribble(
  ~STUDYID, ~USUBJID, ~PARAMCD, ~AVAL, ~AVALC, ~AVISIT, ~ABLFL, ~ANRIND,
  "TEST01", "PAT01", "PARAM01", 10.12, NA, "Baseline", "Y", "NORMAL",
  "TEST01", "PAT01", "PARAM01", 9.700, NA, "Day 7", "N", "LOW",
  "TEST01", "PAT01", "PARAM01", 15.01, NA, "Day 14", "N", "HIGH",
  "TEST01", "PAT01", "PARAM02", 8.350, NA, "Baseline", "Y", "LOW",
  "TEST01", "PAT01", "PARAM02", NA, NA, "Day 7", "N", NA,
  "TEST01", "PAT01", "PARAM02", 8.350, NA, "Day 14", "N", "LOW",
  "TEST01", "PAT01", "PARAM03", NA, "LOW", "Baseline", "Y", NA,
  "TEST01", "PAT01", "PARAM03", NA, "LOW", "Day 7", "N", NA,
  "TEST01", "PAT01", "PARAM03", NA, "MEDIUM", "Day 14", "N", NA,
  "TEST01", "PAT01", "PARAM04", NA, "HIGH", "Baseline", "Y", NA,
  "TEST01", "PAT01", "PARAM04", NA, "HIGH", "Day 7", "N", NA,
  "TEST01", "PAT01", "PARAM04", NA, "MEDIUM", "Day 14", "N", NA
)

## Derive `BASE` variable from `AVAL`
derive_var_base(
  dataset,
  by_vars = exprs(USUBJID, PARAMCD),
  source_var = AVAL,
  new_var = BASE
)

## Derive `BASEC` variable from `AVALC`
derive_var_base(
  dataset,
  by_vars = exprs(USUBJID, PARAMCD),
  source_var = AVALC,
  new_var = BASEC
)

## Derive `BNRIND` variable from `ANRIND`
derive_var_base(
  dataset,
  by_vars = exprs(USUBJID, PARAMCD),
  source_var = ANRIND,
  new_var = BNRIND
)
```

derive_var_basetype *Derive Basetype Variable*

Description

Baseline Type BASETYPE is needed when there is more than one definition of baseline for a given Analysis Parameter PARAM in the same dataset. For a given parameter, if Baseline Value BASE is populated, and there is more than one definition of baseline, then BASETYPE must be non-null on all records of any type for that parameter. Each value of BASETYPE refers to a definition of baseline that characterizes the value of BASE on that row. Please see section 4.2.1.6 of the ADaM Implementation Guide, version 1.3 for further background.

Usage

```
derive_var_basetype(dataset, basetypes)
```

Arguments

dataset	Input dataset The columns specified in the expressions inside basetypes are required.
basetypes	A <i>named</i> list of expressions created using the <code>rlang::exprs()</code> function The names corresponds to the values of the newly created BASETYPE variables and the expressions are used to subset the input dataset.

Details

Adds the BASETYPE variable to a dataset and duplicates records based upon the provided conditions.

For each element of basetypes the input dataset is subset based upon the provided expression and the BASETYPE variable is set to the name of the expression. Then, all subsets are stacked. Records which do not match any condition are kept and BASETYPE is set to NA.

Value

The input dataset with variable BASETYPE added

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_anrind\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_base\(\)](#), [derive_var_chg\(\)](#), [derive_var_ontrftl\(\)](#), [derive_var_pchg\(\)](#), [derive_var_shift\(\)](#)

Examples

```

library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(rlang)

bds <- tribble(
  ~USUBJID, ~EPOCH,      ~PARAMCD, ~ASEQ, ~AVAL,
  "P01",    "RUN-IN",    "PARAM01", 1, 10.0,
  "P01",    "RUN-IN",    "PARAM01", 2, 9.8,
  "P01",    "DOUBLE-BLIND", "PARAM01", 3, 9.2,
  "P01",    "DOUBLE-BLIND", "PARAM01", 4, 10.1,
  "P01",    "OPEN-LABEL",  "PARAM01", 5, 10.4,
  "P01",    "OPEN-LABEL",  "PARAM01", 6, 9.9,
  "P02",    "RUN-IN",    "PARAM01", 1, 12.1,
  "P02",    "DOUBLE-BLIND", "PARAM01", 2, 10.2,
  "P02",    "DOUBLE-BLIND", "PARAM01", 3, 10.8,
  "P02",    "OPEN-LABEL",  "PARAM01", 4, 11.4,
  "P02",    "OPEN-LABEL",  "PARAM01", 5, 10.8
)

bds_with_basetype <- derive_var_basetype(
  dataset = bds,
  basetypes = exprs(
    "RUN-IN" = EPOCH %in% c("RUN-IN", "STABILIZATION", "DOUBLE-BLIND", "OPEN-LABEL"),
    "DOUBLE-BLIND" = EPOCH %in% c("DOUBLE-BLIND", "OPEN-LABEL"),
    "OPEN-LABEL" = EPOCH == "OPEN-LABEL"
  )
)

# Below print statement will print all 23 records in the data frame
# bds_with_basetype
print(bds_with_basetype, n = Inf)

count(bds_with_basetype, BASETYPE, name = "Number of Records")

# An example where all parameter records need to be included for 2 different
# baseline type derivations (such as LAST and WORST)
bds <- tribble(
  ~USUBJID, ~EPOCH,      ~PARAMCD, ~ASEQ, ~AVAL,
  "P01",    "RUN-IN",    "PARAM01", 1, 10.0,
  "P01",    "RUN-IN",    "PARAM01", 2, 9.8,
  "P01",    "DOUBLE-BLIND", "PARAM01", 3, 9.2,
  "P01",    "DOUBLE-BLIND", "PARAM01", 4, 10.1
)

bds_with_basetype <- derive_var_basetype(
  dataset = bds,
  basetypes = exprs(
    "LAST" = TRUE,
    "WORST" = TRUE
  )
)

```

```

)

print(bds_with_basetype, n = Inf)

count(bds_with_basetype, BASETYPE, name = "Number of Records")

```

derive_var_chg	<i>Derive Change from Baseline</i>
----------------	------------------------------------

Description

Derive change from baseline (CHG) in a BDS dataset

Usage

```
derive_var_chg(dataset)
```

Arguments

dataset The input dataset. Required variables are AVAL and BASE.

Details

Change from baseline is calculated by subtracting the baseline value from the analysis value.

Value

The input dataset with an additional column named CHG

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_anrind\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_basetype\(\)](#), [derive_var_base\(\)](#), [derive_var_ontrtfl\(\)](#), [derive_var_pchg\(\)](#), [derive_var_shift\(\)](#)

Examples

```

library(tibble)

advs <- tribble(
  ~USUBJID, ~PARAMCD, ~AVAL, ~ABLFL, ~BASE,
  "P01", "WEIGHT", 80, "Y", 80,
  "P01", "WEIGHT", 80.8, "", 80,
  "P01", "WEIGHT", 81.4, "", 80,
  "P02", "WEIGHT", 75.3, "Y", 75.3,
  "P02", "WEIGHT", 76, "", 75.3
)

derive_var_chg(advs)

```

`derive_var_confirmation_flag`*Derive Confirmation Flag*

Description

[Deprecated]

This function is *deprecated*, please use `derive_var_joined_exist_flag()` instead.

Usage

```
derive_var_confirmation_flag(  
  dataset,  
  by_vars,  
  order,  
  new_var,  
  tmp_obs_nr_var = NULL,  
  join_vars,  
  join_type,  
  first_cond = NULL,  
  filter,  
  true_value = "Y",  
  false_value = NA_character_,  
  check_type = "warning"  
)
```

Arguments

<code>dataset</code>	Input dataset The variables specified by the <code>by_vars</code> and <code>join_vars</code> parameter are expected.
<code>by_vars</code>	By variables The specified variables are used as by variables for joining the input dataset with itself.
<code>order</code>	Order The observations are ordered by the specified order.
<code>new_var</code>	New variable The specified variable is added to the input dataset.
<code>tmp_obs_nr_var</code>	Temporary observation number The specified variable is added to the input dataset and set to the observation number with respect to <code>order</code> . For each by group (<code>by_vars</code>) the observation number starts with 1. The variable can be used in the conditions (<code>filter</code> , <code>first_cond</code>). It is not included in the output dataset. It can be used to flag consecutive observations or the last observation (see last example below).

join_vars	<p>Variables to keep from joined dataset</p> <p>The variables needed from the other observations should be specified for this parameter. The specified variables are added to the joined dataset with suffix ".join". For example to flag all observations with AVALC == "Y" and AVALC == "Y" for at least one subsequent visit join_vars = exprs(AVALC, AVISITN) and filter = AVALC == "Y" & AVALC.join == "Y" & AVISITN < AVISITN.join could be specified.</p> <p>The *.join variables are not included in the output dataset.</p>
join_type	<p>Observations to keep after joining</p> <p>The argument determines which of the joined observations are kept with respect to the original observation. For example, if join_type = "after" is specified all observations after the original observations are kept.</p> <p>For example for confirmed response or BOR in the oncology setting or confirmed deterioration in questionnaires the confirmatory assessment must be after the assessment to be flagged. Thus join_type = "after" could be used.</p> <p>Whereas, sometimes you might allow for confirmatory observations to occur prior to the observation to be flagged. For example, to flag AEs occurring on or after seven days before a COVID AE. Thus join_type = "all" could be used.</p> <p><i>Permitted Values:</i> "before", "after", "all"</p>
first_cond	<p>Condition for selecting range of data</p> <p>If this argument is specified, the other observations are restricted up to the first observation where the specified condition is fulfilled. If the condition is not fulfilled for any of the other observations, no observations are considered, i.e., the observation is not flagged.</p> <p>This parameter should be specified if filter contains summary functions which should not apply to all observations but only up to the confirmation assessment. For an example see the third example below.</p>
filter	<p>Condition for selecting observations</p> <p>The filter is applied to the joined dataset for flagging the confirmed observations. The condition can include summary functions. The joined dataset is grouped by the original observations. I.e., the summary function are applied to all observations up to the confirmation observation. For example, filter = AVALC == "CR" & all(AVALC.join %in% c("CR", "NE")) & count_vals(var = AVALC.join, val = "NE") <= 1 selects observations with response "CR" and for all observations up to the confirmation observation the response is "CR" or "NE" and there is at most one "NE".</p>
true_value	<p>Value of new_var for flagged observations</p> <p><i>Default:</i> "Y"</p>
false_value	<p>Value of new_var for observations not flagged</p> <p><i>Default:</i> NA_character_</p>
check_type	<p>Check uniqueness?</p> <p>If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order.</p> <p><i>Default:</i> "warning"</p> <p><i>Permitted Values:</i> "none", "warning", "error"</p>

Details

The following steps are performed to produce the output dataset.

Step 1:

The input dataset is joined with itself by the variables specified for `by_vars`. From the right hand side of the join only the variables specified for `join_vars` are kept. The suffix ".join" is added to these variables.

For example, for `by_vars = USUBJID`, `join_vars = exprs(AVISITN, AVALC)` and input dataset

```
# A tibble: 2 x 4
  USUBJID AVISITN AVALC  AVAL
<chr>    <dbl> <chr> <dbl>
1         1 Y      1
1         2 N      0
```

the joined dataset is

```
A tibble: 4 x 6
  USUBJID AVISITN AVALC  AVAL AVISITN.join AVALC.join
<chr>    <dbl> <chr> <dbl> <dbl> <chr>
1         1 Y      1      1 Y
1         1 Y      1      2 N
1         2 N      0      1 Y
1         2 N      0      2 N
```

Step 2:

The joined dataset is restricted to observations with respect to `join_type` and `order`.

The dataset from the example in the previous step with `join_type = "after"` and `order = exprs(AVISITN)` is restricted to

```
A tibble: 4 x 6
  USUBJID AVISITN AVALC  AVAL AVISITN.join AVALC.join
<chr>    <dbl> <chr> <dbl> <dbl> <chr>
1         1 Y      1      1 2 N
```

Step 3:

If `first_cond` is specified, for each observation of the input dataset the joined dataset is restricted to observations up to the first observation where `first_cond` is fulfilled (the observation fulfilling the condition is included). If for an observation of the input dataset the condition is not fulfilled, the observation is removed.

Step 4:

The joined dataset is grouped by the observations from the input dataset and restricted to the observations fulfilling the condition specified by `filter`.

Step 5:

The first observation of each group is selected

Step 6:

The variable specified by `new_var` is added to the input dataset. It is set to `true_value` for all observations which were selected in the previous step. For the other observations it is set to `false_value`.

Value

The input dataset with the variable specified by new_var added.

See Also

Other deprecated: `derive_derived_param()`, `derive_param_first_event()`, `derive_var_agegr_fda()`, `derive_var_disposition_status()`, `derive_var_worst_flag()`, `derive_vars_disposition_reason()`, `derive_vars_merged_dtm()`, `derive_vars_merged_dt()`, `filter_confirmation()`, `format_eoxxstt_default()`, `format_reason_default()`, `vars()`

derive_var_disposition_status

Derive a Disposition Status at a Specific Timepoint

Description**[Deprecated]**

This function is *deprecated*, Please define your own function and use that as input for the cat_fun argument in `derive_var_merged_cat()` instead.

Derive a disposition status from the the relevant records in the disposition domain.

Usage

```
derive_var_disposition_status(
  dataset,
  dataset_ds,
  new_var,
  status_var,
  format_new_var = format_eoxxstt_default,
  filter_ds,
  subject_keys = get_admiral_option("subject_keys")
)
```

Arguments

dataset	Input dataset.
dataset_ds	Dataset containing the disposition information (e.g.: ds). It must contain: <ul style="list-style-type: none"> • STUDYID, USUBJID, • The variable(s) specified in the status_var • The variables used in filter_ds.
new_var	Name of the disposition status variable. A variable name is expected (e.g. EOSSTT).

status_var	The variable used to derive the disposition status. A variable name is expected (e.g. DSDECOD).
format_new_var	The format used to derive the status. Default: format_eoxxstt_default() defined as: <pre>format_eoxxstt_default <- function(status) { case_when(status %in% c("SCREEN FAILURE", "SCREENING NOT COMPLETED") ~ "NOT STARTED", status == "COMPLETED" ~ "COMPLETED", !status %in% c("COMPLETED", "SCREEN FAILURE", "SCREENING NOT COMPLETED") & !is.na(status) ~ "DISCONTINUED", TRUE ~ "ONGOING") }</pre> where status is the status_var.
filter_ds	Filter condition for the disposition data. one observation per patient. An error is issued otherwise. Permitted Values: logical expression.
subject_keys	Variables to uniquely identify a subject A list of expressions where the expressions are symbols as returned by exprs() is expected.

Value

The input dataset with the disposition status (new_var) added. new_var is derived based on the values given in status_var and according to the format defined by format_new_var (e.g. when the default format is used, the function will derive new_var as: "NOT STARTED" if status is "SCREEN FAILURE" or "SCREENING NOT COMPLETED", "COMPLETED" if status_var == "COMPLETED", "DISCONTINUED" if status is not in ("COMPLETED", "SCREEN FAILURE", "SCREENING NOT COMPLETED") nor NA, "ONGOING" otherwise).

See Also

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

derive_var_dthcaus *Derive Death Cause*

Description

Derive death cause (DTHCAUS) and add traceability variables if required.

Usage

```
derive_var_dthcaus(
  dataset,
  ...,
  source_datasets,
  subject_keys = get_admiral_option("subject_keys")
)
```

Arguments

dataset	Input dataset. The variables specified by subject_keys are required.
...	Objects of class "dthcaus_source" created by dthcaus_source() .
source_datasets	A named list containing datasets in which to search for the death cause
subject_keys	Variables to uniquely identify a subject A list of expressions where the expressions are symbols as returned by exprs() is expected.

Details

This function derives DTHCAUS along with the user-defined traceability variables, if required. If a subject has death info from multiple sources, the one from the source with the earliest death date will be used. If dates are equivalent, the first source will be kept, so the user should provide the inputs in the preferred order.

Value

The input dataset with DTHCAUS variable added.

See Also

[dthcaus_source\(\)](#)

ADSL Functions that returns variable appended to dataset: [derive_var_age_years\(\)](#), [derive_var_extreme_dtm\(\)](#), [derive_var_extreme_dt\(\)](#), [derive_vars_aage\(\)](#), [derive_vars_period\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

adsl <- tribble(
  ~STUDYID, ~USUBJID,
  "STUDY01", "PAT01",
  "STUDY01", "PAT02",
  "STUDY01", "PAT03"
)
```

```

ae <- tribble(
  ~STUDYID, ~USUBJID, ~AESEQ, ~AEDECOD, ~AEOUT, ~AEDTHDTC,
  "STUDY01", "PAT01", 12, "SUDDEN DEATH", "FATAL", "2021-04-04"
) %>%
  mutate(
    AEDTHDT = ymd(AEDTHDTC)
  )
ds <- tribble(
  ~STUDYID, ~USUBJID, ~DSSEQ, ~DSDECOD, ~DSTERM, ~DSSTDTC,
  "STUDY01", "PAT02", 1, "INFORMED CONSENT OBTAINED", "INFORMED CONSENT OBTAINED", "2021-04-03",
  "STUDY01", "PAT02", 2, "RANDOMIZATION", "RANDOMIZATION", "2021-04-11",
  "STUDY01", "PAT02", 3, "DEATH", "DEATH DUE TO PROGRESSION OF DISEASE", "2022-02-01",
  "STUDY01", "PAT03", 1, "DEATH", "POST STUDY REPORTING OF DEATH", "2022-03-03"
) %>%
  mutate(
    DSSTDTC = ymd(DSSTDTC)
  )

# Derive `DTHCAUS` only - for on-study deaths only
src_ae <- dthcaus_source(
  dataset_name = "ae",
  filter = AEOUT == "FATAL",
  date = AEDTHDT,
  mode = "first",
  dthcaus = AEDECOD
)

src_ds <- dthcaus_source(
  dataset_name = "ds",
  filter = DSDECOD == "DEATH" & grepl("DEATH DUE TO", DSTERM),
  date = DSSTDTC,
  mode = "first",
  dthcaus = DSTERM
)

derive_var_dthcaus(ads1, src_ae, src_ds, source_datasets = list(ae = ae, ds = ds))

# Derive `DTHCAUS` and add traceability variables - for on-study deaths only
src_ae <- dthcaus_source(
  dataset_name = "ae",
  filter = AEOUT == "FATAL",
  date = AEDTHDT,
  mode = "first",
  dthcaus = AEDECOD,
  traceability_vars = exprs(DTHDOM = "AE", DTHSEQ = AESEQ)
)

src_ds <- dthcaus_source(
  dataset_name = "ds",
  filter = DSDECOD == "DEATH" & grepl("DEATH DUE TO", DSTERM),
  date = DSSTDTC,
  mode = "first",
  dthcaus = DSTERM,

```

```

    traceability_vars = exprs(DTHDOM = "DS", DTHSEQ = DSSEQ)
  )

derive_var_dthcaus(adsl, src_ae, src_ds, source_datasets = list(ae = ae, ds = ds))

# Derive `DTHCAUS` as above - now including post-study deaths with different `DTHCAUS` value
src_ae <- dthcaus_source(
  dataset_name = "ae",
  filter = AEOUT == "FATAL",
  date = AEDTHDT,
  mode = "first",
  dthcaus = AEDECOD,
  traceability_vars = exprs(DTHDOM = "AE", DTHSEQ = AESEQ)
)

src_ds <- dthcaus_source(
  dataset_name = "ds",
  filter = DSDECOD == "DEATH" & grepl("DEATH DUE TO", DSTERM),
  date = DSSTDT,
  mode = "first",
  dthcaus = DSTERM,
  traceability_vars = exprs(DTHDOM = "DS", DTHSEQ = DSSEQ)
)

src_ds_post <- dthcaus_source(
  dataset_name = "ds",
  filter = DSDECOD == "DEATH" & DSTERM == "POST STUDY REPORTING OF DEATH",
  date = DSSTDT,
  mode = "first",
  dthcaus = "POST STUDY: UNKNOWN CAUSE",
  traceability_vars = exprs(DTHDOM = "DS", DTHSEQ = DSSEQ)
)

derive_var_dthcaus(adsl, src_ae, src_ds, src_ds_post, source_datasets = list(ae = ae, ds = ds))

```

derive_var_extreme_dt *Derive First or Last Date from Multiple Sources*

Description

Add the first or last date from multiple sources to the dataset, e.g., the last known alive date (LSTALVDT).

Usage

```

derive_var_extreme_dt(
  dataset,
  new_var,
  ...,
  source_datasets,

```

```

mode,
subject_keys = get_admiral_option("subject_keys")
)

```

Arguments

dataset	Input dataset The variables specified by subject_keys are required.
new_var	Name of variable to create
...	Source(s) of dates. One or more date_source() objects are expected.
source_datasets	A named list containing datasets in which to search for the first or last date
mode	Selection mode (first or last) If "first" is specified, the first date for each subject is selected. If "last" is specified, the last date for each subject is selected. Permitted Values: "first", "last"
subject_keys	Variables to uniquely identify a subject A list of expressions where the expressions are symbols as returned by exprs() is expected.

Details

The following steps are performed to create the output dataset:

1. For each source dataset the observations as specified by the filter element are selected and observations where date is NA are removed. Then for each patient the first or last observation (with respect to date and mode) is selected.
2. The new variable is set to the variable specified by the date element.
3. The variables specified by the traceability_vars element are added.
4. The selected observations of all source datasets are combined into a single dataset.
5. For each patient the first or last observation (with respect to the new variable and mode) from the single dataset is selected and the new variable is merged to the input dataset.
6. The time part is removed from the new variable.

Value

The input dataset with the new variable added.

See Also

[date_source\(\)](#), [derive_var_extreme_dtm\(\)](#), [derive_vars_merged\(\)](#)

ADSL Functions that returns variable appended to dataset: [derive_var_age_years\(\)](#), [derive_var_dthcaus\(\)](#), [derive_var_extreme_dtm\(\)](#), [derive_vars_aage\(\)](#), [derive_vars_period\(\)](#)

Examples

```

library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data("admiral_dm")
data("admiral_ae")
data("admiral_lb")
data("admiral_adsl")

# derive last known alive date (LSTALVDT)
ae_start <- date_source(
  dataset_name = "ae",
  date = AESTDT
)
ae_end <- date_source(
  dataset_name = "ae",
  date = AEENDT
)

ae_ext <- admiral_ae %>%
  derive_vars_dt(
    dtc = AESTDTC,
    new_vars_prefix = "AEST",
    highest_imputation = "M"
  ) %>%
  derive_vars_dt(
    dtc = AEENDTC,
    new_vars_prefix = "AEEN",
    highest_imputation = "M"
  )

lb_date <- date_source(
  dataset_name = "lb",
  date = LBDT,
  filter = !is.na(LBDT),
)

lb_ext <- derive_vars_dt(
  admiral_lb,
  dtc = LBDTC,
  new_vars_prefix = "LB"
)

adsl_date <- date_source(dataset_name = "adsl", date = TRTEDT)

admiral_dm %>%
  derive_var_extreme_dt(
    new_var = LSTALVDT,
    ae_start, ae_end, lb_date, adsl_date,
    source_datasets = list(
      adsl = admiral_adsl,
      ae = ae_ext,
      lb = lb_ext
    )
  )

```

```

    ),
    mode = "last"
  ) %>%
  select(USUBJID, LSTALVDT)

# derive last alive date and traceability variables
ae_start <- date_source(
  dataset_name = "ae",
  date = AESTDT,
  traceability_vars = exprs(
    LALVDOM = "AE",
    LALVSEQ = AESEQ,
    LALVVAR = "AESTDTC"
  )
)

ae_end <- date_source(
  dataset_name = "ae",
  date = AEENDT,
  traceability_vars = exprs(
    LALVDOM = "AE",
    LALVSEQ = AESEQ,
    LALVVAR = "AEENDTC"
  )
)

lb_date <- date_source(
  dataset_name = "lb",
  date = LBDT,
  filter = !is.na(LBDT),
  traceability_vars = exprs(
    LALVDOM = "LB",
    LALVSEQ = LBSEQ,
    LALVVAR = "LBDTC"
  )
)

adsl_date <- date_source(
  dataset_name = "adsl",
  date = TRTEDT,
  traceability_vars = exprs(
    LALVDOM = "ADSL",
    LALVSEQ = NA_integer_,
    LALVVAR = "TRTEDT"
  )
)

admiral_dm %>%
  derive_var_extreme_dt(
    new_var = LSTALVDT,
    ae_start, ae_end, lb_date, adsl_date,
    source_datasets = list(
      adsl = admiral_adsl,
      ae = ae_ext,

```

```

        lb = lb_ext
      ),
      mode = "last"
    ) %>%
  select(USUBJID, LSTALVDT, LALVDOM, LALVSEQ, LALVVAR)

```

 derive_var_extreme_dtm

Derive First or Last Datetime from Multiple Sources

Description

Add the first or last datetime from multiple sources to the dataset, e.g., the last known alive datetime (LSTALVDTM).

Usage

```

derive_var_extreme_dtm(
  dataset,
  new_var,
  ...,
  source_datasets,
  mode,
  subject_keys = get_admiral_option("subject_keys")
)

```

Arguments

dataset	Input dataset The variables specified by subject_keys are required.
new_var	Name of variable to create
...	Source(s) of dates. One or more date_source() objects are expected.
source_datasets	A named list containing datasets in which to search for the first or last date
mode	Selection mode (first or last) If "first" is specified, the first date for each subject is selected. If "last" is specified, the last date for each subject is selected. Permitted Values: "first", "last"
subject_keys	Variables to uniquely identify a subject A list of expressions where the expressions are symbols as returned by exprs() is expected.

Details

The following steps are performed to create the output dataset:

1. For each source dataset the observations as specified by the `filter` element are selected and observations where date is NA are removed. Then for each patient the first or last observation (with respect to date and mode) is selected.
2. The new variable is set to the variable specified by the `date` element. If this is a date variable (rather than datetime), then the time is imputed as "00:00:00".
3. The variables specified by the `traceability_vars` element are added.
4. The selected observations of all source datasets are combined into a single dataset.
5. For each patient the first or last observation (with respect to the new variable and mode) from the single dataset is selected and the new variable is merged to the input dataset.

Value

The input dataset with the new variable added.

See Also

[date_source\(\)](#), [derive_var_extreme_dt\(\)](#), [derive_vars_merged\(\)](#)

ADSL Functions that returns variable appended to dataset: [derive_var_age_years\(\)](#), [derive_var_dthcaus\(\)](#), [derive_var_extreme_dt\(\)](#), [derive_vars_aage\(\)](#), [derive_vars_period\(\)](#)

Examples

```
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data("admiral_dm")
data("admiral_ae")
data("admiral_lb")
data("admiral_ads1")

# derive last known alive datetime (LSTALVDTM)
ae_start <- date_source(
  dataset_name = "ae",
  date = AESTDTM
)
ae_end <- date_source(
  dataset_name = "ae",
  date = AEENDTM
)

ae_ext <- admiral_ae %>%
  derive_vars_dtm(
    dtc = AESTDTC,
    new_vars_prefix = "AEST",
    highest_imputation = "M"
  ) %>%
  derive_vars_dtm(
```



```

    dtc = AEENDTC,
    new_vars_prefix = "AEEN",
    highest_imputation = "M"
  )

lb_date <- date_source(
  dataset_name = "lb",
  date = LBDTM,
  filter = !is.na(LBDTM)
)

lb_ext <- derive_vars_dtm(
  admiral_lb,
  dtc = LBDTC,
  new_vars_prefix = "LB"
)

adsl_date <- date_source(dataset_name = "adsl", date = TRTEDTM)

admiral_dm %>%
  derive_var_extreme_dtm(
    new_var = LSTALVDTM,
    ae_start, ae_end, lb_date, adsl_date,
    source_datasets = list(
      adsl = admiral_adsl,
      ae = ae_ext, lb = lb_ext
    ),
    mode = "last"
  ) %>%
  select(USUBJID, LSTALVDTM)

# derive last alive datetime and traceability variables
ae_start <- date_source(
  dataset_name = "ae",
  date = AESTDTM,
  traceability_vars = exprs(
    LALVDOM = "AE",
    LALVSEQ = AESEQ,
    LALVVAR = "AESTDTC"
  )
)

ae_end <- date_source(
  dataset_name = "ae",
  date = AEENDTM,
  traceability_vars = exprs(
    LALVDOM = "AE",
    LALVSEQ = AESEQ,
    LALVVAR = "AEENDTC"
  )
)

lb_date <- date_source(
  dataset_name = "lb",

```

```

    date = LBDTM,
    filter = !is.na(LBDTM),
    traceability_vars = exprs(
      LALVDOM = "LB",
      LALVSEQ = LBSEQ,
      LALVVAR = "LBDTC"
    )
  )
)

adsl_date <- date_source(
  dataset_name = "adsl",
  date = TRTEDTM,
  traceability_vars = exprs(
    LALVDOM = "ADSL",
    LALVSEQ = NA_integer_,
    LALVVAR = "TRTEDTM"
  )
)

admiral_dm %>%
  derive_var_extreme_dtm(
    new_var = LSTALVDTM,
    ae_start, ae_end, lb_date, adsl_date,
    source_datasets = list(
      adsl = admiral_adsl,
      ae = ae_ext,
      lb = lb_ext
    ),
    mode = "last"
  ) %>%
  select(USUBJID, LSTALVDTM, LALVDOM, LALVSEQ, LALVVAR)

```

```
derive_var_extreme_flag
```

Add a Variable Flagging the First or Last Observation Within Each By Group

Description

Add a variable flagging the first or last observation within each by group

Usage

```

derive_var_extreme_flag(
  dataset,
  by_vars,
  order,
  new_var,
  mode,
  check_type = "warning"
)

```

Arguments

dataset	Input dataset The variables specified by the by_vars parameter are expected.
by_vars	Grouping variables Permitted Values: list of variables
order	Sort order The first or last observation is determined with respect to the specified order. Permitted Values: list of variables or functions of variables
new_var	Variable to add The specified variable is added to the output dataset. It is set to "Y" for the first or last observation (depending on the mode) of each by group. Permitted Values: list of name-value pairs
mode	Flag mode Determines of the first or last observation is flagged. Permitted Values: "first", "last"
check_type	Check uniqueness? If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order. Default: "warning" Permitted Values: "none", "warning", "error"

Details

For each group (with respect to the variables specified for the by_vars parameter), new_var is set to "Y" for the first or last observation (with respect to the order specified for the order parameter and the flag mode specified for the mode parameter). Only observations included by the filter parameter are considered for flagging. Otherwise, new_var is set to NA. Thus, the direction of "worst" is considered fixed for all parameters in the dataset depending on the order and the mode, i.e. for every parameter the first or last record will be flagged across the whole dataset.

Value

The input dataset with the new flag variable added

See Also

[derive_var_worst_flag\(\)](#)

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```

library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data("admiral_vs")

# Flag last value for each patient, test, and visit, baseline observations are ignored
admiral_vs %>%
  restrict_derivation(
    derivation = derive_var_extreme_flag,
    args = params(
      by_vars = exprs(USUBJID, VSTESTCD, VISIT),
      order = exprs(VSTPTNUM),
      new_var = LASTFL,
      mode = "last"
    ),
    filter = VISIT != "BASELINE"
  ) %>%
  arrange(USUBJID, VSTESTCD, VISITNUM, VSTPTNUM) %>%
  select(USUBJID, VSTESTCD, VISIT, VSTPTNUM, VSSTRESN, LASTFL)

# Baseline (ABLFL) examples:

input <- tribble(
  ~STUDYID, ~USUBJID, ~PARAMCD, ~AVISIT, ~ADT, ~AVAL, ~DTYPE,
  "TEST01", "PAT01", "PARAM01", "BASELINE", as.Date("2021-04-27"), 15.0, NA,
  "TEST01", "PAT01", "PARAM01", "BASELINE", as.Date("2021-04-25"), 14.0, NA,
  "TEST01", "PAT01", "PARAM01", "BASELINE", as.Date("2021-04-23"), 15.0, "AVERAGE",
  "TEST01", "PAT01", "PARAM01", "WEEK 1", as.Date("2021-04-27"), 10.0, "AVERAGE",
  "TEST01", "PAT01", "PARAM01", "WEEK 2", as.Date("2021-04-30"), 12.0, NA,
  "TEST01", "PAT02", "PARAM01", "SCREENING", as.Date("2021-04-27"), 15.0, "AVERAGE",
  "TEST01", "PAT02", "PARAM01", "BASELINE", as.Date("2021-04-25"), 14.0, "AVERAGE",
  "TEST01", "PAT02", "PARAM01", "BASELINE", as.Date("2021-04-23"), 15.0, "AVERAGE",
  "TEST01", "PAT02", "PARAM01", "WEEK 1", as.Date("2021-04-27"), 10.0, "AVERAGE",
  "TEST01", "PAT02", "PARAM01", "WEEK 2", as.Date("2021-04-30"), 12.0, "AVERAGE",
  "TEST01", "PAT01", "PARAM02", "SCREENING", as.Date("2021-04-27"), 15.0, "AVERAGE",
  "TEST01", "PAT01", "PARAM02", "SCREENING", as.Date("2021-04-25"), 14.0, "AVERAGE",
  "TEST01", "PAT01", "PARAM02", "SCREENING", as.Date("2021-04-23"), 15.0, NA,
  "TEST01", "PAT01", "PARAM02", "BASELINE", as.Date("2021-04-27"), 10.0, "AVERAGE",
  "TEST01", "PAT01", "PARAM02", "WEEK 2", as.Date("2021-04-30"), 12.0, NA,
  "TEST01", "PAT02", "PARAM02", "SCREENING", as.Date("2021-04-27"), 15.0, NA,
  "TEST01", "PAT02", "PARAM02", "BASELINE", as.Date("2021-04-25"), 14.0, NA,
  "TEST01", "PAT02", "PARAM02", "WEEK 1", as.Date("2021-04-23"), 15.0, NA,
  "TEST01", "PAT02", "PARAM02", "WEEK 1", as.Date("2021-04-27"), 10.0, NA,
  "TEST01", "PAT02", "PARAM02", "BASELINE", as.Date("2021-04-30"), 12.0, NA
)

# Last observation
restrict_derivation(
  input,
  derivation = derive_var_extreme_flag,
  args = params(

```

```
    by_vars = exprs(USUBJID, PARAMCD),
    order = exprs(ADT),
    new_var = ABLFL,
    mode = "last"
  ),
  filter = AVISIT == "BASELINE"
)

# Worst observation - Direction = High
restrict_derivation(
  input,
  derivation = derive_var_extreme_flag,
  args = params(
    by_vars = exprs(USUBJID, PARAMCD),
    order = exprs(AVAL, ADT),
    new_var = ABLFL,
    mode = "last"
  ),
  filter = AVISIT == "BASELINE"
)

# Worst observation - Direction = Lo
restrict_derivation(
  input,
  derivation = derive_var_extreme_flag,
  args = params(
    by_vars = exprs(USUBJID, PARAMCD),
    order = exprs(desc(AVAL), ADT),
    new_var = ABLFL,
    mode = "last"
  ),
  filter = AVISIT == "BASELINE"
)

# Average observation
restrict_derivation(
  input,
  derivation = derive_var_extreme_flag,
  args = params(
    by_vars = exprs(USUBJID, PARAMCD),
    order = exprs(ADT, desc(AVAL)),
    new_var = ABLFL,
    mode = "last"
  ),
  filter = AVISIT == "BASELINE" & DTYPE == "AVERAGE"
)

# OCCURDS Examples
data("admiral_ae")

# Most severe AE first occurrence per patient
admiral_ae %>%
  mutate(
```

```

    TEMP_AESEVN =
      as.integer(factor(AESEV, levels = c("SEVERE", "MODERATE", "MILD")))
  ) %>%
  derive_var_extreme_flag(
    new_var = AOCCIFL,
    by_vars = exprs(USUBJID),
    order = exprs(TEMP_AESEVN, AESTDY, AESEQ),
    mode = "first"
  ) %>%
  arrange(USUBJID, AESTDY, AESEQ) %>%
  select(USUBJID, AEDECOD, AESEV, AESTDY, AESEQ, AOCCIFL)

# Most severe AE first occurrence per patient per body system
admiral_ae %>%
  mutate(
    TEMP_AESEVN =
      as.integer(factor(AESEV, levels = c("SEVERE", "MODERATE", "MILD")))
  ) %>%
  derive_var_extreme_flag(
    new_var = AOCCSIFL,
    by_vars = exprs(USUBJID, AEBODSYS),
    order = exprs(TEMP_AESEVN, AESTDY, AESEQ),
    mode = "first"
  ) %>%
  arrange(USUBJID, AESTDY, AESEQ) %>%
  select(USUBJID, AEBODSYS, AESEV, AESTDY, AOCCSIFL)

```

derive_var_joined_exist_flag

Derives a Flag Based on an Existing Flag

Description

Derive a flag which depends on other observations of the dataset. For example, flagging events which need to be confirmed by a second event.

Usage

```

derive_var_joined_exist_flag(
  dataset,
  by_vars,
  order,
  new_var,
  tmp_obs_nr_var = NULL,
  join_vars,
  join_type,
  first_cond = NULL,
  filter,
  true_value = "Y",

```

```

    false_value = NA_character_,
    check_type = "warning"
  )

```

Arguments

dataset	Input dataset The variables specified by the <code>by_vars</code> and <code>join_vars</code> parameter are expected.
by_vars	By variables The specified variables are used as by variables for joining the input dataset with itself.
order	Order The observations are ordered by the specified order.
new_var	New variable The specified variable is added to the input dataset.
tmp_obs_nr_var	Temporary observation number The specified variable is added to the input dataset and set to the observation number with respect to <code>order</code> . For each by group (<code>by_vars</code>) the observation number starts with 1. The variable can be used in the conditions (<code>filter</code> , <code>first_cond</code>). It is not included in the output dataset. It can be used to flag consecutive observations or the last observation (see last example below).
join_vars	Variables to keep from joined dataset The variables needed from the other observations should be specified for this parameter. The specified variables are added to the joined dataset with suffix ".join". For example to flag all observations with <code>AVALC == "Y"</code> and <code>AVALC == "Y"</code> for at least one subsequent visit <code>join_vars = exprs(AVALC, AVISITN)</code> and <code>filter = AVALC == "Y" & AVALC.join == "Y" & AVISITN < AVISITN.join</code> could be specified. The <code>*.join</code> variables are not included in the output dataset.
join_type	Observations to keep after joining The argument determines which of the joined observations are kept with respect to the original observation. For example, if <code>join_type = "after"</code> is specified all observations after the original observations are kept. For example for confirmed response or BOR in the oncology setting or confirmed deterioration in questionnaires the confirmatory assessment must be after the assessment to be flagged. Thus <code>join_type = "after"</code> could be used. Whereas, sometimes you might allow for confirmatory observations to occur prior to the observation to be flagged. For example, to flag AEs occurring on or after seven days before a COVID AE. Thus <code>join_type = "all"</code> could be used. <i>Permitted Values:</i> "before", "after", "all"
first_cond	Condition for selecting range of data If this argument is specified, the other observations are restricted up to the first observation where the specified condition is fulfilled. If the condition is not fulfilled for any of the other observations, no observations are considered, i.e., the observation is not flagged.

This parameter should be specified if `filter` contains summary functions which should not apply to all observations but only up to the confirmation assessment. For an example see the third example below.

<code>filter</code>	<p>Condition for selecting observations</p> <p>The filter is applied to the joined dataset for flagging the confirmed observations. The condition can include summary functions. The joined dataset is grouped by the original observations. I.e., the summary function are applied to all observations up to the confirmation observation. For example, <code>filter = AVALC == "CR" & all(AVALC.join %in% c("CR", "NE")) & count_vals(var = AVALC.join, val = "NE") <= 1</code> selects observations with response "CR" and for all observations up to the confirmation observation the response is "CR" or "NE" and there is at most one "NE".</p>
<code>true_value</code>	<p>Value of <code>new_var</code> for flagged observations</p> <p><i>Default:</i> "Y"</p>
<code>false_value</code>	<p>Value of <code>new_var</code> for observations not flagged</p> <p><i>Default:</i> NA_character_</p>
<code>check_type</code>	<p>Check uniqueness?</p> <p>If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order.</p> <p><i>Default:</i> "warning"</p> <p><i>Permitted Values:</i> "none", "warning", "error"</p>

Details

An example usage might be flagging if a patient received two required medications within a certain timeframe of each other.

In the oncology setting, for example, the function could be used to flag if a response value can be confirmed by an other assessment. This is commonly used in endpoints such as best overall response.

The following steps are performed to produce the output dataset.

Step 1:

The input dataset is joined with itself by the variables specified for `by_vars`. From the right hand side of the join only the variables specified for `join_vars` are kept. The suffix ".join" is added to these variables.

For example, for `by_vars = USUBJID`, `join_vars = exprs(AVISITN, AVALC)` and input dataset

```
# A tibble: 2 x 4
  USUBJID AVISITN AVALC  AVAL
<chr>    <dbl> <chr> <dbl>
1         1 Y      1
1         2 N      0
```

the joined dataset is


```
A tibble: 4 x 6
  USUBJID AVISITN AVALC  AVAL AVISITN.join AVALC.join
<chr>    <dbl> <chr> <dbl>         <dbl> <chr>
1         1 Y      1         1         1 Y
1         1 Y      1         1         2 N
1         2 N      0         0         1 Y
1         2 N      0         0         2 N
```

Step 2:

The joined dataset is restricted to observations with respect to `join_type` and order.

The dataset from the example in the previous step with `join_type = "after"` and `order = exprs(AVISITN)` is restricted to

```
A tibble: 4 x 6
  USUBJID AVISITN AVALC  AVAL AVISITN.join AVALC.join
<chr>    <dbl> <chr> <dbl>         <dbl> <chr>
1         1 Y      1         1         2 N
```

Step 3:

If `first_cond` is specified, for each observation of the input dataset the joined dataset is restricted to observations up to the first observation where `first_cond` is fulfilled (the observation fulfilling the condition is included). If for an observation of the input dataset the condition is not fulfilled, the observation is removed.

Step 4:

The joined dataset is grouped by the observations from the input dataset and restricted to the observations fulfilling the condition specified by `filter`.

Step 5:

The first observation of each group is selected

Step 6:

The variable specified by `new_var` is added to the input dataset. It is set to `true_value` for all observations which were selected in the previous step. For the other observations it is set to `false_value`.

Value

The input dataset with the variable specified by `new_var` added.

See Also

[filter_joined\(\)](#)

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```

library(tibble)
library(admiral)

# flag observations with a duration longer than 30 and
# at, after, or up to 7 days before a COVID AE (ACOVFL == "Y")
adae <- tribble(
  ~USUBJID, ~ADY, ~ACOVFL, ~ADURN,
  "1",      10, "N",      1,
  "1",      21, "N",     50,
  "1",      23, "Y",     14,
  "1",      32, "N",     31,
  "1",      42, "N",     20,
  "2",      11, "Y",     13,
  "2",      23, "N",      2,
  "3",      13, "Y",     12,
  "4",      14, "N",     32,
  "4",      21, "N",     41
)

derive_var_joined_exist_flag(
  adae,
  new_var = ALCOVFL,
  by_vars = exprs(USUBJID),
  join_vars = exprs(ACOVFL, ADY),
  join_type = "all",
  order = exprs(ADY),
  filter = ADURN > 30 & ACOVFL.join == "Y" & ADY >= ADY.join - 7
)

# flag observations with AVALC == "Y" and AVALC == "Y" at one subsequent visit
data <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      1,      "Y",
  "1",      2,      "N",
  "1",      3,      "Y",
  "1",      4,      "N",
  "2",      1,      "Y",
  "2",      2,      "N",
  "3",      1,      "Y",
  "4",      1,      "N",
  "4",      2,      "N",
)

derive_var_joined_exist_flag(
  data,
  by_vars = exprs(USUBJID),
  new_var = CONFFL,
  join_vars = exprs(AVALC, AVISITN),
  join_type = "after",
  order = exprs(AVISITN),
  filter = AVALC == "Y" & AVALC.join == "Y" & AVISITN < AVISITN.join
)

```

```

)

# select observations with AVALC == "CR", AVALC == "CR" at a subsequent visit,
# only "CR" or "NE" in between, and at most one "NE" in between
data <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      1,      "PR",
  "1",      2,      "CR",
  "1",      3,      "NE",
  "1",      4,      "CR",
  "1",      5,      "NE",
  "2",      1,      "CR",
  "2",      2,      "PR",
  "2",      3,      "CR",
  "3",      1,      "CR",
  "4",      1,      "CR",
  "4",      2,      "NE",
  "4",      3,      "NE",
  "4",      4,      "CR",
  "4",      5,      "PR"
)

derive_var_joined_exist_flag(
  data,
  by_vars = exprs(USUBJID),
  join_vars = exprs(AVALC),
  join_type = "after",
  order = exprs(AVISITN),
  new_var = CONFFL,
  first_cond = AVALC.join == "CR",
  filter = AVALC == "CR" & all(AVALC.join %in% c("CR", "NE")) &
    count_vals(var = AVALC.join, val = "NE") <= 1
)

# flag observations with AVALC == "PR", AVALC == "CR" or AVALC == "PR"
# at a subsequent visit at least 20 days later, only "CR", "PR", or "NE"
# in between, at most one "NE" in between, and "CR" is not followed by "PR"
data <- tribble(
  ~USUBJID, ~ADY, ~AVALC,
  "1",      6, "PR",
  "1",     12, "CR",
  "1",     24, "NE",
  "1",     32, "CR",
  "1",     48, "PR",
  "2",      3, "PR",
  "2",     21, "CR",
  "2",     33, "PR",
  "3",     11, "PR",
  "4",      7, "PR",
  "4",     12, "NE",
  "4",     24, "NE",
  "4",     32, "PR",
  "4",     55, "PR"
)

```

```

)

derive_var_joined_exist_flag(
  data,
  by_vars = exprs(USUBJID),
  join_vars = exprs(AVALC, ADY),
  join_type = "after",
  order = exprs(ADY),
  new_var = CONFFL,
  first_cond = AVALC.join %in% c("CR", "PR") & ADY.join - ADY >= 20,
  filter = AVALC == "PR" &
    all(AVALC.join %in% c("CR", "PR", "NE")) &
    count_vals(var = AVALC.join, val = "NE") <= 1 &
    (
      min_cond(var = ADY.join, cond = AVALC.join == "CR") >
      max_cond(var = ADY.join, cond = AVALC.join == "PR") |
      count_vals(var = AVALC.join, val = "CR") == 0
    )
)

# flag observations with CRIT1FL == "Y" at two consecutive visits or at the last visit
data <- tribble(
  ~USUBJID, ~AVISITN, ~CRIT1FL,
  "1",      1,      "Y",
  "1",      2,      "N",
  "1",      3,      "Y",
  "1",      5,      "N",
  "2",      1,      "Y",
  "2",      3,      "Y",
  "2",      5,      "N",
  "3",      1,      "Y",
  "4",      1,      "Y",
  "4",      2,      "N",
)

derive_var_joined_exist_flag(
  data,
  by_vars = exprs(USUBJID),
  new_var = CONFFL,
  tmp_obs_nr_var = tmp_obs_nr,
  join_vars = exprs(CRIT1FL),
  join_type = "all",
  order = exprs(AVISITN),
  filter = CRIT1FL == "Y" & CRIT1FL.join == "Y" &
    (tmp_obs_nr + 1 == tmp_obs_nr.join | tmp_obs_nr == max(tmp_obs_nr.join))
)

```

derive_var_last_dose_amt

Derive Last Dose Amount

Description

Add a variable for dose amount from the last dose to the input dataset.

Usage

```
derive_var_last_dose_amt(
  dataset,
  dataset_ex,
  filter_ex = NULL,
  by_vars = exprs(STUDYID, USUBJID),
  dose_id = exprs(),
  dose_date,
  analysis_date,
  single_dose_condition = (EXDOSFRQ == "ONCE"),
  new_var,
  dose_var = EXDOSE,
  traceability_vars = NULL
)
```

Arguments

dataset	Input dataset. The variables specified by the <code>by_vars</code> and <code>analysis_date</code> parameters are expected.
dataset_ex	Input EX dataset. The variables specified by the <code>by_vars</code> , <code>dose_date</code> , <code>new_vars</code> parameters, and source variables from <code>traceability_vars</code> parameter are expected.
filter_ex	Filtering condition applied to EX dataset. For example, it can be used to filter for valid dose. Defaults to NULL.
by_vars	Variables to join by (created by <code>rlang::exprs()</code>).
dose_id	Variables to identify unique dose (created by <code>rlang::exprs()</code>). Defaults to empty <code>exprs()</code> .
dose_date	The EX dose date variable. A date or date-time object is expected.
analysis_date	The analysis date variable. A date or date-time object is expected.
single_dose_condition	The condition for checking if <code>dataset_ex</code> is single dose. An error is issued if the condition is not true. Defaults to <code>(EXDOSFRQ == "ONCE")</code> .
new_var	The new variable added to dataset.
dose_var	The EX source dose amount variable. Defaults to EXDOSE.
traceability_vars	A named list returned by <code>exprs()</code> listing the traceability variables, e.g. <code>exprs(LDOSEDOM = "EX", LDOSESEQ = EXSEQ)</code> . The left-hand side (names of the list elements) gives the names of the traceability variables in the returned dataset. The right-hand side (values of the list elements) gives the values of the traceability variables in the returned dataset. These can be either strings or symbols referring to existing variables.

Details

The last dose amount is derived as the dose amount where the maximum dose_date is lower to or equal to the analysis_date per by_vars for each observation in dataset.

If dose information is aggregated (i.e. is a dosing frequency other than "ONCE" over a period defined by a start and end date) the function create_single_dose_dataset() can be used to generate single doses from aggregate dose information and satisfy single_dose_condition.

Value

Input dataset with additional column new_var.

See Also

[derive_vars_last_dose\(\)](#), [create_single_dose_dataset\(\)](#)

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data(admiral_ae)
data(ex_single)

ex_single <- derive_vars_dtm(
  head(ex_single, 100),
  dtc = EXENDTC,
  new_vars_prefix = "EXEN",
  flag_imputation = "none"
)

adae <- admiral_ae %>%
  head(100) %>%
  derive_vars_dtm(
    dtc = AESTDTC,
    new_vars_prefix = "AST",
    highest_imputation = "M"
  )

adae %>%
  derive_var_last_dose_amt(
    dataset_ex = ex_single,
    filter_ex = (EXDOSE > 0 | (EXDOSE == 0 & grepl("PLACEBO", EXTRT))) &
      !is.na(EXENDTM),
    dose_date = EXENDTM,
    analysis_date = ASTDTM,
```

```

    new_var = LDOSE,
    dose_var = EXDOSE
  ) %>%
  select(STUDYID, USUBJID, AESEQ, AESTDTC, LDOSE)

# or with traceability variables
adae %>%
  derive_var_last_dose_amt(
    dataset_ex = ex_single,
    filter_ex = (EXDOSE > 0 | (EXDOSE == 0 & grepl("PLACEBO", EXTRT))) &
      !is.na(EXENDTM),
    dose_date = EXENDTM,
    analysis_date = ASTDTM,
    new_var = LDOSE,
    dose_var = EXDOSE,
    traceability_vars = exprs(
      LDOSEDOM = "EX",
      LDOSESEQ = EXSEQ,
      LDOSEVAR = "EXDOSE"
    )
  ) %>%
  select(STUDYID, USUBJID, AESEQ, AESTDTC, LDOSEDOM, LDOSESEQ, LDOSEVAR, LDOSE)

```

 derive_var_last_dose_date

Derive Last Dose Date-Time

Description

Add a variable for the dose date or datetime of the last dose to the input dataset.

Usage

```

derive_var_last_dose_date(
  dataset,
  dataset_ex,
  filter_ex = NULL,
  by_vars = exprs(STUDYID, USUBJID),
  dose_id = exprs(),
  dose_date,
  analysis_date,
  single_dose_condition = (EXDOSFRQ == "ONCE"),
  new_var,
  output_datetime = TRUE,
  traceability_vars = NULL
)

```

Arguments

dataset	Input dataset. The variables specified by the <code>by_vars</code> and <code>analysis_date</code> parameters are expected.
dataset_ex	Input EX dataset. The variables specified by the <code>by_vars</code> , <code>dose_date</code> , <code>new_vars</code> parameters, and source variables from <code>traceability_vars</code> parameter are expected.
filter_ex	Filtering condition applied to EX dataset. For example, it can be used to filter for valid dose. Defaults to NULL.
by_vars	Variables to join by (created by <code>rlang::exprs()</code>).
dose_id	Variables to identify unique dose (created by <code>rlang::exprs()</code>). Defaults to empty <code>exprs()</code> .
dose_date	The EX dose date variable. A date or date-time object is expected.
analysis_date	The analysis date variable. A date or date-time object is expected.
single_dose_condition	The condition for checking if <code>dataset_ex</code> is single dose. An error is issued if the condition is not true. Defaults to <code>(EXDOSFRQ == "ONCE")</code> .
new_var	The new date or datetime variable added to dataset.
output_datetime	Display <code>new_var</code> as datetime or as date only. Defaults to TRUE.
traceability_vars	A named list returned by <code>exprs()</code> listing the traceability variables, e.g. <code>exprs(LDOSEDOM = "EX", LDOSESEQ = EXSEQ)</code> . The left-hand side (names of the list elements) gives the names of the traceability variables in the returned dataset. The right-hand side (values of the list elements) gives the values of the traceability variables in the returned dataset. These can be either strings or symbols referring to existing variables.

Details

The last dose date is derived as the maximum dose date where the `dose_date` is lower to or equal to the `analysis_date` per `by_vars` for each observation in `dataset`. When `output_datetime` is TRUE and time is missing, then the last dose date time is imputed to `00:00:00`. However, if date is missing, then no imputation is done.

If dose information is aggregated (i.e. is a dosing frequency other than "ONCE" over a period defined by a start and end date) the function `create_single_dose_dataset()` can be used to generate single doses from aggregate dose information and satisfy `single_dose_condition`.

Value

Input dataset with additional column `new_var`.

See Also

[derive_vars_last_dose\(\)](#), [create_single_dose_dataset\(\)](#)

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_grp\(\)](#),


```

derive_var_merged_cat(), derive_var_merged_character(), derive_var_merged_exist_flag(),
derive_var_merged_summary(), derive_var_obs_number(), derive_var_relative_flag(),
derive_vars_joined(), derive_vars_last_dose(), derive_vars_merged_lookup(), derive_vars_merged(),
derive_vars_transposed(), get_summary_records()

```

Examples

```

library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data(admiral_ae)
data(ex_single)

ex_single <- derive_vars_dtm(
  head(ex_single, 100),
  dtc = EXENDTC,
  new_vars_prefix = "EXEN",
  flag_imputation = "none"
)

adae <- admiral_ae %>%
  head(100) %>%
  derive_vars_dtm(
    dtc = AESTDTC,
    new_vars_prefix = "AST",
    highest_imputation = "M"
  )

adae %>%
  derive_var_last_dose_date(
    dataset_ex = ex_single,
    filter_ex = (EXDOSE > 0 | (EXDOSE == 0 & grepl("PLACEBO", EXTRT))) &
      !is.na(EXENDTM),
    dose_date = EXENDTM,
    analysis_date = ASTDTM,
    new_var = LDOSEDTM,
    traceability_vars = exprs(LDOSEDOM = "EX", LDOSESEQ = EXSEQ, LDOSEVAR = "EXDOSE")
  ) %>%
  select(STUDYID, USUBJID, AESEQ, AESTDTC, LDOSEDOM, LDOSESEQ, LDOSEVAR, LDOSEDTM)

```

derive_var_last_dose_grp

Derive Last Dose with User-Defined Groupings

Description

Add a variable for user-defined dose grouping of the last dose to the input dataset.

Usage

```

derive_var_last_dose_grp(
  dataset,
  dataset_ex,
  filter_ex = NULL,
  by_vars = exprs(STUDYID, USUBJID),
  dose_id = exprs(),
  dose_date,
  analysis_date,
  single_dose_condition = (EXDOSFRQ == "ONCE"),
  new_var,
  grp_brks,
  grp_lbls,
  include_lowest = TRUE,
  right = TRUE,
  dose_var = EXDOSE,
  traceability_vars = NULL
)

```

Arguments

dataset	Input dataset. The variables specified by the <code>by_vars</code> and <code>analysis_date</code> parameters are expected.
dataset_ex	Input EX dataset. The variables specified by the <code>by_vars</code> , <code>dose_date</code> , <code>new_vars</code> parameters, and source variables from <code>traceability_vars</code> parameter are expected.
filter_ex	Filtering condition applied to EX dataset. For example, it can be used to filter for valid dose. Defaults to <code>NULL</code> .
by_vars	Variables to join by (created by <code>rlang::exprs</code>).
dose_id	Variables to identify unique dose (created by <code>rlang::exprs</code>). Defaults to empty <code>exprs()</code> .
dose_date	The EX dose date variable. A date or date-time object is expected.
analysis_date	The analysis date variable. A date or date-time object is expected.
single_dose_condition	The condition for checking if <code>dataset_ex</code> is single dose. An error is issued if the condition is not true. Defaults to <code>(EXDOSFRQ == "ONCE")</code> .
new_var	The output variable defined by the user.
grp_brks	User supplied breaks to apply to groups. Refer to <code>breaks</code> parameter in <code>cut()</code> for details.
grp_lbls	User supplied labels to apply to groups. Refer to <code>labels</code> parameter in <code>cut()</code> for details.
include_lowest	logical, indicating if a value equal to the lowest (or highest, for <code>right = FALSE</code>) 'breaks' value should be included. Refer to <code>include.lowest</code> parameter in <code>cut()</code> for details.

right	Logical, indicating if the intervals should be closed on the right (and open on the left) or vice versa. Refer to right parameter in cut() for details.
dose_var	The source dose amount variable. Defaults to EXDOSE.
traceability_vars	A named list returned by <code>exprs()</code> listing the traceability variables, e.g. <code>exprs(LDOSEDOM = "EX", LDOSESEQ = EXSEQ)</code> . The left-hand side (names of the list elements) gives the names of the traceability variables in the returned dataset. The right-hand side (values of the list elements) gives the values of the traceability variables in the returned dataset. These can be either strings or symbols referring to existing variables.

Details

Last dose is the dose with maximum dose_date that is lower to or equal to the analysis_date per by_vars for each observation in dataset. The last dose group is then derived by user-defined grouping, which groups dose_var as specified in grp_brks, and returns grp_lbls as the values for new_var.

If dose information is aggregated (i.e. is a dosing frequency other than "ONCE" over a period defined by a start and end date) the function create_single_dose_dataset() can be used to generate single doses from aggregate dose information and satisfy single_dose_condition.

Value

Input dataset with additional column new_var.

See Also

[derive_vars_last_dose\(\)](#), [cut\(\)](#), [create_single_dose_dataset\(\)](#)

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data(admiral_ae)
data(ex_single)

ex_single <- derive_vars_dtm(
  head(ex_single, 100),
  dtc = EXSTDTC,
  new_vars_prefix = "EXST",
  flag_imputation = "none"
)
```

```

adae <- admiral_ae %>%
  head(100) %>%
  derive_vars_dtm(
    dtc = AESTDTC,
    new_vars_prefix = "AST",
    highest_imputation = "M"
  )

adae %>%
  derive_var_last_dose_grp(
    dataset_ex = ex_single,
    filter_ex = (EXDOSE > 0 | (EXDOSE == 0 & grepl("PLACEBO", EXTRT))) &
      !is.na(EXSTDTM),
    by_vars = exprs(STUDYID, USUBJID),
    dose_date = EXSTDTM,
    new_var = LDGRP,
    grp_brks = c(0, 20, 40, 60),
    grp_lbls = c("Low", "Medium", "High"),
    include_lowest = TRUE,
    right = TRUE,
    dose_var = EXDOSE,
    analysis_date = ASTDTM,
    traceability_vars = exprs(LDOSEDOM = "EX", LDOSESEQ = EXSEQ, LDOSEVAR = "EXENDTC")
  ) %>%
  select(USUBJID, LDGRP, LDOSEDOM, LDOSESEQ, LDOSEVAR)

```

derive_var_merged_cat *Merge a Categorization Variable*

Description

Merge a categorization variable from a dataset to the input dataset. The observations to merge can be selected by a condition and/or selecting the first or last observation for each by group.

Usage

```

derive_var_merged_cat(
  dataset,
  dataset_add,
  by_vars,
  order = NULL,
  new_var,
  source_var,
  cat_fun,
  filter_add = NULL,
  mode = NULL,
  missing_value = NA_character_
)

```

Arguments

dataset	<p>Input dataset</p> <p>The variables specified by the <code>by_vars</code> argument are expected.</p>
dataset_add	<p>Additional dataset</p> <p>The variables specified by the <code>by_vars</code>, the <code>source_var</code>, and the <code>order</code> argument are expected.</p>
by_vars	<p>Grouping variables</p> <p>The input dataset and the selected observations from the additional dataset are merged by the specified by variables. The by variables must be a unique key of the selected observations. Variables from the additional dataset can be renamed by naming the element, i.e., <code>by_vars = exprs(<name in input dataset> = <name in additional dataset>)</code> similar to the <code>dplyr</code> joins.</p> <p><i>Permitted Values:</i> list of variables created by <code>exprs()</code></p>
order	<p>Sort order</p> <p>If the argument is set to a non-null value, for each by group the first or last observation from the additional dataset is selected with respect to the specified order.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code>, e.g., <code>exprs(ADT, desc(AVAL))</code> or NULL</p>
new_var	<p>New variable</p> <p>The specified variable is added to the additional dataset and set to the categorized values, i.e., <code>cat_fun(<source variable>)</code>.</p>
source_var	Source variable
cat_fun	<p>Categorization function</p> <p>A function must be specified for this argument which expects the values of the source variable as input and returns the categorized values.</p>
filter_add	<p>Filter for additional dataset (<code>dataset_add</code>)</p> <p>Only observations fulfilling the specified condition are taken into account for merging. If the argument is not specified, all observations are considered.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> a condition</p>
mode	<p>Selection mode</p> <p>Determines if the first or last observation is selected. If the <code>order</code> argument is specified, <code>mode</code> must be non-null.</p> <p>If the <code>order</code> argument is not specified, the <code>mode</code> argument is ignored.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> "first", "last", NULL</p>
missing_value	<p>Values used for missing information</p> <p>The new variable is set to the specified value for all by groups without observations in the additional dataset.</p> <p><i>Default:</i> <code>NA_character_</code></p>

Details

1. The additional dataset is restricted to the observations matching the `filter_add` condition.
2. The categorization variable is added to the additional dataset.
3. If `order` is specified, for each by group the first or last observation (depending on mode) is selected.
4. The categorization variable is merged to the input dataset.

Value

The output dataset contains all observations and variables of the input dataset and additionally the variable specified for `new_var` derived from the additional dataset (`dataset_add`).

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(admiral.test)
library(dplyr, warn.conflicts = FALSE)
data("admiral_dm")
data("admiral_vs")

wgt_cat <- function(wgt) {
  case_when(
    wgt < 50 ~ "low",
    wgt > 90 ~ "high",
    TRUE ~ "normal"
  )
}

derive_var_merged_cat(
  admiral_dm,
  dataset_add = admiral_vs,
  by_vars = exprs(STUDYID, USUBJID),
  order = exprs(VSDTC, VSSEQ),
  filter_add = VSTESTCD == "WEIGHT" & substr(VISIT, 1, 9) == "SCREENING",
  new_var = WGTBLCAT,
  source_var = VSSTRESN,
  cat_fun = wgt_cat,
  mode = "last"
) %>%
  select(STUDYID, USUBJID, AGE, AGEU, WGTBLCAT)

# defining a value for missing VS data
```

```

derive_var_merged_cat(
  admiral_dm,
  dataset_add = admiral_vs,
  by_vars = exprs(STUDYID, USUBJID),
  order = exprs(VSDTC, VSSEQ),
  filter_add = VSTESTCD == "WEIGHT" & substr(VISIT, 1, 9) == "SCREENING",
  new_var = WGTBLCAT,
  source_var = VSSTRESN,
  cat_fun = wgt_cat,
  mode = "last",
  missing_value = "MISSING"
) %>%
  select(STUDYID, USUBJID, AGE, AGEU, WGTBLCAT)

```

derive_var_merged_character

Merge a Character Variable

Description

Merge a character variable from a dataset to the input dataset. The observations to merge can be selected by a condition and/or selecting the first or last observation for each by group.

Usage

```

derive_var_merged_character(
  dataset,
  dataset_add,
  by_vars,
  order = NULL,
  new_var,
  source_var,
  case = NULL,
  filter_add = NULL,
  mode = NULL,
  missing_value = NA_character_
)

```

Arguments

dataset	Input dataset The variables specified by the by_vars argument are expected.
dataset_add	Additional dataset The variables specified by the by_vars, the source_var, and the order argument are expected.

by_vars	<p>Grouping variables</p> <p>The input dataset and the selected observations from the additional dataset are merged by the specified by variables. The by variables must be a unique key of the selected observations. Variables from the additional dataset can be renamed by naming the element, i.e., <code>by_vars = exprs(<name in input dataset> = <name in additional dataset>)</code> similar to the dplyr joins.</p> <p><i>Permitted Values:</i> list of variables created by <code>exprs()</code></p>
order	<p>Sort order</p> <p>If the argument is set to a non-null value, for each by group the first or last observation from the additional dataset is selected with respect to the specified order.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code>, e.g., <code>exprs(ADT, desc(AVAL))</code> or NULL</p>
new_var	<p>New variable</p> <p>The specified variable is added to the additional dataset and set to the transformed value with respect to the case argument.</p>
source_var	Source variable
case	<p>Change case</p> <p>Changes the case of the values of the new variable.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> NULL, "lower", "upper", "title"</p>
filter_add	<p>Filter for additional dataset (<code>dataset_add</code>)</p> <p>Only observations fulfilling the specified condition are taken into account for merging. If the argument is not specified, all observations are considered.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> a condition</p>
mode	<p>Selection mode</p> <p>Determines if the first or last observation is selected. If the order argument is specified, mode must be non-null.</p> <p>If the order argument is not specified, the mode argument is ignored.</p> <p><i>Default:</i> NULL</p> <p><i>Permitted Values:</i> "first", "last", NULL</p>
missing_value	<p>Values used for missing information</p> <p>The new variable is set to the specified value for all by groups without observations in the additional dataset.</p> <p><i>Default:</i> <code>NA_character_</code></p> <p><i>Permitted Value:</i> A character scalar</p>

Details

1. The additional dataset is restricted to the observations matching the `filter_add` condition.
2. The (transformed) character variable is added to the additional dataset.

3. If order is specified, for each by group the first or last observation (depending on mode) is selected.
4. The character variable is merged to the input dataset.

Value

The output dataset contains all observations and variables of the input dataset and additionally the variable specified for new_var derived from the additional dataset (dataset_add).

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(admiral.test)
library(dplyr, warn.conflicts = FALSE)
data("admiral_dm")
data("admiral_ds")

derive_var_merged_character(
  admiral_dm,
  dataset_add = admiral_ds,
  by_vars = exprs(STUDYID, USUBJID),
  new_var = DISPSTAT,
  filter_add = DSCAT == "DISPOSITION EVENT",
  source_var = DSDECOD,
  case = "title"
) %>%
  select(STUDYID, USUBJID, AGE, AGEU, DISPSTAT)
```

derive_var_merged_exist_flag

Merge an Existence Flag

Description

Adds a flag variable to the input dataset which indicates if there exists at least one observation in another dataset fulfilling a certain condition.

Usage

```

derive_var_merged_exist_flag(
  dataset,
  dataset_add,
  by_vars,
  new_var,
  condition,
  true_value = "Y",
  false_value = NA_character_,
  missing_value = NA_character_,
  filter_add = NULL
)

```

Arguments

dataset	Input dataset The variables specified by the <code>by_vars</code> argument are expected.
dataset_add	Additional dataset The variables specified by the <code>by_vars</code> argument are expected.
by_vars	Grouping variables <i>Permitted Values:</i> list of variables
new_var	New variable The specified variable is added to the input dataset.
condition	Condition The condition is evaluated at the additional dataset (<code>dataset_add</code>). For all by groups where it evaluates as TRUE at least once the new variable is set to the true value (<code>true_value</code>). For all by groups where it evaluates as FALSE or NA for all observations the new variable is set to the false value (<code>false_value</code>). The new variable is set to the missing value (<code>missing_value</code>) for by groups not present in the additional dataset.
true_value	True value <i>Default:</i> "Y"
false_value	False value <i>Default:</i> NA_character_
missing_value	Values used for missing information The new variable is set to the specified value for all by groups without observations in the additional dataset. <i>Default:</i> NA_character_ <i>Permitted Value:</i> A character scalar
filter_add	Filter for additional data Only observations fulfilling the specified condition are taken into account for flagging. If the argument is not specified, all observations are considered. <i>Permitted Values:</i> a condition

Details

1. The additional dataset is restricted to the observations matching the `filter_add` condition.
2. The new variable is added to the input dataset and set to the true value (`true_value`) if for the `by` group at least one observation exists in the (restricted) additional dataset where the condition evaluates to TRUE. It is set to the false value (`false_value`) if for the `by` group at least one observation exists and for all observations the condition evaluates to FALSE or NA. Otherwise, it is set to the missing value (`missing_value`).

Value

The output dataset contains all observations and variables of the input dataset and additionally the variable specified for `new_var` derived from the additional dataset (`dataset_add`).

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(admiral.test)
library(dplyr, warn.conflicts = FALSE)
data("admiral_dm")
data("admiral_ae")
derive_var_merged_exist_flag(
  admiral_dm,
  dataset_add = admiral_ae,
  by_vars = exprs(STUDYID, USUBJID),
  new_var = AERELFL,
  condition = AEREL == "PROBABLE"
) %>%
  select(STUDYID, USUBJID, AGE, AGEU, AERELFL)

data("admiral_vs")
derive_var_merged_exist_flag(
  admiral_dm,
  dataset_add = admiral_vs,
  by_vars = exprs(STUDYID, USUBJID),
  filter_add = VSTESTCD == "WEIGHT" & VSBLFL == "Y",
  new_var = WTBLHIFL,
  condition = VSSTRESN > 90,
  false_value = "N",
  missing_value = "M"
) %>%
  select(STUDYID, USUBJID, AGE, AGEU, WTBLHIFL)
```

 derive_var_merged_summary

Merge a Summary Variable

Description

Merge a summary variable from a dataset to the input dataset.

Usage

```
derive_var_merged_summary(
  dataset,
  dataset_add,
  by_vars,
  new_var,
  filter_add = NULL,
  analysis_var,
  summary_fun
)
```

Arguments

dataset	Input dataset The variables specified by the <code>by_vars</code> argument are expected.
dataset_add	Additional dataset The variables specified by the <code>by_vars</code> and the <code>analysis_var</code> arguments are expected.
by_vars	Grouping variables The values of <code>analysis_var</code> are summarized by the specified variables. The summarized values are merged to the input dataset (<code>dataset</code>) by the specified by variables. <i>Permitted Values:</i> list of variables created by <code>exprs()</code>
new_var	Variable to add The specified variable is added to the input dataset (<code>dataset</code>) and set to the summarized values.
filter_add	Filter for additional dataset (<code>dataset_add</code>) Only observations fulfilling the specified condition are taken into account for summarizing. If the argument is not specified, all observations are considered. <i>Permitted Values:</i> a condition
analysis_var	Analysis variable The values of the specified variable are summarized by the function specified for <code>summary_fun</code> .
summary_fun	Summary function The specified function that takes as input <code>analysis_var</code> and performs the calculation. This can include built-in functions as well as user defined functions, for example <code>mean</code> or <code>function(x) mean(x, na.rm = TRUE)</code> .

Details

1. The records from the additional dataset (`dataset_add`) are restricted to those matching the `filter_add` condition.
2. The values of the analysis variable (`analysis_var`) are summarized by the summary function (`summary_fun`) for each by group (`by_vars`) in the additional dataset (`dataset_add`).
3. The summarized values are merged to the input dataset as a new variable (`new_var`). For observations without a matching observation in the additional dataset the new variable is set to NA. Observations in the additional dataset which have no matching observation in the input dataset are ignored.

Value

The output dataset contains all observations and variables of the input dataset and additionally the variable specified for `new_var`.

See Also

[derive_summary_records\(\)](#), [get_summary_records\(\)](#)

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(tibble)

# Add a variable for the mean of AVAL within each visit
adbds <- tribble(
  ~USUBJID, ~AVISIT, ~ASEQ, ~AVAL,
  "1",      "WEEK 1",  1,    10,
  "1",      "WEEK 1",  2,    NA,
  "1",      "WEEK 2",  3,    NA,
  "1",      "WEEK 3",  4,    42,
  "1",      "WEEK 4",  5,    12,
  "1",      "WEEK 4",  6,    12,
  "1",      "WEEK 4",  7,    15,
  "2",      "WEEK 1",  1,    21,
  "2",      "WEEK 4",  2,    22
)

derive_var_merged_summary(
  adbds,
  dataset_add = adbds,
  by_vars = exprs(USUBJID, AVISIT),
  new_var = MEANVIS,
  analysis_var = AVAL,
  summary_fun = function(x) mean(x, na.rm = TRUE)
```

```

)

# Add a variable listing the lesion ids at baseline
adsl <- tribble(
  ~USUBJID,
  "1",
  "2",
  "3"
)

adtr <- tribble(
  ~USUBJID, ~AVISIT, ~LESIONID,
  "1", "BASELINE", "INV-T1",
  "1", "BASELINE", "INV-T2",
  "1", "BASELINE", "INV-T3",
  "1", "BASELINE", "INV-T4",
  "1", "WEEK 1", "INV-T1",
  "1", "WEEK 1", "INV-T2",
  "1", "WEEK 1", "INV-T4",
  "2", "BASELINE", "INV-T1",
  "2", "BASELINE", "INV-T2",
  "2", "BASELINE", "INV-T3",
  "2", "WEEK 1", "INV-T1",
  "2", "WEEK 1", "INV-N1"
)

derive_var_merged_summary(
  adsl,
  dataset_add = adtr,
  by_vars = exprs(USUBJID),
  filter_add = AVISIT == "BASELINE",
  new_var = LESIONSBL,
  analysis_var = LESIONID,
  summary_fun = function(x) paste(x, collapse = ", ")
)

```

derive_var_obs_number *Adds a Variable Numbering the Observations Within Each By Group*

Description

Adds a variable numbering the observations within each by group

Usage

```

derive_var_obs_number(
  dataset,
  by_vars = NULL,
  order = NULL,

```

```

    new_var = ASEQ,
    check_type = "none"
)

```

Arguments

dataset	Input dataset The variables specified by the order and the by_vars parameter are expected.
by_vars	Grouping variables Permitted Values: list of variables
order	Sort order Within each by group the observations are ordered by the specified order. Permitted Values: list of variables or functions of variables
new_var	Name of variable to create The new variable is set to the observation number for each by group. The numbering starts with 1. Default: ASEQ
check_type	Check uniqueness? If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order. Default: "none" Permitted Values: "none", "warning", "error"

Details

For each group (with respect to the variables specified for the by_vars parameter) the first or last observation (with respect to the order specified for the order parameter and the mode specified for the mode parameter) is included in the output dataset.

Value

A dataset containing all observations and variables of the input dataset and additionally the variable specified by the new_var parameter.

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```

library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data("admiral_vs")

admiral_vs %>%
  select(USUBJID, VSTESTCD, VISITNUM, VSTPTNUM) %>%
  derive_var_obs_number(
    by_vars = exprs(USUBJID, VSTESTCD),
    order = exprs(VISITNUM, desc(VSTPTNUM))
  )

```

derive_var_ontrtfl *Derive On-Treatment Flag Variable*

Description

Derive on-treatment flag (ONTRTFL) in an ADaM dataset with a single assessment date (e.g ADT) or event start and end dates (e.g. ASTDT/AENDT).

Usage

```

derive_var_ontrtfl(
  dataset,
  new_var = ONTRTFL,
  start_date,
  end_date = NULL,
  ref_start_date,
  ref_end_date = NULL,
  ref_end_window = 0,
  ignore_time_for_ref_end_date = TRUE,
  filter_pre_timepoint = NULL,
  span_period = NULL
)

```

Arguments

dataset	Input dataset. Required columns are start_date, end_date, ref_start_date and ref_end_date.
new_var	On-treatment flag variable name to be created. Default is ONTRTFL.
start_date	The start date (e.g. AESDT) or assessment date (e.g. ADT) Required; A date or date-time object column is expected. Refer to derive_vars_dt() to impute and derive a date from a date character vector to a date object.

end_date	<p>The end date of assessment/event (e.g. AENDT) A date or date-time object column is expected.</p> <p>Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object.</p> <p>Optional; Default is null. If the used and date value is missing on an observation, it is assumed the medication is ongoing and ONTRTFL is set to "Y".</p>
ref_start_date	<p>The lower bound of the on-treatment period Required; A date or date-time object column is expected.</p> <p>Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object.</p>
ref_end_date	<p>The upper bound of the on-treatment period A date or date-time object column is expected.</p> <p>Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object.</p> <p>Optional; This can be null and everything after <code>ref_start_date</code> will be considered on-treatment. Default is NULL.</p>
ref_end_window	<p>A window to add to the upper bound <code>ref_end_date</code> measured in days (e.g. 7 if 7 days should be added to the upper bound) Optional; default is 0.</p>
ignore_time_for_ref_end_date	<p>If the argument is set to TRUE, the time part is ignored for checking if the event occurred more than <code>ref_end_window</code> days after reference end date.</p> <p><i>Permitted Values:</i> TRUE, FALSE</p>
filter_pre_timepoint	<p>An expression to filter observations as not on-treatment when <code>date = ref_start_date</code>. For example, if observations where <code>VSTPT = PRE</code> should not be considered on-treatment when <code>date = ref_start_date</code>, <code>filter_pre_timepoint</code> should be used to denote when the on-treatment flag should be set to null. Optional; default is NULL.</p>
span_period	<p>A "Y" scalar character. If "Y", events that started prior to the <code>ref_start_date</code> and are ongoing or end after the <code>ref_start_date</code> are flagged as "Y". Optional; default is NULL.</p>

Details

On-Treatment is calculated by determining whether the assessment date or start/stop dates fall between 2 dates. The following logic is used to assign on-treatment = "Y":

1. `start_date` is missing and `ref_start_date` is non-missing
2. No timepoint filter is provided (`filter_pre_timepoint`) and both `start_date` and `ref_start_date` are non-missing and `start_date = ref_start_date`
3. A timepoint is provided (`filter_pre_timepoint`) and both `start_date` and `ref_start_date` are non-missing and `start_date = ref_start_date` and the filter provided in `filter_pre_timepoint` is not true.
4. `ref_end_date` is not provided and `ref_start_date < start_date`
5. `ref_end_date` is provided and `ref_start_date < start_date <= ref_end_date + ref_end_window`.

If the `end_date` is provided and the `end_date < ref_start_date` then the ONTRTFL is set to NULL. This would be applicable to cases where the `start_date` is missing and ONTRTFL has been assigned as "Y" above.

If the `span_period` is specified as "Y", this allows the user to assign ONTRTFL as "Y" to cases where the record started prior to the `ref_start_date` and was ongoing or ended after the `ref_start_date`.

Any date imputations needed should be done prior to calling this function.

Value

The input dataset with an additional column named ONTRTFL with a value of "Y" or NA

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_anrind\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_basetype\(\)](#), [derive_var_base\(\)](#), [derive_var_chg\(\)](#), [derive_var_pchg\(\)](#), [derive_var_shift\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate, warn.conflicts = FALSE)

advs <- tribble(
  ~USUBJID, ~ADT,          ~TRTSDT,          ~TRTEDT,
  "P01",    ymd("2020-02-24"), ymd("2020-01-01"), ymd("2020-03-01"),
  "P02",    ymd("2020-01-01"), ymd("2020-01-01"), ymd("2020-03-01"),
  "P03",    ymd("2019-12-31"), ymd("2020-01-01"), ymd("2020-03-01")
)
derive_var_ontrtfl(
  advs,
  start_date = ADT,
  ref_start_date = TRTSDT,
  ref_end_date = TRTEDT
)

advs <- tribble(
  ~USUBJID, ~ADT,          ~TRTSDT,          ~TRTEDT,
  "P01",    ymd("2020-07-01"), ymd("2020-01-01"), ymd("2020-03-01"),
  "P02",    ymd("2020-04-30"), ymd("2020-01-01"), ymd("2020-03-01"),
  "P03",    ymd("2020-03-15"), ymd("2020-01-01"), ymd("2020-03-01")
)
derive_var_ontrtfl(
  advs,
  start_date = ADT,
  ref_start_date = TRTSDT,
  ref_end_date = TRTEDT,
  ref_end_window = 60
)

advs <- tribble(
```

```

~USUBJID, ~ADTM, ~TRTSDTM, ~TRTEDTM,
"P01", ymd_hm("2020-01-02T12:00"), ymd_hm("2020-01-01T12:00"), ymd_hm("2020-03-01T12:00"),
"P02", ymd("2020-01-01"), ymd_hm("2020-01-01T12:00"), ymd_hm("2020-03-01T12:00"),
"P03", ymd("2019-12-31"), ymd_hm("2020-01-01T12:00"), ymd_hm("2020-03-01T12:00"),
) %>%
mutate(TPT = c(NA, "PRE", NA))
derive_var_ontrtfl(
  advs,
  start_date = ADTM,
  ref_start_date = TRTSDTM,
  ref_end_date = TRTEDTM,
  filter_pre_timepoint = TPT == "PRE"
)

advs <- tribble(
  ~USUBJID, ~ASTDT, ~TRTSDT, ~TRTEDT, ~AENDT,
  "P01", ymd("2020-03-15"), ymd("2020-01-01"), ymd("2020-03-01"), ymd("2020-12-01"),
  "P02", ymd("2019-04-30"), ymd("2020-01-01"), ymd("2020-03-01"), ymd("2020-03-15"),
  "P03", ymd("2019-04-30"), ymd("2020-01-01"), ymd("2020-03-01"), NA,
)
derive_var_ontrtfl(
  advs,
  start_date = ASTDT,
  end_date = AENDT,
  ref_start_date = TRTSDT,
  ref_end_date = TRTEDT,
  ref_end_window = 60,
  span_period = "Y"
)

advs <- tribble(
  ~USUBJID, ~ASTDT, ~AP01SDT, ~AP01EDT, ~AENDT,
  "P01", ymd("2020-03-15"), ymd("2020-01-01"), ymd("2020-03-01"), ymd("2020-12-01"),
  "P02", ymd("2019-04-30"), ymd("2020-01-01"), ymd("2020-03-01"), ymd("2020-03-15"),
  "P03", ymd("2019-04-30"), ymd("2020-01-01"), ymd("2020-03-01"), NA,
)
derive_var_ontrtfl(
  advs,
  new_var = ONTR01FL,
  start_date = ASTDT,
  end_date = AENDT,
  ref_start_date = AP01SDT,
  ref_end_date = AP01EDT,
  span_period = "Y"
)

```

derive_var_pchg

Derive Percent Change from Baseline

Description

Derive percent change from baseline (PCHG) in a BDS dataset

Usage

```
derive_var_pchg(dataset)
```

Arguments

dataset The input dataset. Required variables are AVAL and BASE.

Details

Percent change from baseline is calculated by dividing change from baseline by the absolute value of the baseline value and multiplying the result by 100.

Value

The input dataset with an additional column named PCHG

See Also

[derive_var_chg\(\)](#)

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_anrind\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_basetype\(\)](#), [derive_var_base\(\)](#), [derive_var_chg\(\)](#), [derive_var_ontrtfl\(\)](#), [derive_var_shift\(\)](#)

Examples

```
library(tibble)

advs <- tribble(
  ~USUBJID, ~PARAMCD, ~AVAL, ~ABLFL, ~BASE,
  "P01",    "WEIGHT", 80,    "Y",    80,
  "P01",    "WEIGHT", 80.8,  "",    80,
  "P01",    "WEIGHT", 81.4,  "",    80,
  "P02",    "WEIGHT", 75.3,  "Y",   75.3,
  "P02",    "WEIGHT", 76,    "",    75.3
)
derive_var_pchg(advs)
```

```
derive_var_relative_flag
```

Flag Observations Before or After a Condition is Fulfilled

Description

Flag all observations before or after the observation where a specified condition is fulfilled for each by group. For example, the function could be called to flag for each subject all observations before the first disease progression or to flag all AEs after a specific AE.

Usage

```

derive_var_relative_flag(
  dataset,
  by_vars,
  order,
  new_var,
  condition,
  mode,
  selection,
  inclusive,
  flag_no_ref_groups = TRUE,
  check_type = "warning"
)

```

Arguments

dataset	Input dataset The variables specified by the order and the by_vars argument are expected.
by_vars	Grouping variables <i>Permitted Values:</i> list of variables created by <code>exprs()</code>
order	Sort order Within each by group the observations are ordered by the specified order. <i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code> , e.g., <code>exprs(ADT, desc(AVAL))</code>
new_var	New variable The variable is added to the input dataset and set to "Y" for all observations before or after the condition is fulfilled. For all other observations it is set to NA.
condition	Condition for Reference Observation The specified condition determines the reference observation. In the output dataset all observations before or after (selection argument) the reference observation are flagged.
mode	Selection mode (first or last) If "first" is specified, for each by group the observations before or after (selection argument) the observation where the condition (condition argument) is fulfilled the <i>first</i> time is flagged in the output dataset. If "last" is specified, for each by group the observations before or after (selection argument) the observation where the condition (condition argument) is fulfilled the <i>last</i> time is flagged in the output dataset. <i>Permitted Values:</i> "first", "last"
selection	Flag observations before or after the reference observation? <i>Permitted Values:</i> "before", "after"
inclusive	Flag the reference observation? <i>Permitted Values:</i> TRUE, FALSE

flag_no_ref_groups	Should by groups without reference observation be flagged? <i>Permitted Values:</i> TRUE, FALSE
check_type	Check uniqueness? If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order. <i>Permitted Values:</i> "none", "warning", "error"

Details

For each by group (by_vars argument) the observations before or after (selection argument) the observations where the condition (condition argument) is fulfilled the first or last time (order argument and mode argument) is flagged in the output dataset.

Value

The input dataset with the new variable (new_var) added

See Also

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#), [get_summary_records\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)

# Flag all AEs after the first COVID AE
adae <- tribble(
  ~USUBJID, ~ASTDY, ~ACOVFL, ~AESEQ,
  "1",      2, NA,      1,
  "1",      5, "Y",     2,
  "1",      5, NA,      3,
  "1",     17, NA,      4,
  "1",     27, "Y",     5,
  "1",     32, NA,      6,
  "2",      8, NA,      1,
  "2",     11, NA,      2,
)

derive_var_relative_flag(
  adae,
  by_vars = exprs(USUBJID),
  order = exprs(ASTDY, AESEQ),
  new_var = PSTCOVFL,
```

```

    condition = ACOVFL == "Y",
    mode = "first",
    selection = "after",
    inclusive = FALSE,
    flag_no_ref_groups = FALSE
  )

response <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      0,        "PR",
  "1",      1,        "CR",
  "1",      2,        "CR",
  "1",      3,        "SD",
  "1",      4,        "NE",
  "2",      0,        "SD",
  "2",      1,        "PD",
  "2",      2,        "PD",
  "3",      0,        "SD",
  "4",      0,        "SD",
  "4",      1,        "PR",
  "4",      2,        "PD",
  "4",      3,        "SD",
  "4",      4,        "PR"
)

# Flag observations up to first PD for each patient
response %>%
  derive_var_relative_flag(
    by_vars = exprs(USUBJID),
    order = exprs(AVISITN),
    new_var = ANL02FL,
    condition = AVALC == "PD",
    mode = "first",
    selection = "before",
    inclusive = TRUE
  )

# Flag observations up to first PD excluding baseline (AVISITN = 0) for each patient
response %>%
  restrict_derivation(
    derivation = derive_var_relative_flag,
    args = params(
      by_vars = exprs(USUBJID),
      order = exprs(AVISITN),
      new_var = ANL02FL,
      condition = AVALC == "PD",
      mode = "first",
      selection = "before",
      inclusive = TRUE
    ),
    filter = AVISITN > 0
  ) %>%
  arrange(USUBJID, AVISITN)

```

derive_var_shift *Derive Shift*

Description

Derives a character shift variable containing concatenated shift in values based on user-defined pairing, e.g., shift from baseline to analysis value, shift from baseline grade to analysis grade, ...

Usage

```
derive_var_shift(
  dataset,
  new_var,
  from_var,
  to_var,
  na_val = "NULL",
  sep_val = " to "
)
```

Arguments

dataset	Input dataset The columns specified by from_var and the to_var parameters are expected.
new_var	Name of the character shift variable to create.
from_var	Variable containing value to shift from.
to_var	Variable containing value to shift to.
na_val	Character string to replace missing values in from_var or to_var. Default: "NULL"
sep_val	Character string to concatenate values of from_var and to_var. Default: " to "

Details

new_var is derived by concatenating the values of from_var to values of to_var (e.g. "NORMAL to HIGH"). When from_var or to_var has missing value, the missing value is replaced by na_val (e.g. "NORMAL to NULL").

Value

The input dataset with the character shift variable added

See Also

BDS-Findings Functions that returns variable appended to dataset: [derive_var_analysis_ratio\(\)](#), [derive_var_anrind\(\)](#), [derive_var_atoxgr_dir\(\)](#), [derive_var_atoxgr\(\)](#), [derive_var_basetype\(\)](#), [derive_var_base\(\)](#), [derive_var_chg\(\)](#), [derive_var_ontrtfl\(\)](#), [derive_var_pchg\(\)](#)

Examples

```

library(tibble)

data <- tribble(
  ~USUBJID, ~PARAMCD, ~AVAL, ~ABLFL, ~BNRIND, ~ANRIND,
  "P01", "ALB", 33, "Y", "LOW", "LOW",
  "P01", "ALB", 38, NA, "LOW", "NORMAL",
  "P01", "ALB", NA, NA, "LOW", NA,
  "P02", "ALB", 37, "Y", "NORMAL", "NORMAL",
  "P02", "ALB", 49, NA, "NORMAL", "HIGH",
  "P02", "SODIUM", 147, "Y", "HIGH", "HIGH"
)

data %>%
  convert_blanks_to_na() %>%
  derive_var_shift(
    new_var = SHIFT1,
    from_var = BNRIND,
    to_var = ANRIND
  )

# or only populate post-baseline records
data %>%
  convert_blanks_to_na() %>%
  restrict_derivation(
    derivation = derive_var_shift,
    args = params(
      new_var = SHIFT1,
      from_var = BNRIND,
      to_var = ANRIND
    ),
    filter = is.na(ABLFL)
  )

```

derive_var_trtdurd *Derive Total Treatment Duration (Days)*

Description

Derives total treatment duration (days) (TRTDURD)

Usage

```
derive_var_trtdurd(dataset, start_date = TRTSDT, end_date = TRTEDT)
```

Arguments

dataset	Input dataset
---------	---------------

The columns specified by the start_date and the end_date parameter are expected.

start_date	The start date A date or date-time object is expected. Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object. Default: TRTSDT
end_date	The end date A date or date-time object is expected. Refer to <code>derive_vars_dt()</code> to impute and derive a date from a date character vector to a date object. Default: TRTEDT

Details

The total treatment duration is derived as the number of days from start to end date plus one.

Value

The input dataset with TRTDURD added

See Also

[derive_vars_duration\(\)](#)

Date/Time Derivation Functions that returns variable appended to dataset: [derive_vars_dtm_to_dt\(\)](#), [derive_vars_dtm_to_tm\(\)](#), [derive_vars_dtm\(\)](#), [derive_vars_dt\(\)](#), [derive_vars_duration\(\)](#), [derive_vars_dy\(\)](#)

Examples

```
library(tibble)
library(lubridate)

data <- tribble(
  ~TRTSDT, ~TRTEDT,
  ymd("2020-01-01"), ymd("2020-02-24")
)

derive_var_trtdurd(data)
```

derive_var_trtemfl *Derive Treatment-emergent Flag*

Description

Derive treatment emergent analysis flag (e.g., TRTEMFL).

Usage

```

derive_var_trtemfl(
  dataset,
  new_var = TRTEMFL,
  start_date = ASTDTM,
  end_date = AENDTM,
  trt_start_date = TRTSDTM,
  trt_end_date = NULL,
  end_window = NULL,
  ignore_time_for_trt_end = TRUE,
  initial_intensity = NULL,
  intensity = NULL
)

```

Arguments

dataset	Input dataset The variables specified by start_date, end_date, trt_start_date, trt_end_date, initial_intensity, and intensity are expected.
new_var	New variable
start_date	Event start date <i>Permitted Values:</i> A symbol referring to a date or datetime variable of the input dataset
end_date	Event end date <i>Permitted Values:</i> A symbol referring to a date or datetime variable of the input dataset
trt_start_date	Treatment start date <i>Permitted Values:</i> A symbol referring to a date or datetime variable of the input dataset
trt_end_date	Treatment end date <i>Permitted Values:</i> A symbol referring to a date or datetime variable of the input dataset or NULL
end_window	If the argument is specified, events starting more than the specified number of days after end of treatment, are not flagged. <i>Permitted Values:</i> A non-negative integer or NULL
ignore_time_for_trt_end	If the argument is set to TRUE, the time part is ignored for checking if the event occurred more than end_window days after end of treatment. <i>Permitted Values:</i> TRUE, FALSE
initial_intensity	Initial severity/intensity or toxicity This derivation assumes AE data collection method as single record per AE with “initial” and “most extreme” severity/intensity recorded separately.

intensity	<p>If the argument is specified, events which start before treatment start and end after treatment start (or are ongoing) and worsened (i.e., the intensity is greater than the initial intensity), are flagged.</p> <p>The values of the specified variable must be comparable with the usual comparison operators. I.e., if the intensity is greater than the initial intensity <code>initial_intensity < intensity</code> must evaluate to TRUE.</p> <p><i>Permitted Values:</i> A symbol referring to a variable of the input dataset or NULL</p>
	<p>Severity/intensity or toxicity</p> <p>If the argument is specified, events which start before treatment start and end after treatment start (or are ongoing) and worsened (i.e., the intensity is greater than the initial intensity), are flagged.</p> <p>The values of the specified variable must be comparable with the usual comparison operators. I.e., if the intensity is greater than the initial intensity <code>initial_intensity < intensity</code> must evaluate to TRUE.</p> <p><i>Permitted Values:</i> A symbol referring to a variable of the input dataset or NULL</p>

Details

For the derivation of the new variable the following cases are considered in this order. The first case which applies, defines the value of the variable.

- *not treated:* If `trt_start_date` is NA, it is set to `NA_character_`.
- *event before treatment:* If `end_date` is before `trt_start_date` (and `end_date` is not NA), it is set to `NA_character_`.
- *no event date:* If `start_date` is NA, it is set to "Y" as in such cases it is usually considered more conservative to assume the event was treatment-emergent.
- *event started during treatment:*
 - if `end_window` is not specified: if `start_date` is on or after `trt_start_date`, it is set to "Y",
 - if `end_window` is specified: if `start_date` is on or after `trt_start_date` and `start_date` is on or before `trt_end_date + end_window days`, it is set to "Y",
- *event started before treatment and (possibly) worsened on treatment:*
 - if `initial_intensity` and `intensity` is specified: if `initial_intensity < intensity` and `start_date` is before `trt_start_date` and `end_date` is on or after `trt_start_date` or `end_date` is NA, it is set to "Y".
- Otherwise it is set to `NA_character_`.

Value

The input dataset with the variable specified by `new_var` added

See Also

OCCDS Functions: [derive_vars_atc\(\)](#), [derive_vars_query\(\)](#), [get_terms_from_db\(\)](#)

Examples

```

library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

adae <- expected <- tribble(
  ~USUBJID, ~ASTDTM,          ~AENDTM,          ~AEITOXGR, ~AETOXGR,
  # before treatment
  "1",      "2021-12-13T20:15", "2021-12-15T12:45", "1",      "1",
  "1",      "2021-12-14T20:15", "2021-12-14T22:00", "1",      "3",
  # starting before treatment and ending during treatment
  "1",      "2021-12-30T20:00", "2022-01-14T11:00", "1",      "3",
  "1",      "2021-12-31T20:15", "2022-01-01T01:23", "1",      "1",
  # starting during treatment
  "1",      "2022-01-01T12:00", "2022-01-02T23:25", "3",      "4",
  # after treatment
  "1",      "2022-05-10T11:00", "2022-05-10T13:05", "2",      "2",
  "1",      "2022-05-11T11:00", "2022-05-11T13:05", "2",      "2",
  # missing dates
  "1",      "",                "",                "3",      "4",
  "1",      "2021-12-30T09:00", "",                "3",      "4",
  "1",      "2021-12-30T11:00", "",                "3",      "3",
  "1",      "",                "2022-01-04T09:00", "3",      "4",
  "1",      "",                "2021-12-24T19:00", "3",      "4",
  "1",      "",                "2022-06-04T09:00", "3",      "4",
  # without treatment
  "2",      "",                "2021-12-03T12:00", "1",      "2",
  "2",      "2021-12-01T12:00", "2021-12-03T12:00", "1",      "2",
  "2",      "2021-12-06T18:00", "",                "1",      "2"
) %>%
  mutate(
    ASTDTM = ymd_hm(ASTDTM),
    AENDTM = ymd_hm(AENDTM),
    TRTSDTM = if_else(USUBJID == "1", ymd_hm("2022-01-01T01:01"), ymd_hms("")),
    TRTEDTM = if_else(USUBJID == "1", ymd_hm("2022-04-30T23:59"), ymd_hms(""))
  )

# derive TRTEMFL without considering treatment end and worsening
derive_var_trtemfl(adae) %>% select(ASTDTM, AENDTM, TRTSDTM, TRTEMFL)

# derive TRTEM2FL taking treatment end and worsening into account
derive_var_trtemfl(
  adae,
  new_var = TRTEM2FL,
  trt_end_date = TRTEDTM,
  end_window = 10,
  initial_intensity = AEITOXGR,
  intensity = AETOXGR
) %>% select(ASTDTM, AENDTM, AEITOXGR, AETOXGR, TRTEM2FL)

```

derive_var_worst_flag *Adds a Variable Flagging the Maximal / Minimal Value Within a Group of Observations*

Description

[Deprecated]

This function is *deprecated*. Please use `slice_derivation()` / `derive_var_extreme_flag()` to derive extreme flags and adjust the order argument.

Usage

```
derive_var_worst_flag(
  dataset,
  by_vars,
  order,
  new_var,
  param_var,
  analysis_var,
  worst_high,
  worst_low,
  check_type = "warning"
)
```

Arguments

dataset	Input dataset. Variables specified by <code>by_vars</code> , <code>order</code> , <code>param_var</code> , and <code>analysis_var</code> are expected.
by_vars	Grouping variables Permitted Values: list of variables
order	Sort order. Used to determine maximal / minimal observation if they are not unique, see Details section for more information.
new_var	Variable to add to the dataset. It is set "Y" for the maximal / minimal observation of each group, see Details section for more information.
param_var	Variable with the parameter values for which the maximal / minimal value is calculated.
analysis_var	Variable with the measurement values for which the maximal / minimal value is calculated.
worst_high	Character with <code>param_var</code> values specifying the parameters referring to "high". Use <code>character(0)</code> if not required.
worst_low	Character with <code>param_var</code> values specifying the parameters referring to "low". Use <code>character(0)</code> if not required.

`check_type` Check uniqueness?
If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order.
Default: "warning"
Permitted Values: "none", "warning", "error"

Details

For each group with respect to the variables specified by the `by_vars` parameter, the maximal / minimal observation of `analysis_var` is labeled in the `new_var` column as "Y", if its `param_var` is in `worst_high / worst_low`. Otherwise, it is assigned NA. If there is more than one such maximal / minimal observation, the first one with respect to the order specified by the `order` parameter is flagged. The direction of "worst" depends on the definition of worst for a specified parameters in the arguments `worst_high / worst_low`, i.e. for some parameters the highest value is the worst and for others the worst is the lowest value.

Value

The input dataset with the new flag variable added.

See Also

[derive_var_extreme_flag\(\)](#)

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

desc

dplyr desc

Description

See `dplyr::desc` for details.

dose_freq_lookup	<i>Pre-Defined Dose Frequencies</i>
------------------	-------------------------------------

Description

These pre-defined dose frequencies are sourced from **CDISC**. The number of rows to generate using `create_single_dose_dataset()` arguments `start_date` and `end_date` is derived from `DOSE_COUNT`, `DOSE_WINDOW`, and `CONVERSION_FACTOR` with appropriate functions from `lubridate`.

Usage

```
dose_freq_lookup
```

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 86 rows and 5 columns.

Details

`NCI_CODE` and `CDISC_VALUE` are included from the **CDISC** source for traceability.

`DOSE_COUNT` represents the number of doses received in one single unit of `DOSE_WINDOW`. For example, for `CDISC_VALUE=="10 DAYS PER MONTH"`, `DOSE_WINDOW=="MONTH"` and `DOSE_COUNT==10`. Similarly, for `CDISC_VALUE=="EVERY 2 WEEKS"`, `DOSE_WINDOW=="WEEK"` and `DOSE_COUNT==0.5` (to yield one dose every two weeks).

`CONVERSION_FACTOR` is used to convert `DOSE_WINDOW` units "WEEK", "MONTH", and "YEAR" to the unit "DAY".

For example, for `CDISC_VALUE=="10 DAYS PER MONTH"`, `CONVERSION_FACTOR` is 0.0329. One day of a month is assumed to be 1 / 30.4375 of a month (one day is assumed to be 1/365.25 of a year). Given only `start_date` and `end_date` in the aggregate dataset, `CONVERSION_FACTOR` is used to calculate specific dates for `start_date` and `end_date` in the resulting single dose dataset for the doses that occur. In such cases, doses are assumed to occur at evenly spaced increments over the interval.

To see the entire table in the console, run `print(dose_freq_lookup)`.

See Also

[create_single_dose_dataset\(\)](#)

Other metadata: [atoxgr_criteria_ctcv4](#), [atoxgr_criteria_ctcv5](#)

dthcaus_source *Create a dthcaus_source Object*

Description

Create a dthcaus_source Object

Usage

```
dthcaus_source(
  dataset_name,
  filter,
  date,
  order = NULL,
  mode = "first",
  dthcaus,
  traceability_vars = NULL
)
```

Arguments

dataset_name	The name of the dataset, i.e. a string, used to search for the death cause.
filter	An expression used for filtering dataset.
date	A date or datetime variable to be used for sorting dataset.
order	Sort order
	Additional variables to be used for sorting the dataset which is ordered by the date and order. Can be used to avoid duplicate record warning. <i>Default:</i> NULL <i>Permitted Values:</i> list of variables or desc(<variable>) function calls created by exprs(), e.g., exprs(ADT, desc(AVAL)) or NULL
mode	One of "first" or "last". Either the "first" or "last" observation is preserved from the dataset which is ordered by date.
dthcaus	A variable name or a string literal — if a variable name, e.g., AEDECOD, it is the variable in the source dataset to be used to assign values to DTHCAUS; if a string literal, e.g. "Adverse Event", it is the fixed value to be assigned to DTHCAUS.
traceability_vars	A named list returned by <code>exprs()</code> listing the traceability variables, e.g. <code>exprs(DTHDOM = "DS", DTHSEQ = DSSEQ)</code> . The left-hand side (names of the list elements) gives the names of the traceability variables in the returned dataset. The right-hand side (values of the list elements) gives the values of the traceability variables in the returned dataset. These can be either strings or symbols referring to existing variables.

Value

An object of class "dthcaus_source".

See Also

[derive_var_dthcaus\(\)](#)

Source Objects: [basket_select\(\)](#), [censor_source\(\)](#), [date_source\(\)](#), [death_event](#), [event_source\(\)](#), [event\(\)](#), [query\(\)](#), [tte_source\(\)](#)

Examples

```
# Deaths sourced from AE
src_ae <- dthcaus_source(
  dataset_name = "ae",
  filter = AEOUT == "FATAL",
  date = AEDTHDT,
  mode = "first",
  dthcaus = AEDECOD
)

# Deaths sourced from DS
src_ds <- dthcaus_source(
  dataset_name = "ds",
  filter = DSDECOD == "DEATH",
  date = DSSTDT,
  mode = "first",
  dthcaus = DSTERM
)
```

dtm_level

Create a dtm_level object

Description

Create a dtm_level object

Usage

```
dtm_level(level)
```

Arguments

level Datetime level
Permitted Values: "Y" (year, highest level), "M" (month), "D" (day), "h" (hour),
 "m" (minute), "s" (second, lowest level), "n" (none)

Details

A dtm_level object is an ordered factor, i.e., two objects can be compared.

Value

A dtm_level object

See Also

Utilities used for date imputation: [dt_level\(\)](#), [get_imputation_target_date\(\)](#), [get_imputation_target_time\(\)](#), [get_partialdatetime\(\)](#), [restrict_imputed_dtc_dtm\(\)](#), [restrict_imputed_dtc_dt\(\)](#)

dt_level	<i>Create a dt_level object</i>
----------	---------------------------------

Description

Create a dt_level object

Usage

```
dt_level(level)
```

Arguments

level	Date level <i>Permitted Values:</i> "Y" (year, highest level), "M" (month), "D" (day), "n" (none, lowest level)
-------	--

Details

A dt_level object is an ordered factor, i.e., two objects can be compared.

Value

A dt_level object

See Also

Utilities used for date imputation: [dtm_level\(\)](#), [get_imputation_target_date\(\)](#), [get_imputation_target_time\(\)](#), [get_partialdatetime\(\)](#), [restrict_imputed_dtc_dtm\(\)](#), [restrict_imputed_dtc_dt\(\)](#)

event	<i>Create a event Object</i>
-------	------------------------------

Description

The event object is used to define events as input for the `derive_extreme_event()` function.

Usage

```
event(condition, set_values_to = NULL)
```

Arguments

- `condition` An unquoted condition for selecting the observations, which will contribute to the extreme event.
- `set_values_to` A named list returned by `exprs()` defining the variables to be set for the extreme answer, e.g. `exprs(PARAMCD = "WSP", PARAM = "Worst Sleeping Problems")`. The values must be a symbol, a character string, a numeric value, or NA.

Value

An object of class `event`

See Also

[derive_extreme_event\(\)](#)

Source Objects: [basket_select\(\)](#), [censor_source\(\)](#), [date_source\(\)](#), [death_event](#), [dthcaus_source\(\)](#), [event_source\(\)](#), [query\(\)](#), [tte_source\(\)](#)

event_source	<i>Create an event_source Object</i>
--------------	--------------------------------------

Description

`event_source` objects are used to define events as input for the `derive_param_tte()` function.

Usage

```
event_source(dataset_name, filter = NULL, date, set_values_to = NULL)
```

Arguments

- `dataset_name` The name of the source dataset
The name refers to the dataset provided by the `source_datasets` parameter of `derive_param_tte()`.
- `filter` An unquoted condition for selecting the observations from dataset which are events or possible censoring time points.
- `date` A variable providing the date of the event or censoring. A date, or a datetime can be specified. An unquoted symbol is expected.
Refer to `derive_vars_dt()` to impute and derive a date from a date character vector to a date object.
- `set_values_to` A named list returned by `exprs()` defining the variables to be set for the event or censoring, e.g. `exprs(EVENTDESC = "DEATH", SRCDOM = "ADSL", SRCVAR = "DTHDT")`. The values must be a symbol, a character string, a numeric value, or NA.

Value

An object of class `event_source`, inheriting from class `tte_source`

See Also

[derive_param_tte\(\)](#), [censor_source\(\)](#)

Source Objects: [basket_select\(\)](#), [censor_source\(\)](#), [date_source\(\)](#), [death_event](#), [dthcaus_source\(\)](#), [event\(\)](#), [query\(\)](#), [tte_source\(\)](#)

Examples

```
# Death event

event_source(
  dataset_name = "adsl",
  filter = DTHFL == "Y",
  date = DTHDT,
  set_values_to = exprs(
    EVNTDESC = "DEATH",
    SRCDOM = "ADSL",
    SRCVAR = "DTHDT"
  )
)
```

example_qs

Example QS Dataset

Description

An example QS dataset based on the examples from the CDISC ADaM Supplements [Generalized Anxiety Disorder 7-Item Version 2 \(GAD-7\)](#) and [Geriatric Depression Scale Short Form \(GDS-SF\)](#).

Usage

```
example_qs
```

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 176 rows and 11 columns.

Source

Created by (https://github.com/pharmaverse/admiral/blob/main/inst/example_scripts/example_qs.R)

See Also

Other datasets: [admiral_adlb](#), [admiral_adsl](#), [ex_single](#), [queries_mh](#), [queries](#)

exprs	<i>rlang exprs</i>
-------	--------------------

Description

See `rlang::exprs` for details.

extend_source_datasets	<i>Add By Groups to All Datasets if Necessary</i>
------------------------	---

Description

The function ensures that the by variables are contained in all source datasets.

Usage

```
extend_source_datasets(source_datasets, by_vars)
```

Arguments

source_datasets	Source datasets A named list of datasets is expected. Each dataset must contain either all by variables or none of the by variables.
by_vars	By variables

Details

1. The by groups are determined as the union of the by groups occurring in the source datasets.
2. For all source datasets which do not contain the by variables the source dataset is replaced by the cartesian product of the source dataset and the by groups.

Value

The list of extended source datasets

See Also

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_terms\(\)](#), [assert_valid_queries\(\)](#), [filter_date_sources\(\)](#), [format.basket_select\(\)](#), [list_tte_source_objects\(\)](#), [params\(\)](#), [validate_basket_select\(\)](#), [validate_query\(\)](#)

Examples

```

library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

adsl <- tribble(
  ~USUBJID, ~TRTSDT,          ~EOSDT,
  "01",     ymd("2020-12-06"), ymd("2021-03-06"),
  "02",     ymd("2021-01-16"), ymd("2021-02-03")
) %>%
  mutate(STUDYID = "AB42")

ae <- tribble(
  ~USUBJID, ~AESTDTC,          ~AESEQ, ~AEDECOD,
  "01",     "2021-01-03T10:56", 1,      "Flu",
  "01",     "2021-03-04",       2,      "Cough",
  "01",     "2021",             3,      "Flu"
) %>%
  mutate(STUDYID = "AB42")

extend_source_datasets(
  source_datasets = list(adsl = adsl, ae = ae),
  by_vars = exprs(AEDECOD)
)

```

extract_duplicate_records

Extract Duplicate Records

Description

Extract Duplicate Records

Usage

```
extract_duplicate_records(dataset, by_vars)
```

Arguments

dataset	A data frame
by_vars	A list of variables created using <code>exprs()</code> identifying groups of records in which to look for duplicates

Value

A data frame of duplicate records within dataset

See Also

Utilities for Dataset Checking: [get_duplicates_dataset\(\)](#), [get_many_to_one_dataset\(\)](#), [get_one_to_many_dataset\(\)](#)

Examples

```
data(admiral_adsl)

# Duplicate the first record
adsl <- rbind(admiral_adsl[1L, ], admiral_adsl)

extract_duplicate_records(adsl, exprs(USUBJID))
```

extract_unit

Extract Unit From Parameter Description

Description

Extract the unit of a parameter from a description like "Param (unit)".

Usage

```
extract_unit(x)
```

Arguments

x A parameter description

Value

A string

See Also

Utilities used within Derivation functions: [call_user_fun\(\)](#), [get_not_mapped\(\)](#), [signal_duplicate_records\(\)](#)

Examples

```
extract_unit("Height (cm)")

extract_unit("Diastolic Blood Pressure (mmHg)")
```

ex_single	<i>Single Dose Exposure Dataset</i>
-----------	-------------------------------------

Description

A derived dataset with single dose per date.

Usage

```
ex_single
```

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 22439 rows and 17 columns.

Source

Derived from the `ex` dataset using `{admiral}` and `{dplyr}` (https://github.com/pharmaverse/admiral/blob/main/inst/example_scripts/derive_single_dose.R)

See Also

Other datasets: [admiral_adlb](#), [admiral_adsl](#), [example_qs](#), [queries_mh](#), [queries](#)

filter_confirmation	<i>Filter Confirmed Observations</i>
---------------------	--------------------------------------

Description**[Deprecated]**

This function is *deprecated*, please use `filter_joined()` instead.

Usage

```
filter_confirmation(  
  dataset,  
  by_vars,  
  join_vars,  
  join_type,  
  first_cond = NULL,  
  order,  
  tmp_obs_nr_var = NULL,  
  filter,  
  check_type = "warning"  
)
```

Arguments

dataset	Input dataset
by_vars	The variables specified for by_vars, join_vars, and order are expected. By variables The specified variables are used as by variables for joining the input dataset with itself.
join_vars	Variables to keep from joined dataset The variables needed from the other observations should be specified for this parameter. The specified variables are added to the joined dataset with suffix ".join". For example to select all observations with AVALC == "Y" and AVISITN == "Y" for at least one subsequent visit join_vars = exprs(AVALC, AVISITN) and filter = AVALC == "Y" & AVALC.join == "Y" & AVISITN < AVISITN.join could be specified. The *.join variables are not included in the output dataset.
join_type	Observations to keep after joining The argument determines which of the joined observations are kept with respect to the original observation. For example, if join_type = "after" is specified all observations after the original observations are kept. <i>Permitted Values:</i> "before", "after", "all"
first_cond	Condition for selecting range of data If this argument is specified, the other observations are restricted up to the first observation where the specified condition is fulfilled. If the condition is not fulfilled for any of the subsequent observations, all observations are removed.
order	Order The observations are ordered by the specified order.
tmp_obs_nr_var	Temporary observation number The specified variable is added to the input dataset and set to the observation number with respect to order. For each by group (by_vars) the observation number starts with 1. The variable can be used in the conditions (filter, first_cond). It is not included in the output dataset. It can be used to select consecutive observations or the last observation (see last example below).
filter	Condition for selecting observations The filter is applied to the joined dataset for selecting the confirmed observations. The condition can include summary functions. The joined dataset is grouped by the original observations. I.e., the summary function are applied to all observations up to the confirmation observation. For example in the oncology setting when using this function for confirmed best overall response, filter = AVALC == "CR" & all(AVALC.join %in% c("CR", "NE")) & count_vals(var = AVALC.join, val = "NE") <= 1 selects observations with response "CR" and for all observations up to the confirmation observation the response is "CR" or "NE" and there is at most one "NE".
check_type	Check uniqueness? If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order.

Default: "none"

Permitted Values: "none", "warning", "error"

Details

The following steps are performed to produce the output dataset.

Step 1:

The input dataset is joined with itself by the variables specified for `by_vars`. From the right hand side of the join only the variables specified for `join_vars` are kept. The suffix ".join" is added to these variables.

For example, for `by_vars = USUBJID`, `join_vars = exprs(AVISITN, AVALC)` and input dataset

```
# A tibble: 2 x 4
  USUBJID AVISITN AVALC  AVAL
<chr>    <dbl> <chr> <dbl>
1         1 Y      1
1         2 N      0
```

the joined dataset is

```
A tibble: 4 x 6
  USUBJID AVISITN AVALC  AVAL AVISITN.join AVALC.join
<chr>    <dbl> <chr> <dbl>    <dbl> <chr>
1         1 Y      1         1 Y
1         1 Y      1         2 N
1         2 N      0         1 Y
1         2 N      0         2 N
```

Step 2:

The joined dataset is restricted to observations with respect to `join_type` and `order`.

The dataset from the example in the previous step with `join_type = "after"` and `order = exprs(AVISITN)` is restricted to

```
A tibble: 4 x 6
  USUBJID AVISITN AVALC  AVAL AVISITN.join AVALC.join
<chr>    <dbl> <chr> <dbl>    <dbl> <chr>
1         1 Y      1         2 N
```

Step 3:

If `first_cond` is specified, for each observation of the input dataset the joined dataset is restricted to observations up to the first observation where `first_cond` is fulfilled (the observation fulfilling the condition is included). If for an observation of the input dataset the condition is not fulfilled, the observation is removed.

Step 4:

The joined dataset is grouped by the observations from the input dataset and restricted to the observations fulfilling the condition specified by `filter`.

Step 5:

The first observation of each group is selected and the `*.join` variables are dropped.

Value

A subset of the observations of the input dataset. All variables of the input dataset are included in the output dataset.

See Also

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

filter_date_sources *Select the First or Last Date from Several Sources*

Description

Select for each subject the first or last observation with respect to a date from a list of sources.

Usage

```
filter_date_sources(
  sources,
  source_datasets,
  by_vars,
  create_datetime = FALSE,
  subject_keys,
  mode
)
```

Arguments

sources	Sources A list of <code>tte_source()</code> objects is expected.
source_datasets	Source datasets A named list of datasets is expected. The <code>dataset_name</code> field of <code>tte_source()</code> refers to the dataset provided in the list.
by_vars	By variables If the parameter is specified, for each by group the observations are selected separately.
create_datetime	Create datetime variable? If set to TRUE, variables ADTM is created. Otherwise, variables ADT is created.
subject_keys	Variables to uniquely identify a subject A list of symbols created using <code>exprs()</code> is expected.

mode Selection mode (first or last)
 If "first" is specified, for each subject the first observation with respect to the date is included in the output dataset. If "last" is specified, the last observation is included in the output dataset.
 Permitted Values: "first", "last"

Details

The following steps are performed to create the output dataset:

1. For each source dataset the observations as specified by the filter element are selected. Then for each patient the first or last observation (with respect to date) is selected.
2. The ADT variable is set to the variable specified by the date element. If the date variable is a datetime variable, only the datepart is copied. If the source variable is a character variable, it is converted to a date. If the date is incomplete, it is imputed as the first possible date.
3. The CNSR is added and set to the value of the censor element.
4. The selected observations of all source datasets are combined into a single dataset.
5. For each patient the first or last observation (with respect to the ADT variable) from the single dataset is selected.

Value

A dataset with one observation per subject as described in the "Details" section.

See Also

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_terms\(\)](#), [assert_valid_queries\(\)](#), [extend_source_datasets\(\)](#), [format.basket_select\(\)](#), [list_tte_source_objects\(\)](#), [params\(\)](#), [validate_basket_select\(\)](#), [validate_query\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(lubridate)

adsl <- tribble(
  ~USUBJID, ~TRTSDT,          ~EOSDT,
  "01",     ymd("2020-12-06"), ymd("2021-03-06"),
  "02",     ymd("2021-01-16"), ymd("2021-02-03")
) %>%
  mutate(STUDYID = "AB42")

ae <- tribble(
  ~USUBJID, ~AESTDTC,      ~AESEQ, ~AEDECOD,
  "01",     "2021-01-03",  1,        "Flu",
  "01",     "2021-03-04",  2,        "Cough",
  "01",     "2021-01-01",  3,        "Flu"
) %>%
```

```

mutate(
  STUDYID = "AB42",
  AESTDT = ymd(AESTDTC)
)

ttae <- event_source(
  dataset_name = "ae",
  date = AESTDT,
  set_values_to = exprs(
    EVNTDESC = "AE",
    SRCDOM = "AE",
    SRCVAR = "AESTDTC",
    SRCSEQ = AESEQ
  )
)

filter_date_sources(
  sources = list(ttae),
  source_datasets = list(adsl = adsl, ae = ae),
  by_vars = exprs(AEDECOD),
  create_datetime = FALSE,
  subject_keys = get_admiral_option("subject_keys"),
  mode = "first"
)

```

filter_extreme

Filter the First or Last Observation for Each By Group

Description

Filters the first or last observation for each by group.

Usage

```
filter_extreme(dataset, by_vars = NULL, order, mode, check_type = "warning")
```

Arguments

dataset	Input dataset The variables specified by the order and the by_vars parameter are expected.
by_vars	Grouping variables <i>Default:</i> NULL <i>Permitted Values:</i> list of variables created by exprs()
order	Sort order Within each by group the observations are ordered by the specified order. <i>Permitted Values:</i> list of variables or desc(<variable>) function calls created by exprs(), e.g., exprs(ADT, desc(AVAL))

mode	<p>Selection mode (first or last)</p> <p>If "first" is specified, the first observation of each by group is included in the output dataset. If "last" is specified, the last observation of each by group is included in the output dataset.</p> <p><i>Permitted Values:</i> "first", "last"</p>
check_type	<p>Check uniqueness?</p> <p>If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order.</p> <p><i>Default:</i> "warning"</p> <p><i>Permitted Values:</i> "none", "warning", "error"</p>

Details

For each group (with respect to the variables specified for the `by_vars` parameter) the first or last observation (with respect to the order specified for the `order` parameter and the mode specified for the `mode` parameter) is included in the output dataset.

Value

A dataset containing the first or last observation of each by group

See Also

Utilities for Filtering Observations: [count_vals\(\)](#), [filter_joined\(\)](#), [filter_relative\(\)](#), [max_cond\(\)](#), [min_cond\(\)](#)

Examples

```
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data("admiral_ex")

# Select first dose for each patient
admiral_ex %>%
  filter_extreme(
    by_vars = exprs(USUBJID),
    order = exprs(EXSEQ),
    mode = "first"
  ) %>%
  select(USUBJID, EXSEQ)

# Select highest dose for each patient on the active drug
admiral_ex %>%
  filter(EXTRT != "PLACEBO") %>%
  filter_extreme(
    by_vars = exprs(USUBJID),
    order = exprs(EXDOSE),
    mode = "last",
    check_type = "none"
  )
```

```
) %>%
select(USUBJID, EXTRT, EXDOSE)
```

 filter_joined

Filter Observations Taking Other Observations into Account

Description

The function filters observation using a condition taking other observations into account. For example, it could select all observations with `AVALC == "Y"` and `AVALC == "Y"` for at least one subsequent observation. The input dataset is joined with itself to enable conditions taking variables from both the current observation and the other observations into account. The suffix ".join" is added to the variables from the subsequent observations.

An example usage might be checking if a patient received two required medications within a certain timeframe of each other.

In the oncology setting, for example, we use such processing to check if a response value can be confirmed by a subsequent assessment. This is commonly used in endpoints such as best overall response.

Usage

```
filter_joined(
  dataset,
  by_vars,
  join_vars,
  join_type,
  first_cond = NULL,
  order,
  tmp_obs_nr_var = NULL,
  filter,
  check_type = "warning"
)
```

Arguments

dataset	Input dataset The variables specified for <code>by_vars</code> , <code>join_vars</code> , and <code>order</code> are expected.
by_vars	By variables The specified variables are used as by variables for joining the input dataset with itself.
join_vars	Variables to keep from joined dataset The variables needed from the other observations should be specified for this parameter. The specified variables are added to the joined dataset with suffix ".join". For example to select all observations with <code>AVALC == "Y"</code> and <code>AVALC == "Y"</code> for at least one subsequent visit <code>join_vars = exprs(AVALC, AVISITN)</code>

	and filter = AVALC == "Y" & AVALC.join == "Y" & AVISITN < AVISITN.join could be specified.
	The *.join variables are not included in the output dataset.
join_type	Observations to keep after joining The argument determines which of the joined observations are kept with respect to the original observation. For example, if join_type = "after" is specified all observations after the original observations are kept. <i>Permitted Values:</i> "before", "after", "all"
first_cond	Condition for selecting range of data If this argument is specified, the other observations are restricted up to the first observation where the specified condition is fulfilled. If the condition is not fulfilled for any of the subsequent observations, all observations are removed.
order	Order The observations are ordered by the specified order.
tmp_obs_nr_var	Temporary observation number The specified variable is added to the input dataset and set to the observation number with respect to order. For each by group (by_vars) the observation number starts with 1. The variable can be used in the conditions (filter, first_cond). It is not included in the output dataset. It can be used to select consecutive observations or the last observation (see last example below).
filter	Condition for selecting observations The filter is applied to the joined dataset for selecting the confirmed observations. The condition can include summary functions. The joined dataset is grouped by the original observations. I.e., the summary function are applied to all observations up to the confirmation observation. For example in the oncology setting when using this function for confirmed best overall response, filter = AVALC == "CR" & all(AVALC.join %in% c("CR", "NE")) & count_vals(var = AVALC.join, val = "NE") <= 1 selects observations with response "CR" and for all observations up to the confirmation observation the response is "CR" or "NE" and there is at most one "NE".
check_type	Check uniqueness? If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order. <i>Default:</i> "none" <i>Permitted Values:</i> "none", "warning", "error"

Details

The following steps are performed to produce the output dataset.

Step 1:

The input dataset is joined with itself by the variables specified for by_vars. From the right hand side of the join only the variables specified for join_vars are kept. The suffix ".join" is added to these variables.

For example, for by_vars = USUBJID, join_vars = exprs(AVISITN, AVALC) and input dataset

```
# A tibble: 2 x 4
  USUBJID AVISITN AVALC  AVAL
<chr>    <dbl> <chr> <dbl>
1         1 Y      1
1         2 N      0
```

the joined dataset is

```
A tibble: 4 x 6
  USUBJID AVISITN AVALC  AVAL AVISITN.join AVALC.join
<chr>    <dbl> <chr> <dbl>      <dbl> <chr>
1         1 Y      1          1 Y
1         1 Y      1          2 N
1         2 N      0          1 Y
1         2 N      0          2 N
```

Step 2:

The joined dataset is restricted to observations with respect to `join_type` and `order`.

The dataset from the example in the previous step with `join_type = "after"` and `order = exprs(AVISITN)` is restricted to

```
A tibble: 4 x 6
  USUBJID AVISITN AVALC  AVAL AVISITN.join AVALC.join
<chr>    <dbl> <chr> <dbl>      <dbl> <chr>
1         1 Y      1          2 N
1         1 Y      1          1 Y
1         2 N      0          1 Y
1         2 N      0          2 N
```

Step 3:

If `first_cond` is specified, for each observation of the input dataset the joined dataset is restricted to observations up to the first observation where `first_cond` is fulfilled (the observation fulfilling the condition is included). If for an observation of the input dataset the condition is not fulfilled, the observation is removed.

Step 4:

The joined dataset is grouped by the observations from the input dataset and restricted to the observations fulfilling the condition specified by `filter`.

Step 5:

The first observation of each group is selected and the `*`.join variables are dropped.

Value

A subset of the observations of the input dataset. All variables of the input dataset are included in the output dataset.

See Also

[count_vals\(\)](#), [min_cond\(\)](#), [max_cond\(\)](#)

Utilities for Filtering Observations: [count_vals\(\)](#), [filter_extreme\(\)](#), [filter_relative\(\)](#), [max_cond\(\)](#), [min_cond\(\)](#)

Examples

```

library(tibble)
library(admiral)

# filter observations with a duration longer than 30 and
# on or after 7 days before a COVID AE (ACOVFL == "Y")
adae <- tribble(
  ~USUBJID, ~ADY, ~ACOVFL, ~ADURN,
  "1",      10, "N",      1,
  "1",      21, "N",     50,
  "1",      23, "Y",     14,
  "1",      32, "N",     31,
  "1",      42, "N",     20,
  "2",      11, "Y",     13,
  "2",      23, "N",      2,
  "3",      13, "Y",     12,
  "4",      14, "N",     32,
  "4",      21, "N",     41
)

filter_joined(
  adae,
  by_vars = exprs(USUBJID),
  join_vars = exprs(ACOVFL, ADY),
  join_type = "all",
  order = exprs(ADY),
  filter = ADURN > 30 & ACOVFL.join == "Y" & ADY >= ADY.join - 7
)

# filter observations with AVALC == "Y" and AVALC == "Y" at a subsequent visit
data <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      1,      "Y",
  "1",      2,      "N",
  "1",      3,      "Y",
  "1",      4,      "N",
  "2",      1,      "Y",
  "2",      2,      "N",
  "3",      1,      "Y",
  "4",      1,      "N",
  "4",      2,      "N",
)

filter_joined(
  data,
  by_vars = exprs(USUBJID),
  join_vars = exprs(AVALC, AVISITN),
  join_type = "after",
  order = exprs(AVISITN),
  filter = AVALC == "Y" & AVALC.join == "Y" & AVISITN < AVISITN.join
)

```

```

# select observations with AVALC == "CR", AVALC == "CR" at a subsequent visit,
# only "CR" or "NE" in between, and at most one "NE" in between
data <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      1,      "PR",
  "1",      2,      "CR",
  "1",      3,      "NE",
  "1",      4,      "CR",
  "1",      5,      "NE",
  "2",      1,      "CR",
  "2",      2,      "PR",
  "2",      3,      "CR",
  "3",      1,      "CR",
  "4",      1,      "CR",
  "4",      2,      "NE",
  "4",      3,      "NE",
  "4",      4,      "CR",
  "4",      5,      "PR"
)

filter_joined(
  data,
  by_vars = exprs(USUBJID),
  join_vars = exprs(AVALC),
  join_type = "after",
  order = exprs(AVISITN),
  first_cond = AVALC.join == "CR",
  filter = AVALC == "CR" & all(AVALC.join %in% c("CR", "NE")) &
    count_vals(var = AVALC.join, val = "NE") <= 1
)

# select observations with AVALC == "PR", AVALC == "CR" or AVALC == "PR"
# at a subsequent visit at least 20 days later, only "CR", "PR", or "NE"
# in between, at most one "NE" in between, and "CR" is not followed by "PR"
data <- tribble(
  ~USUBJID, ~ADY, ~AVALC,
  "1",      6, "PR",
  "1",     12, "CR",
  "1",     24, "NE",
  "1",     32, "CR",
  "1",     48, "PR",
  "2",      3, "PR",
  "2",     21, "CR",
  "2",     33, "PR",
  "3",     11, "PR",
  "4",      7, "PR",
  "4",     12, "NE",
  "4",     24, "NE",
  "4",     32, "PR",
  "4",     55, "PR"
)

```

```

filter_joined(
  data,
  by_vars = exprs(USUBJID),
  join_vars = exprs(AVALC, ADY),
  join_type = "after",
  order = exprs(ADY),
  first_cond = AVALC.join %in% c("CR", "PR") & ADY.join - ADY >= 20,
  filter = AVALC == "PR" &
    all(AVALC.join %in% c("CR", "PR", "NE")) &
    count_vals(var = AVALC.join, val = "NE") <= 1 &
    (
      min_cond(var = ADY.join, cond = AVALC.join == "CR") >
      max_cond(var = ADY.join, cond = AVALC.join == "PR") |
      count_vals(var = AVALC.join, val = "CR") == 0
    )
)

# select observations with CRIT1FL == "Y" at two consecutive visits or at the last visit
data <- tribble(
  ~USUBJID, ~AVISITN, ~CRIT1FL,
  "1",      1,      "Y",
  "1",      2,      "N",
  "1",      3,      "Y",
  "1",      5,      "N",
  "2",      1,      "Y",
  "2",      3,      "Y",
  "2",      5,      "N",
  "3",      1,      "Y",
  "4",      1,      "Y",
  "4",      2,      "N",
)

filter_joined(
  data,
  by_vars = exprs(USUBJID),
  tmp_obs_nr_var = tmp_obs_nr,
  join_vars = exprs(CRIT1FL),
  join_type = "all",
  order = exprs(AVISITN),
  filter = CRIT1FL == "Y" & CRIT1FL.join == "Y" &
    (tmp_obs_nr + 1 == tmp_obs_nr.join | tmp_obs_nr == max(tmp_obs_nr.join))
)

```

filter_relative

Filter the Observations Before or After a Condition is Fulfilled

Description

Filters the observations before or after the observation where a specified condition is fulfilled for each by group. For example, the function could be called to select for each subject all observations

before the first disease progression.

Usage

```
filter_relative(
  dataset,
  by_vars,
  order,
  condition,
  mode,
  selection,
  inclusive,
  keep_no_ref_groups = TRUE,
  check_type = "warning"
)
```

Arguments

dataset	Input dataset The variables specified by the order and the by_vars parameter are expected.
by_vars	Grouping variables <i>Permitted Values:</i> list of variables created by <code>exprs()</code>
order	Sort order Within each by group the observations are ordered by the specified order. <i>Permitted Values:</i> list of variables or <code>desc(<variable>)</code> function calls created by <code>exprs()</code> , e.g., <code>exprs(ADT, desc(AVAL))</code>
condition	Condition for Reference Observation The specified condition determines the reference observation. The output dataset contains all observations before or after (<code>selection</code> parameter) the reference observation.
mode	Selection mode (first or last) If "first" is specified, for each by group the observations before or after (<code>selection</code> parameter) the observation where the condition (<code>condition</code> parameter) is fulfilled the <i>first</i> time is included in the output dataset. If "last" is specified, for each by group the observations before or after (<code>selection</code> parameter) the observation where the condition (<code>condition</code> parameter) is fulfilled the <i>last</i> time is included in the output dataset. <i>Permitted Values:</i> "first", "last"
selection	Select observations before or after the reference observation? <i>Permitted Values:</i> "before", "after"
inclusive	Include the reference observation? <i>Permitted Values:</i> TRUE, FALSE
keep_no_ref_groups	Should by groups without reference observation be kept? <i>Permitted Values:</i> TRUE, FALSE

check_type Check uniqueness?
 If "warning" or "error" is specified, the specified message is issued if the observations of the input dataset are not unique with respect to the by variables and the order.
Permitted Values: "none", "warning", "error"

Details

For each by group (by_vars parameter) the observations before or after (selection parameter) the observations where the condition (condition parameter) is fulfilled the first or last time (order parameter and mode parameter) is included in the output dataset.

Value

A dataset containing for each by group the observations before or after the observation where the condition was fulfilled the first or last time

See Also

Utilities for Filtering Observations: [count_vals\(\)](#), [filter_extreme\(\)](#), [filter_joined\(\)](#), [max_cond\(\)](#), [min_cond\(\)](#)

Examples

```
library(tibble)

response <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      1,      "PR",
  "1",      2,      "CR",
  "1",      3,      "CR",
  "1",      4,      "SD",
  "1",      5,      "NE",
  "2",      1,      "SD",
  "2",      2,      "PD",
  "2",      3,      "PD",
  "3",      1,      "SD",
  "4",      1,      "SD",
  "4",      2,      "PR",
  "4",      3,      "PD",
  "4",      4,      "SD",
  "4",      5,      "PR"
)

# Select observations up to first PD for each patient
response %>%
  filter_relative(
    by_vars = exprs(USUBJID),
    order = exprs(AVISITN),
    condition = AVALC == "PD",
    mode = "first",
```

```

        selection = "before",
        inclusive = TRUE
    )

# Select observations after last CR, PR, or SD for each patient
response %>%
  filter_relative(
    by_vars = exprs(USUBJID),
    order = exprs(AVISITN),
    condition = AVALC %in% c("CR", "PR", "SD"),
    mode = "last",
    selection = "after",
    inclusive = FALSE
  )

# Select observations from first response to first PD
response %>%
  filter_relative(
    by_vars = exprs(USUBJID),
    order = exprs(AVISITN),
    condition = AVALC %in% c("CR", "PR"),
    mode = "first",
    selection = "after",
    inclusive = TRUE,
    keep_no_ref_groups = FALSE
  ) %>%
  filter_relative(
    by_vars = exprs(USUBJID),
    order = exprs(AVISITN),
    condition = AVALC == "PD",
    mode = "first",
    selection = "before",
    inclusive = TRUE
  )

```

format.basket_select *Returns a Character Representation of a basket_select() Object*

Description

The function returns a character representation of a basket_select() object. It can be used for error messages for example.

Usage

```

## S3 method for class 'basket_select'
format(x, ...)

```


Arguments

x	A basket_select() object
...	Not used

Value

A character representation of the basket_select() object

See Also

[basket_select\(\)](#)

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_terms\(\)](#), [assert_valid_queries\(\)](#), [extend_source_datasets\(\)](#), [filter_date_sources\(\)](#), [list_tte_source_objects\(\)](#), [params\(\)](#), [validate_basket_select\(\)](#), [validate_query\(\)](#)

Examples

```
format(basket_select(id = 42, scope = "NARROW", type = "smq"))
```

```
format_eoxxstt_default
```

Default Format for Disposition Status

Description**[Deprecated]**

This function is *deprecated*. This function is a default for derive_var_disposition_status() for the format_new_var argument. Please define your own function and use that as input for the cat_fun argument in derive_var_merged_cat() instead.

Define a function to map the disposition status. To be used as an input for derive_var_disposition_status().

Usage

```
format_eoxxstt_default(status)
```

Arguments

status	the disposition variable used for the mapping (e.g. DSDECOD).
--------	---

Details

Usually this function can not be used with %>%.

Value

A character vector derived based on the values given in status: "NOT STARTED" if status is "SCREEN FAILURE" or "SCREENING NOT COMPLETED", "COMPLETED" if status is "COMPLETED", "DISCONTINUED" if status is not in ("COMPLETED", "SCREEN FAILURE", "SCREENING NOT COMPLETED") nor NA, "ONGOING" otherwise.

See Also

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_reason_default\(\)](#), [vars\(\)](#)

format_reason_default *Default Format for the Disposition Reason*

Description**[Deprecated]**

This function is *deprecated*. This function is a default for `derive_vars_disposition_reason()` for the `format_new_vars` argument. Please use `derive_vars_merged()` and specify the `filter_add` argument to derive the respective variables.

Define a function to map the disposition reason, to be used as a parameter in `derive_vars_disposition_reason()`.

Usage

```
format_reason_default(reason, reason_spe = NULL)
```

Arguments

reason	the disposition variable used for the mapping (e.g. DSDECOD).
reason_spe	the disposition variable used for the mapping of the details if required (e.g. DSTERM).

Details

`format_reason_default(DSDECOD)` returns DSDECOD when DSDECOD is not 'COMPLETED' nor NA.
`format_reason_default(DSDECOD, DSTERM)` returns DSTERM when DSDECOD is equal to 'OTHER'.
 Usually this function can not be used with %>%.

Value

A character vector

See Also

[derive_vars_disposition_reason\(\)](#)

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [vars\(\)](#)

get_admiral_option *Get the Value of an Admiral Option*

Description

Get the Value of an Admiral Option Which Can Be Modified for Advanced Users.

Usage

```
get_admiral_option(option)
```

Arguments

option	A character scalar of commonly used admiral function inputs. As of now, support only available for "subject_keys" or "force_admiral_vars". See set_admiral_options() for a description of the options.
--------	--

Details

This function allows flexibility for function inputs that may need to be repeated multiple times in a script, such as `subject_keys`.

Value

The value of the specified option.

See Also

[set_admiral_options\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_tte\(\)](#), [derive_var_dthcaus\(\)](#), [derive_var_extreme_dtm\(\)](#), [derive_vars_period\(\)](#), [create_period_dataset\(\)](#)

Other admiral_options: [set_admiral_options\(\)](#)

Examples

```
library(admiral.test)
library(dplyr, warn.conflicts = FALSE)
data("admiral_vs")
data("admiral_dm")

# Merging all dm variables to vs
derive_vars_merged(
  admiral_vs,
  dataset_add = select(admiral_dm, -DOMAIN),
  by_vars = get_admiral_option("subject_keys")
) %>%
  select(STUDYID, USUBJID, VSTESTCD, VISIT, VSTPT, VSSTRESN, AGE, AGEU)
```

get_duplicates_dataset

Get Duplicate Records that Led to a Prior Error

Description

Get Duplicate Records that Led to a Prior Error

Usage

```
get_duplicates_dataset()
```

Details

Many admiral function check that the input dataset contains only one record per `by_vars` group and throw an error otherwise. The `get_duplicates_dataset()` function allows one to retrieve the duplicate records that lead to an error.

Note that the function always returns the dataset of duplicates from the last error that has been thrown in the current R session. Thus, after restarting the R sessions `get_duplicates_dataset()` will return NULL and after a second error has been thrown, the dataset of the first error can no longer be accessed (unless it has been saved in a variable).

Value

A `data.frame` or NULL

See Also

Utilities for Dataset Checking: [extract_duplicate_records\(\)](#), [get_many_to_one_dataset\(\)](#), [get_one_to_many_dataset\(\)](#)

Examples

```
data(admiral_adsl)

# Duplicate the first record
adsl <- rbind(admiral_adsl[1L, ], admiral_adsl)

signal_duplicate_records(adsl, exprs(USUBJID), cnd_type = "warning")

get_duplicates_dataset()
```

```
get_imputation_target_date
```

Get Date Imputation Targets

Description

Get Date Imputation Targets

Usage

```
get_imputation_target_date(date_imputation, month)
```

Arguments

date_imputation

The value to impute the day/month when a datepart is missing.

A character value is expected, either as a

- format with month and day specified as "mm-dd": e.g. "06-15" for the 15th of June,
- or as a keyword: "first", "mid", "last" to impute to the first/mid/last day/month.

month

Month component of the partial date

Details

- For date_imputation = "first" "0000", "01", "01" are returned.
- For date_imputation = "mid" "xxxx", "06", "30" if month is NA and "15" otherwise are returned.
- For date_imputation = "last" "9999", "12", "31" are returned.
- For date_imputation = "<mm>-<dd>" "xxxx", "<mm>", "<dd>" are returned.

"xxxx" indicates that the component is undefined. If an undefined component occurs in the imputed DTC value, the imputed DTC value is set to NA_character_ in the imputation functions.

Value

A list of character vectors. The elements of the list are named "year", "month", "day".

See Also

[impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Utilities used for date imputation: [dt_level\(\)](#), [dtm_level\(\)](#), [get_imputation_target_time\(\)](#), [get_partialdatetime\(\)](#), [restrict_imputed_dtc_dtm\(\)](#), [restrict_imputed_dtc_dt\(\)](#)

get_imputation_target_time

Get Time Imputation Targets

Description

Get Time Imputation Targets

Usage

```
get_imputation_target_time(time_imputation)
```

Arguments

time_imputation

The value to impute the time when a timepart is missing.

A character value is expected, either as a

- format with hour, min and sec specified as "hh:mm:ss": e.g. "00:00:00" for the start of the day,
- or as a keyword: "first", "last" to impute to the start/end of a day.

Details

- For time_imputation = "first" "00", "00", "00" are returned.
- For time_imputation = "last" "23", "59", "59" are returned.
- For time_imputation = "<hh>:<mm>:<ss>" "<hh>", "<mm>", "<ss>" are returned.

Value

A list of character vectors. The elements of the list are named "hour", "minute", "second".

See Also

[impute_dtc_dtm\(\)](#)

Utilities used for date imputation: [dt_level\(\)](#), [dtm_level\(\)](#), [get_imputation_target_date\(\)](#), [get_partialdatetime\(\)](#), [restrict_imputed_dtc_dtm\(\)](#), [restrict_imputed_dtc_dt\(\)](#)

`get_many_to_one_dataset`*Get Many to One Values that Led to a Prior Error*

Description

Get Many to One Values that Led to a Prior Error

Usage

```
get_many_to_one_dataset()
```

Details

If `assert_one_to_one()` detects an issue, the many to one values are stored in a dataset. This dataset can be retrieved by `get_many_to_one_dataset()`.

Note that the function always returns the many to one values from the last error that has been thrown in the current R session. Thus, after restarting the R sessions `get_many_to_one_dataset()` will return NULL and after a second error has been thrown, the dataset of the first error can no longer be accessed (unless it has been saved in a variable).

Value

A `data.frame` or NULL

See Also

Utilities for Dataset Checking: [extract_duplicate_records\(\)](#), [get_duplicates_dataset\(\)](#), [get_one_to_many_dataset\(\)](#)

Examples

```
library(admiraldev, warn.conflicts = FALSE)
data(admiral_adsl)

try(
  assert_one_to_one(admiral_adsl, exprs(SITEID), exprs(STUDYID))
)

get_many_to_one_dataset()
```

get_not_mapped	<i>Get list of records not mapped from the lookup table.</i>
----------------	--

Description

Get list of records not mapped from the lookup table.

Usage

```
get_not_mapped()
```

Value

A data.frame or NULL

See Also

Utilities used within Derivation functions: [call_user_fun\(\)](#), [extract_unit\(\)](#), [signal_duplicate_records\(\)](#)

get_one_to_many_dataset	<i>Get One to Many Values that Led to a Prior Error</i>
-------------------------	---

Description

Get One to Many Values that Led to a Prior Error

Usage

```
get_one_to_many_dataset()
```

Details

If `assert_one_to_one()` detects an issue, the one to many values are stored in a dataset. This dataset can be retrieved by `get_one_to_many_dataset()`.

Note that the function always returns the one to many values from the last error that has been thrown in the current R session. Thus, after restarting the R sessions `get_one_to_many_dataset()` will return NULL and after a second error has been thrown, the dataset of the first error can no longer be accessed (unless it has been saved in a variable).

Value

A data.frame or NULL

See Also

Utilities for Dataset Checking: [extract_duplicate_records\(\)](#), [get_duplicates_dataset\(\)](#), [get_many_to_one_dataset\(\)](#)

Examples

```
library(admiraldev, warn.conflicts = FALSE)
data(admiral_adsl)

try(
  assert_one_to_one(admiral_adsl, exprs(STUDYID), exprs(SITEID))
)

get_one_to_many_dataset()
```

get_partialdatetime *Parse DTC variable and Determine Components*

Description

Parse DTC variable and Determine Components

Usage

```
get_partialdatetime(dtc)
```

Arguments

dtc	The '--DTC' date to parse A character date is expected in a format like yyyy-mm-dd or yyyy-mm-ddThh:mm:ss. Trailing components can be omitted and - is a valid value for any component.
-----	---

Details

The function can be replaced by the parttime parser once it is available.

Value

A list of character vectors. The elements of the list are named "year", "month", "day", "hour", "minute", and "second". Missing components are set to NA_character_.

See Also

[impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Utilities used for date imputation: [dt_level\(\)](#), [dtm_level\(\)](#), [get_imputation_target_date\(\)](#), [get_imputation_target_time\(\)](#), [restrict_imputed_dtc_dtm\(\)](#), [restrict_imputed_dtc_dt\(\)](#)

get_summary_records *Create Summary Records*

Description

It is not uncommon to have an analysis need whereby one needs to derive an analysis value (AVAL) from multiple records. The ADaM basic dataset structure variable DTYPE is available to indicate when a new derived records has been added to a dataset.

Usage

```
get_summary_records(
  dataset,
  by_vars,
  filter = NULL,
  analysis_var,
  summary_fun,
  set_values_to = NULL
)
```

Arguments

dataset	A data frame.
by_vars	Variables to consider for generation of groupwise summary records. Providing the names of variables in <code>exprs()</code> will create a groupwise summary and generate summary records for the specified groups.
filter	Filter condition as logical expression to apply during summary calculation. By default, filtering expressions are computed within <code>by_vars</code> as this will help when an aggregating, lagging, or ranking function is involved. For example, <ul style="list-style-type: none"> • <code>filter_rows = (AVAL > mean(AVAL, na.rm = TRUE))</code> will filter all AVAL values greater than mean of AVAL with in <code>by_vars</code>. • <code>filter_rows = (dplyr::n() > 2)</code> will filter n count of <code>by_vars</code> greater than 2.
analysis_var	Analysis variable.
summary_fun	Function that takes as an input the <code>analysis_var</code> and performs the calculation. This can include built-in functions as well as user defined functions, for example <code>mean</code> or <code>function(x) mean(x, na.rm = TRUE)</code> .
set_values_to	A list of variable name-value pairs. Use this argument if you need to change the values of any newly derived records. Set a list of variables to some specified value for the new observation(s) <ul style="list-style-type: none"> • LHS refer to a variable. • RHS refers to the values to set to the variable. This can be a string, a symbol, a numeric value or NA. (e.g. <code>exprs(PARAMCD = "TDOSE", PARCAT1 = "OVERALL")</code>). More general expression are not allowed.

Details

This function only creates derived observations and does not append them to the original dataset observations. If you would like to do this instead, see the `derive_summary_records()` function.

Value

A data frame of derived records.

See Also

`derive_summary_records()`

General Derivation Functions for all ADaMs that returns variable appended to dataset: [derive_var_extreme_flag\(\)](#), [derive_var_joined_exist_flag\(\)](#), [derive_var_last_dose_amt\(\)](#), [derive_var_last_dose_date\(\)](#), [derive_var_last_dose_grp\(\)](#), [derive_var_merged_cat\(\)](#), [derive_var_merged_character\(\)](#), [derive_var_merged_exist_flag\(\)](#), [derive_var_merged_summary\(\)](#), [derive_var_obs_number\(\)](#), [derive_var_relative_flag\(\)](#), [derive_vars_joined\(\)](#), [derive_vars_last_dose\(\)](#), [derive_vars_merged_lookup\(\)](#), [derive_vars_merged\(\)](#), [derive_vars_transposed\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)

adeg <- tribble(
  ~USUBJID, ~EGSEQ, ~PARAM, ~AVISIT, ~EGDTC, ~AVAL, ~TRTA,
  "XYZ-1001", 1, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:50", 385, "",
  "XYZ-1001", 2, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:52", 399, "",
  "XYZ-1001", 3, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:56", 396, "",
  "XYZ-1001", 4, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:45", 384, "Placebo",
  "XYZ-1001", 5, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:48", 393, "Placebo",
  "XYZ-1001", 6, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:51", 388, "Placebo",
  "XYZ-1001", 7, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:45", 385, "Placebo",
  "XYZ-1001", 8, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:48", 394, "Placebo",
  "XYZ-1001", 9, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:51", 402, "Placebo",
  "XYZ-1002", 1, "QTcF Int. (msec)", "Baseline", "2016-02-22T07:58", 399, "",
  "XYZ-1002", 2, "QTcF Int. (msec)", "Baseline", "2016-02-22T07:58", 410, "",
  "XYZ-1002", 3, "QTcF Int. (msec)", "Baseline", "2016-02-22T08:01", 392, "",
  "XYZ-1002", 4, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:50", 401, "Active 20mg",
  "XYZ-1002", 5, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:53", 407, "Active 20mg",
  "XYZ-1002", 6, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:56", 400, "Active 20mg",
  "XYZ-1002", 7, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:50", 412, "Active 20mg",
  "XYZ-1002", 8, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:53", 414, "Active 20mg",
  "XYZ-1002", 9, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:56", 402, "Active 20mg",
)

# Summarize the average of the triplicate ECG interval values (AVAL)
get_summary_records(
  adeg,
  by_vars = exprs(USUBJID, PARAM, AVISIT),
  analysis_var = AVAL,
  summary_fun = function(x) mean(x, na.rm = TRUE),
)
```

```

    set_values_to = exprs(DTYPE = "AVERAGE")
  )

advs <- tribble(
  ~USUBJID, ~VSSEQ, ~PARAM, ~AVAL, ~VSSTRESU, ~VISIT, ~VSDTC,
  "XYZ-001-001", 1164, "Weight", 99, "kg", "Screening", "2018-03-19",
  "XYZ-001-001", 1165, "Weight", 101, "kg", "Run-In", "2018-03-26",
  "XYZ-001-001", 1166, "Weight", 100, "kg", "Baseline", "2018-04-16",
  "XYZ-001-001", 1167, "Weight", 94, "kg", "Week 24", "2018-09-30",
  "XYZ-001-001", 1168, "Weight", 92, "kg", "Week 48", "2019-03-17",
  "XYZ-001-001", 1169, "Weight", 95, "kg", "Week 52", "2019-04-14",
)

# Set new values to any variable. Here, `DTYPE = MAXIMUM` refers to `max()` records
# and `DTYPE = AVERAGE` refers to `mean()` records.
get_summary_records(
  advs,
  by_vars = exprs(USUBJID, PARAM),
  analysis_var = AVAL,
  summary_fun = max,
  set_values_to = exprs(DTYPE = "MAXIMUM")
) %>%
  get_summary_records(
    by_vars = exprs(USUBJID, PARAM),
    analysis_var = AVAL,
    summary_fun = mean,
    set_values_to = exprs(DTYPE = "AVERAGE")
  )

# Sample ADEG dataset with triplicate record for only AVISIT = 'Baseline'
adeg <- tribble(
  ~USUBJID, ~EGSEQ, ~PARAM, ~AVISIT, ~EGDTC, ~AVAL, ~TRTA,
  "XYZ-1001", 1, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:50", 385, "",
  "XYZ-1001", 2, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:52", 399, "",
  "XYZ-1001", 3, "QTcF Int. (msec)", "Baseline", "2016-02-24T07:56", 396, "",
  "XYZ-1001", 4, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:48", 393, "Placebo",
  "XYZ-1001", 5, "QTcF Int. (msec)", "Visit 2", "2016-03-08T09:51", 388, "Placebo",
  "XYZ-1001", 6, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:48", 394, "Placebo",
  "XYZ-1001", 7, "QTcF Int. (msec)", "Visit 3", "2016-03-22T10:51", 402, "Placebo",
  "XYZ-1002", 1, "QTcF Int. (msec)", "Baseline", "2016-02-22T07:58", 399, "",
  "XYZ-1002", 2, "QTcF Int. (msec)", "Baseline", "2016-02-22T07:58", 410, "",
  "XYZ-1002", 3, "QTcF Int. (msec)", "Baseline", "2016-02-22T08:01", 392, "",
  "XYZ-1002", 4, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:53", 407, "Active 20mg",
  "XYZ-1002", 5, "QTcF Int. (msec)", "Visit 2", "2016-03-06T09:56", 400, "Active 20mg",
  "XYZ-1002", 6, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:53", 414, "Active 20mg",
  "XYZ-1002", 7, "QTcF Int. (msec)", "Visit 3", "2016-03-24T10:56", 402, "Active 20mg",
)

# Compute the average of AVAL only if there are more than 2 records within the
# by group
get_summary_records(
  adeg,
  by_vars = exprs(USUBJID, PARAM, AVISIT),

```

```

filter = n() > 2,
analysis_var = AVAL,
summary_fun = function(x) mean(x, na.rm = TRUE),
set_values_to = exprs(DTYPE = "AVERAGE")
)

```

get_terms_from_db

Get Terms from the Queries Database

Description

The function checks if all requirements to access the database are fulfilled (version and access function are available, see `assert_db_requirements()`), reads the terms from the database, and checks if the dataset with the terms is in the expected format (see `assert_terms()`).

Usage

```

get_terms_from_db(
  version,
  fun,
  queries,
  definition,
  expect_query_name = FALSE,
  expect_query_id = FALSE,
  i,
  temp_env
)

```

Arguments

version	Version The version must be non null. Otherwise, an error is issued. The value is passed to the access function (<code>fun</code>).
fun	Access function The access function must be non null. Otherwise, an error is issued. The function is called to retrieve the terms.
queries	Queries List of all queries passed to <code>create_query_data()</code> . It is used for error messages.
definition	Definition of the query The definition is passed to the access function. It defines which terms are returned.
expect_query_name	Is QUERY_NAME expected in the output dataset?
expect_query_id	Is QUERY_ID expected in the output dataset?

i	Index of definition in queries The value is used for error messages.
temp_env	Temporary environment The value is passed to the access function.

Value

Output dataset of the access function

See Also

OCCDS Functions: [derive_var_trtemfl\(\)](#), [derive_vars_atc\(\)](#), [derive_vars_query\(\)](#)

impute_dtc_dt	<i>Impute Partial Date Portion of a '--DTC' Variable</i>
---------------	--

Description

Imputation partial date portion of a '--DTC' variable based on user input.

Usage

```
impute_dtc_dt(
  dtc,
  highest_imputation = "n",
  date_imputation = "first",
  min_dates = NULL,
  max_dates = NULL,
  preserve = FALSE
)
```

Arguments

dtc	The '--DTC' date to impute A character date is expected in a format like yyyy-mm-dd or yyyy-mm-ddThh:mm:ss. Trailing components can be omitted and - is a valid "missing" value for any component.
highest_imputation	Highest imputation level The highest_imputation argument controls which components of the DTC value are imputed if they are missing. All components up to the specified level are imputed. If a component at a higher level than the highest imputation level is missing, NA_character_ is returned. For example, for highest_imputation = "D" "2020" results in NA_character_ because the month is missing.

If "n" is specified no imputation is performed, i.e., if any component is missing, NA_character_ is returned.

If "Y" is specified, date_imputation should be "first" or "last" and min_dates or max_dates should be specified respectively. Otherwise, NA_character_ is returned if the year component is missing.

Default: "n"

Permitted Values: "Y" (year, highest level), "M" (month), "D" (day), "n" (none, lowest level)

date_imputation

The value to impute the day/month when a datepart is missing.

A character value is expected, either as a

- format with month and day specified as "mm-dd": e.g. "06-15" for the 15th of June (The year can not be specified; for imputing the year "first" or "last" together with min_dates or max_dates argument can be used (see examples).),
- or as a keyword: "first", "mid", "last" to impute to the first/mid/last day/month.

The argument is ignored if highest_imputation is less than "D".

Default: "first"

min_dates

Minimum dates

A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example

```
impute_dtc_dtm(
  "2020-11",
  min_dates = list(
    ymd_hms("2020-12-06T12:12:12"),
    ymd_hms("2020-11-11T11:11:11")
  ),
  highest_imputation = "M"
)
```

returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).

max_dates

Maximum dates

A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.

preserve Preserve day if month is missing and day is present
 For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "MID").
 Permitted Values: TRUE, FALSE
 Default: FALSE

Details

Usually this computation function can not be used with %>%.

Value

A character vector

See Also

Date/Time Computation Functions that returns a vector: [compute_dtf\(\)](#), [compute_duration\(\)](#), [compute_tmf\(\)](#), [convert_date_to_dtm\(\)](#), [convert_dtc_to_dtm\(\)](#), [convert_dtc_to_dt\(\)](#), [impute_dtc_dtm\(\)](#)

Examples

```
library(lubridate)

dates <- c(
  "2019-07-18T15:25:40",
  "2019-07-18T15:25",
  "2019-07-18T15",
  "2019-07-18",
  "2019-02",
  "2019",
  "2019",
  "2019---07",
  ""
)

# No date imputation (highest_imputation defaulted to "n")
impute_dtc_dt(dtc = dates)

# Impute to first day/month if date is partial
impute_dtc_dt(
  dtc = dates,
  highest_imputation = "M"
)

# Same as above
impute_dtc_dtm(
  dtc = dates,
  highest_imputation = "M",
  date_imputation = "01-01"
)

# Impute to last day/month if date is partial
```



```

impute_dtc_dt(
  dtc = dates,
  highest_imputation = "M",
  date_imputation = "last",
)

# Impute to mid day/month if date is partial
impute_dtc_dt(
  dtc = dates,
  highest_imputation = "M",
  date_imputation = "mid"
)

# Impute a date and ensure that the imputed date is not before a list of
# minimum dates
impute_dtc_dt(
  "2020-12",
  min_dates = list(
    ymd("2020-12-06"),
    ymd("2020-11-11")
  ),
  highest_imputation = "M"
)

# Impute completely missing dates (only possible if min_dates or max_dates is specified)
impute_dtc_dt(
  c("2020-12", NA_character_),
  min_dates = list(
    ymd("2020-12-06", "2020-01-01"),
    ymd("2020-11-11", NA)
  ),
  highest_imputation = "Y"
)

```

impute_dtc_dtm

Impute Partial Date(-time) Portion of a '--DTC' Variable

Description

Imputation partial date/time portion of a '--DTC' variable. based on user input.

Usage

```

impute_dtc_dtm(
  dtc,
  highest_imputation = "h",
  date_imputation = "first",
  time_imputation = "first",
  min_dates = NULL,
  max_dates = NULL,
)

```

```

    preserve = FALSE
  )

```

Arguments

- dtc** The '--DTC' date to impute
A character date is expected in a format like yyyy-mm-dd or yyyy-mm-ddThh:mm:ss. Trailing components can be omitted and - is a valid "missing" value for any component.
- highest_imputation**
Highest imputation level
The highest_imputation argument controls which components of the DTC value are imputed if they are missing. All components up to the specified level are imputed.
If a component at a higher level than the highest imputation level is missing, NA_character_ is returned. For example, for highest_imputation = "D" "2020" results in NA_character_ because the month is missing.
If "n" is specified, no imputation is performed, i.e., if any component is missing, NA_character_ is returned.
If "Y" is specified, date_imputation should be "first" or "last" and min_dates or max_dates should be specified respectively. Otherwise, NA_character_ is returned if the year component is missing.
Default: "h"
Permitted Values: "Y" (year, highest level), "M" (month), "D" (day), "h" (hour), "m" (minute), "s" (second), "n" (none, lowest level)
- date_imputation**
The value to impute the day/month when a datepart is missing.
A character value is expected, either as a
- format with month and day specified as "mm-dd": e.g. "06-15" for the 15th of June (The year can not be specified; for imputing the year "first" or "last" together with min_dates or max_dates argument can be used (see examples).),
 - or as a keyword: "first", "mid", "last" to impute to the first/mid/last day/month.
- The argument is ignored if highest_imputation is less than "D".
Default: "first".
- time_imputation**
The value to impute the time when a timepart is missing.
A character value is expected, either as a
- format with hour, min and sec specified as "hh:mm:ss": e.g. "00:00:00" for the start of the day,
 - or as a keyword: "first", "last" to impute to the start/end of a day.
- The argument is ignored if highest_imputation = "n".
Default: "first".

min_dates	<p>Minimum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example</p> <pre>impute_dtc_dtm("2020-11", min_dates = list(ymd_hms("2020-12-06T12:12:12"), ymd_hms("2020-11-11T11:11:11")), highest_imputation = "M")</pre> <p>returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).</p> <p>For date variables (not datetime) in the list the time is imputed to "00:00:00". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.</p>
max_dates	<p>Maximum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.</p> <p>For date variables (not datetime) in the list the time is imputed to "23:59:59". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.</p>
preserve	<p>Preserve day if month is missing and day is present</p> <p>For example "2019--07" would return "2019-06-07" if preserve = TRUE (and date_imputation = "mid").</p> <p>Permitted Values: TRUE, FALSE</p> <p><i>Default:</i> FALSE</p>

Details

Usually this computation function can not be used with %>%.

Value

A character vector

See Also

Date/Time Computation Functions that returns a vector: [compute_dtf\(\)](#), [compute_duration\(\)](#), [compute_tmf\(\)](#), [convert_date_to_dtm\(\)](#), [convert_dtc_to_dtm\(\)](#), [convert_dtc_to_dt\(\)](#), [impute_dtc_dt\(\)](#)

Examples

```
library(lubridate)

dates <- c(
  "2019-07-18T15:25:40",
  "2019-07-18T15:25",
  "2019-07-18T15",
  "2019-07-18",
  "2019-02",
  "2019",
  "2019",
  "2019",
  "2019---07",
  ""
)

# No date imputation (highest_imputation defaulted to "h")
# Missing time part imputed with 00:00:00 portion by default
impute_dtc_dtm(dtc = dates)

# No date imputation (highest_imputation defaulted to "h")
# Missing time part imputed with 23:59:59 portion
impute_dtc_dtm(
  dtc = dates,
  time_imputation = "23:59:59"
)

# Same as above
impute_dtc_dtm(
  dtc = dates,
  time_imputation = "last"
)

# Impute to first day/month if date is partial
# Missing time part imputed with 00:00:00 portion by default
impute_dtc_dtm(
  dtc = dates,
  highest_imputation = "M"
)

# same as above
impute_dtc_dtm(
  dtc = dates,
  highest_imputation = "M",
  date_imputation = "01-01"
)

# Impute to last day/month if date is partial
# Missing time part imputed with 23:59:59 portion
```

```
impute_dtc_dtm(  
  dtc = dates,  
  date_imputation = "last",  
  time_imputation = "last"  
)  
  
# Impute to mid day/month if date is partial  
# Missing time part imputed with 00:00:00 portion by default  
impute_dtc_dtm(  
  dtc = dates,  
  highest_imputation = "M",  
  date_imputation = "mid"  
)  
  
# Impute a date and ensure that the imputed date is not before a list of  
# minimum dates  
impute_dtc_dtm(  
  "2020-12",  
  min_dates = list(  
    ymd_hms("2020-12-06T12:12:12"),  
    ymd_hms("2020-11-11T11:11:11")  
  ),  
  highest_imputation = "M"  
)  
  
# Impute completely missing dates (only possible if min_dates or max_dates is specified)  
impute_dtc_dtm(  
  c("2020-12", NA_character_),  
  min_dates = list(  
    ymd_hms("2020-12-06T12:12:12", "2020-01-01T01:01:01"),  
    ymd_hms("2020-11-11T11:11:11", NA)  
  ),  
  highest_imputation = "Y"  
)
```

list_all_templates *List All Available ADaM Templates*

Description

List All Available ADaM Templates

Usage

```
list_all_templates(package = "admiral")
```

Arguments

package The R package in which to look for templates. By default "admiral".

Value

A character vector of all available templates

See Also

Utilities used for examples and template scripts: [use_ad_template\(\)](#)

Examples

```
list_all_templates()
```

```
list_tte_source_objects
```

List all tte_source Objects Available in a Package

Description

List all tte_source Objects Available in a Package

Usage

```
list_tte_source_objects(package = "admiral")
```

Arguments

package The name of the package in which to search for tte_source objects

Value

A data.frame where each row corresponds to one tte_source object or NULL if package does not contain any tte_source objects

See Also

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_terms\(\)](#), [assert_valid_queries\(\)](#), [extend_source_datasets\(\)](#), [filter_date_sources\(\)](#), [format.basket_select\(\)](#), [params\(\)](#), [validate_basket_select\(\)](#), [validate_query\(\)](#)

Examples

```
list_tte_source_objects()
```

max_cond	<i>Maximum Value on a Subset</i>
----------	----------------------------------

Description

The function derives the maximum value of a vector/column on a subset of entries/observations.

Usage

```
max_cond(var, cond)
```

Arguments

var	A vector
cond	A condition

See Also

Utilities for Filtering Observations: [count_vals\(\)](#), [filter_extreme\(\)](#), [filter_joined\(\)](#), [filter_relative\(\)](#), [min_cond\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(admiral)
data <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      1,      "PR",
  "1",      2,      "CR",
  "1",      3,      "NE",
  "1",      4,      "CR",
  "1",      5,      "NE",
  "2",      1,      "CR",
  "2",      2,      "PR",
  "2",      3,      "CR",
)

# In oncology setting, when needing to check the first time a patient had
# a Complete Response (CR) to compare to see if any Partial Response (PR)
# occurred after this add variable indicating if PR occurred after CR
group_by(data, USUBJID) %>% mutate(
  first_cr_vis = min_cond(var = AVISITN, cond = AVALC == "CR"),
  last_pr_vis = max_cond(var = AVISITN, cond = AVALC == "PR"),
  pr_after_cr = last_pr_vis > first_cr_vis
)
```

min_cond	<i>Minimum Value on a Subset</i>
----------	----------------------------------

Description

The function derives the minimum value of a vector/column on a subset of entries/observations.

Usage

```
min_cond(var, cond)
```

Arguments

var	A vector
cond	A condition

See Also

Utilities for Filtering Observations: [count_vals\(\)](#), [filter_extreme\(\)](#), [filter_joined\(\)](#), [filter_relative\(\)](#), [max_cond\(\)](#)

Examples

```
library(tibble)
library(dplyr, warn.conflicts = FALSE)
library(admiral)
data <- tribble(
  ~USUBJID, ~AVISITN, ~AVALC,
  "1",      1,      "PR",
  "1",      2,      "CR",
  "1",      3,      "NE",
  "1",      4,      "CR",
  "1",      5,      "NE",
  "2",      1,      "CR",
  "2",      2,      "PR",
  "2",      3,      "CR",
)

# In oncology setting, when needing to check the first time a patient had
# a Complete Response (CR) to compare to see if any Partial Response (PR)
# occurred after this add variable indicating if PR occurred after CR
group_by(data, USUBJID) %>% mutate(
  first_cr_vis = min_cond(var = AVISITN, cond = AVALC == "CR"),
  last_pr_vis = max_cond(var = AVISITN, cond = AVALC == "PR"),
  pr_after_cr = last_pr_vis > first_cr_vis
)
```

negate_vars	<i>Negate List of Variables</i>
-------------	---------------------------------

Description

The function adds a minus sign as prefix to each variable.

Usage

```
negate_vars(vars = NULL)
```

Arguments

vars List of variables created by `exprs()`

Details

This is useful if a list of variables should be removed from a dataset, e.g., `select(!negate_vars(by_vars))` removes all by variables.

Value

A list of expressions

See Also

Other `utils_` functions: [chr2vars\(\)](#)

Examples

```
negate_vars(exprs(USUBJID, STUDYID))
```

params	<i>Create a Set of Parameters</i>
--------	-----------------------------------

Description

Create a set of variable parameters/function arguments to be used in [call_derivation\(\)](#).

Usage

```
params(...)
```

Arguments

... One or more named arguments

Value

An object of class `params`

See Also

[call_derivation\(\)](#)

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_terms\(\)](#), [assert_valid_queries\(\)](#), [extend_source_datasets\(\)](#), [filter_date_sources\(\)](#), [format.basket_select\(\)](#), [list_tte_source_objects\(\)](#), [validate_basket_select\(\)](#), [validate_query\(\)](#)

Examples

```
library(dplyr, warn.conflicts = FALSE)
library(admiral.test)
data(admiral_ae)
data(admiral_adsl)

adae <- admiral_ae[sample(1:nrow(admiral_ae), 1000), ] %>%
  select(USUBJID, AESTDTC, AEENDTC) %>%
  derive_vars_merged(
    dataset_add = admiral_adsl,
    new_vars = exprs(TRTSDT, TRTEDT),
    by_vars = exprs(USUBJID)
  )

## In order to derive both `ASTDT` and `AENDT` in `ADAE`, one can use `derive_vars_dt()`
adae %>%
  derive_vars_dt(
    new_vars_prefix = "AST",
    dtc = AESTDTC,
    date_imputation = "first",
    min_dates = exprs(TRTSDT),
    max_dates = exprs(TRTEDT)
  ) %>%
  derive_vars_dt(
    new_vars_prefix = "AEN",
    dtc = AEENDTC,
    date_imputation = "last",
    min_dates = exprs(TRTSDT),
    max_dates = exprs(TRTEDT)
  )

## While `derive_vars_dt()` can only add one variable at a time, using `call_derivation()`
## one can add multiple variables in one go.
## The function arguments which are different from a variable to another (e.g. `new_vars_prefix`,
## `dtc`, and `date_imputation`) are specified as a list of `params()` in the `variable_params`
## argument of `call_derivation()`. All other arguments which are common to all variables
## (e.g. `min_dates` and `max_dates`) are specified outside of `variable_params` (i.e. in `...`).
call_derivation(
  dataset = adae,
  derivation = derive_vars_dt,
```

```
variable_params = list(
  params(dtc = AESTDTC, date_imputation = "first", new_vars_prefix = "AST"),
  params(dtc = AEENDTC, date_imputation = "last", new_vars_prefix = "AEN")
),
min_dates = exprs(TRTSDT),
max_dates = exprs(TRTEDT)
)

## The above call using `call_derivation()` is equivalent to the call using `derive_vars_dt()`
## to derive variables `ASTDT` and `AENDT` separately at the beginning.
```

print.adam_templates *Print adam_templates Objects*

Description

Print adam_templates Objects

Usage

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

Arguments

x	A adam_templates object
...	Not used

Value

No return value, called for side effects

See Also

[list_all_templates\(\)](#)

Utilities for printing: [print.source\(\)](#), [print_named_list\(\)](#)

Examples

```
templates <- list_all_templates()
print(templates)
```

print.source	<i>Print source Objects</i>
--------------	-----------------------------

Description

Print source Objects

Usage

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

Arguments

x	An source object
...	If indent = <numeric value> is specified the output is indented by the specified number of characters.

Value

No return value, called for side effects

See Also

Utilities for printing: [print.adam_templates\(\)](#), [print_named_list\(\)](#)

Examples

```
print(death_event)
```

print_named_list	<i>Print Named List</i>
------------------	-------------------------

Description

Print Named List

Usage

```
print_named_list(list, indent = 0)
```

Arguments

list	A named list
indent	Indent The output is indented by the specified number of characters.

Value

No return value, called for side effects

See Also

Utilities for printing: [print.adam_templates\(\)](#), [print.source\(\)](#)

Examples

```
print_named_list(death_event)
```

queries

Queries Dataset

Description

Queries Dataset

Usage

```
queries
```

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 15 rows and 8 columns.

Source

An example of standard query dataset to be used in deriving variables in ADAE and ADCM

See Also

Other datasets: [admiral_adlb](#), [admiral_adsl](#), [ex_single](#), [example_qs](#), [queries_mh](#)

 queries_mh

Queries MH Dataset

Description

Queries MH Dataset

Usage

```
queries_mh
```

Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 14 rows and 8 columns.

Source

An example of standard query MH dataset to be used in deriving variables in ADMH

See Also

Other datasets: [admiral_adlb](#), [admiral_adsl](#), [ex_single](#), [example_qs](#), [queries](#)

 query

Create an query object

Description

A query object defines a query, e.g., a Standard MedDRA Query (SMQ), a Standardized Drug Grouping (SDG), or a customized query (CQ). It is used as input to `create_query_data()`.

Usage

```
query(prefix, name = auto, id = NULL, add_scope_num = FALSE, definition = NULL)
```

Arguments

prefix	The value is used to populate <code>VAR_PREFIX</code> in the output dataset of <code>create_query_data()</code> , e.g., "SMQ03"
name	The value is used to populate <code>QUERY_NAME</code> in the output dataset of <code>create_query_data()</code> . If the <code>auto</code> keyword is specified, the variable is set to the name of the query in the SMQ/SDG database. <i>Permitted Values:</i> A character scalar or the <code>auto</code> keyword. The <code>auto</code> keyword is permitted only for queries which are defined by an <code>basket_select()</code> object.

id	<p>The value is used to populate QUERY_ID in the output dataset of <code>create_query_data()</code>. If the <code>auto</code> keyword is specified, the variable is set to the id of the query in the SMQ/SDG database.</p> <p><i>Permitted Values:</i> A integer scalar or the <code>auto</code> keyword. The <code>auto</code> keyword is permitted only for queries which are defined by an <code>basket_select()</code> object.</p>
add_scope_num	<p>Determines if QUERY_SCOPE_NUM in the output dataset of <code>create_query_data()</code> is populated</p> <p>If the parameter is set to <code>TRUE</code>, the definition must be an <code>basket_select()</code> object.</p> <p><i>Default:</i> <code>FALSE</code></p> <p><i>Permitted Values:</i> <code>TRUE</code>, <code>FALSE</code></p>
definition	<p>Definition of terms belonging to the query</p> <p>There are three different ways to define the terms:</p> <ul style="list-style-type: none"> • An <code>basket_select()</code> object is specified to select a query from the SMQ database. • A data frame with columns <code>TERM_LEVEL</code> and <code>TERM_NAME</code> or <code>TERM_ID</code> can be specified to define the terms of a customized query. The <code>TERM_LEVEL</code> should be set to the name of the variable which should be used to select the terms, e.g., <code>"AEDECOD"</code> or <code>"AELLTCD"</code>. <code>TERM_LEVEL</code> does not need to be constant within a query. For example a query can be based on <code>AEDECOD</code> and <code>AELLT</code>. If <code>TERM_LEVEL</code> refers to a character variable, <code>TERM_NAME</code> should be set to the value the variable. If it refers to a numeric variable, <code>TERM_ID</code> should be set to the value of the variable. If only character variables or only numeric variables are used, <code>TERM_ID</code> or <code>TERM_NAME</code> respectively can be omitted. • A list of data frames and <code>basket_select()</code> objects can be specified to define a customized query based on custom terms and SMQs. The data frames must have the same structure as described for the previous item. <p><i>Permitted Values:</i> an <code>basket_select()</code> object, a data frame, or a list of data frames and <code>basket_select()</code> objects.</p>

Value

An object of class `query`.

See Also

`create_query_data()`, `basket_select()`, [Queries Dataset Documentation](#)

Source Objects: `basket_select()`, `sensor_source()`, `date_source()`, `death_event`, `dthcaus_source()`, `event_source()`, `event()`, `tte_source()`

Examples

```
# create a query for an SMQ
library(tibble)
library(dplyr, warn.conflicts = FALSE)
```

```

# create a query for a SMQ
query(
  prefix = "SMQ02",
  id = auto,
  definition = basket_select(
    name = "Pregnancy and neonatal topics (SMQ)",
    scope = "NARROW",
    type = "smq"
  )
)

# create a query for an SDG
query(
  prefix = "SDG01",
  id = auto,
  definition = basket_select(
    name = "5-aminosalicylates for ulcerative colitis",
    scope = NA_character_,
    type = "sdg"
  )
)

# creating a query for a customized query
cqterms <- tribble(
  ~TERM_NAME, ~TERM_ID,
  "APPLICATION SITE ERYTHEMA", 10003041L,
  "APPLICATION SITE PRURITUS", 10003053L
) %>%
  mutate(TERM_LEVEL = "AEDECOD")

query(
  prefix = "CQ01",
  name = "Application Site Issues",
  definition = cqterms
)

# creating a customized query based on SMQs and additional terms
query(
  prefix = "CQ03",
  name = "Special issues of interest",
  definition = list(
    cqterms,
    basket_select(
      name = "Pregnancy and neonatal topics (SMQ)",
      scope = "NARROW",
      type = "smq"
    ),
    basket_select(
      id = 8050L,
      scope = "BROAD",
      type = "smq"
    )
  )
)

```



```
)
)
```

restrict_derivation *Execute a Derivation on a Subset of the Input Dataset*

Description

Execute a derivation on a subset of the input dataset.

Usage

```
restrict_derivation(dataset, derivation, args = NULL, filter)
```

Arguments

dataset	Input dataset
derivation	Derivation A function that performs a specific derivation is expected. A derivation adds variables or observations to a dataset. The first argument of a derivation must expect a dataset and the derivation must return a dataset. The function must provide the dataset argument and all arguments specified in the <code>params()</code> objects passed to the <code>arg</code> argument. Please note that it is not possible to specify <code>{dplyr}</code> functions like <code>mutate()</code> or <code>summarize()</code> .
args	Arguments of the derivation A <code>params()</code> object is expected.
filter	Filter condition

See Also

[params\(\)](#) [slice_derivation\(\)](#)

Higher Order Functions: [call_derivation\(\)](#), [derivation_slice\(\)](#), [slice_derivation\(\)](#)

Examples

```
library(tibble)

adlb <- tribble(
  ~USUBJID, ~AVISITN, ~AVAL, ~ABLFL,
  "1",      -1,    113, NA_character_,
  "1",      0,    113, "Y",
  "1",      3,    117, NA_character_,
  "2",      0,    95, "Y",
  "3",      0,    111, "Y",
  "3",      1,   101, NA_character_,
```

```

    "3",          2, 123, NA_character_
  )

# Derive BASE for post-baseline records only (derive_var_base() can not be used in this case
# as it requires the baseline observation to be in the input dataset)
restrict_derivation(
  adlb,
  derivation = derive_vars_merged,
  args = params(
    by_vars = exprs(USUBJID),
    dataset_add = adlb,
    filter_add = ABLFL == "Y",
    new_vars = exprs(BASE = AVAL)
  ),
  filter = AVISITN > 0
)

# Derive BASE for baseline and post-baseline records only
restrict_derivation(
  adlb,
  derivation = derive_var_base,
  args = params(
    by_vars = exprs(USUBJID)
  ),
  filter = AVISITN >= 0
) %>%
# Derive CHG for post-baseline records only
restrict_derivation(
  derivation = derive_var_chg,
  filter = AVISITN > 0
)

```

```
restrict_imputed_dtc_dt
```

Restrict Imputed DTC date to Minimum/Maximum Dates

Description

Restrict Imputed DTC date to Minimum/Maximum Dates

Usage

```
restrict_imputed_dtc_dt(dtc, imputed_dtc, min_dates, max_dates)
```

Arguments

dtc	The '--DTC' date to impute A character date is expected in a format like yyyy-mm-dd or yyyy-mm-ddThh:mm:ss. Trailing components can be omitted and - is a valid "missing" value for any component.
-----	--

imputed_dtc	The imputed DTC date
min_dates	<p>Minimum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example</p> <pre> impute_dtc_dtm("2020-11", min_dates = list(ymd_hms("2020-12-06T12:12:12"), ymd_hms("2020-11-11T11:11:11")), highest_imputation = "M") </pre> <p>returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).</p>
max_dates	<p>Maximum dates</p> <p>A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date. Only dates which are in the range of possible dates are considered. A date or date-time object is expected.</p>

Value

- The last of the minimum dates (`min_dates`) which are in the range of the partial DTC date (`dtc`)
- The first of the maximum dates (`max_dates`) which are in the range of the partial DTC date (`dtc`)
- `imputed_dtc` if the partial DTC date (`dtc`) is not in range of any of the minimum or maximum dates.

See Also

[impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Utilities used for date imputation: [dt_level\(\)](#), [dtm_level\(\)](#), [get_imputation_target_date\(\)](#), [get_imputation_target_time\(\)](#), [get_partialdatetime\(\)](#), [restrict_imputed_dtc_dtm\(\)](#)

 restrict_imputed_dtc_dtm

Restrict Imputed DTC date to Minimum/Maximum Dates

Description

Restrict Imputed DTC date to Minimum/Maximum Dates

Usage

```
restrict_imputed_dtc_dtm(dtc, imputed_dtc, min_dates, max_dates)
```

Arguments

dtc The '--DTC' date to impute
 A character date is expected in a format like yyyy-mm-dd or yyyy-mm-ddThh:mm:ss. Trailing components can be omitted and - is a valid "missing" value for any component.

imputed_dtc The imputed DTC date

min_dates Minimum dates

A list of dates is expected. It is ensured that the imputed date is not before any of the specified dates, e.g., that the imputed adverse event start date is not before the first treatment date. Only dates which are in the range of possible dates of the dtc value are considered. The possible dates are defined by the missing parts of the dtc date (see example below). This ensures that the non-missing parts of the dtc date are not changed. A date or date-time object is expected. For example

```
impute_dtc_dtm(
  "2020-11",
  min_dates = list(
    ymd_hms("2020-12-06T12:12:12"),
    ymd_hms("2020-11-11T11:11:11")
  ),
  highest_imputation = "M"
)
```

returns "2020-11-11T11:11:11" because the possible dates for "2020-11" range from "2020-11-01T00:00:00" to "2020-11-30T23:59:59". Therefore "2020-12-06T12:12:12" is ignored. Returning "2020-12-06T12:12:12" would have changed the month although it is not missing (in the dtc date).

For date variables (not datetime) in the list the time is imputed to "00:00:00". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.

max_dates Maximum dates

A list of dates is expected. It is ensured that the imputed date is not after any of the specified dates, e.g., that the imputed date is not after the data cut off date.

Only dates which are in the range of possible dates are considered. A date or date-time object is expected.

For date variables (not datetime) in the list the time is imputed to "23:59:59". Specifying date variables makes sense only if the date is imputed. If only time is imputed, date variables do not affect the result.

Value

- The last of the minimum dates (`min_dates`) which are in the range of the partial DTC date (`dtc`)
- The first of the maximum dates (`max_dates`) which are in the range of the partial DTC date (`dtc`)
- `imputed_dtc` if the partial DTC date (`dtc`) is not in range of any of the minimum or maximum dates.

See Also

[impute_dtc_dtm\(\)](#), [impute_dtc_dt\(\)](#)

Utilities used for date imputation: [dt_level\(\)](#), [dtm_level\(\)](#), [get_imputation_target_date\(\)](#), [get_imputation_target_time\(\)](#), [get_partialdatetime\(\)](#), [restrict_imputed_dtc_dt\(\)](#)

set_admiral_options *Set the Value of Admiral Options*

Description

Set the Values of Admiral Options That Can Be Modified for Advanced Users.

Usage

```
set_admiral_options(subject_keys, force_admiral_vars)
```

Arguments

`subject_keys` Variables to uniquely identify a subject, defaults to `exprs(STUDYID, USUBJID)`. This option is used as default value for the `subject_keys` argument in all admiral functions.

`force_admiral_vars`

If this option is set to `TRUE` (which is the default), the admiral definition of `vars()` is forced. This is just a temporary solution to allow running scripts which use `vars()` in the admiral function calls. It will be removed in a future release.

Details

Modify an admiral option, e.g `subject_keys`, such that it automatically affects downstream function inputs where `get_admiral_option()` is called such as `derive_param_exist_flag()`.

Value

No return value, called for side effects.

See Also

[get_admiral_option\(\)](#), [derive_param_exist_flag\(\)](#), [derive_param_tte\(\)](#), [derive_var_dthcaus\(\)](#), [derive_var_extreme_dtm\(\)](#), [derive_vars_period\(\)](#), [create_period_dataset\(\)](#)

Other `admiral_options`: [get_admiral_option\(\)](#)

Examples

```
library(lubridate)
library(dplyr, warn.conflicts = FALSE)
library(tibble)
set_admiral_options(subject_keys = exprs(STUDYID, USUBJID2))

# Derive a new parameter for measurable disease at baseline
adsl <- tribble(
  ~USUBJID2,
  "1",
  "2",
  "3"
) %>%
  mutate(STUDYID = "XX1234")

tu <- tribble(
  ~USUBJID2, ~VISIT, ~TUSTRESC,
  "1", "SCREENING", "TARGET",
  "1", "WEEK 1", "TARGET",
  "1", "WEEK 5", "TARGET",
  "1", "WEEK 9", "NON-TARGET",
  "2", "SCREENING", "NON-TARGET",
  "2", "SCREENING", "NON-TARGET"
) %>%
  mutate(
    STUDYID = "XX1234",
    TUTESTCD = "TUMIDENT"
  )

derive_param_exist_flag(
  dataset_adsl = adsl,
  dataset_add = tu,
  filter_add = TUTESTCD == "TUMIDENT" & VISIT == "SCREENING",
  condition = TUSTRESC == "TARGET",
  false_value = "N",
  missing_value = "N",
  set_values_to = exprs(
    PARAMCD = "MDIS",
    PARAM = "Measurable Disease at Baseline"
  )
)
```

signal_duplicate_records
Signal Duplicate Records

Description

Signal Duplicate Records

Usage

```
signal_duplicate_records(  
  dataset,  
  by_vars,  
  msg = paste("Dataset contains duplicate records with respect to",  
             enumerate(vars2chr(by_vars))),  
  cnd_type = "error"  
)
```

Arguments

dataset	A data frame
by_vars	A list of variables created using <code>exprs()</code> identifying groups of records in which to look for duplicates
msg	The condition message
cnd_type	Type of condition to signal when detecting duplicate records. One of "message", "warning" or "error". Default is "error".

Value

No return value, called for side effects

See Also

Utilities used within Derivation functions: [call_user_fun\(\)](#), [extract_unit\(\)](#), [get_not_mapped\(\)](#)

Examples

```
data(admiral_adsl)  
  
# Duplicate the first record  
adsl <- rbind(admiral_adsl[1L, ], admiral_adsl)  
  
signal_duplicate_records(adsl, exprs(USUBJID), cnd_type = "message")
```

slice_derivation	<i>Execute a Derivation with Different Arguments for Subsets of the Input Dataset</i>
------------------	---

Description

The input dataset is split into slices (subsets) and for each slice the derivation is called separately. Some or all arguments of the derivation may vary depending on the slice.

Usage

```
slice_derivation(dataset, derivation, args = NULL, ...)
```

Arguments

dataset	Input dataset
derivation	Derivation A function that performs a specific derivation is expected. A derivation adds variables or observations to a dataset. The first argument of a derivation must expect a dataset and the derivation must return a dataset. The function must provide the dataset argument and all arguments specified in the <code>params()</code> objects passed to the <code>arg</code> argument. Please note that it is not possible to specify <code>{dplyr}</code> functions like <code>mutate()</code> or <code>summarize()</code> .
args	Arguments of the derivation A <code>params()</code> object is expected.
...	A <code>derivation_slice()</code> object is expected Each slice defines a subset of the input dataset and some of the parameters for the derivation. The derivation is called on the subset with the parameters specified by the <code>args</code> parameter and the <code>args</code> field of the <code>derivation_slice()</code> object. If a parameter is specified for both, the value in <code>derivation_slice()</code> overwrites the one in <code>args</code> .

Details

For each slice the derivation is called on the subset defined by the `filter` field of the `derivation_slice()` object and with the parameters specified by the `args` parameter and the `args` field of the `derivation_slice()` object. If a parameter is specified for both, the value in `derivation_slice()` overwrites the one in `args`.

- Observations that match with more than one slice are only considered for the first matching slice.
- Observations with no match to any of the slices are included in the output dataset but the derivation is not called for them.

Value

The input dataset with the variables derived by the derivation added

See Also

[params\(\)](#) [restrict_derivation\(\)](#)

Higher Order Functions: [call_derivation\(\)](#), [derivation_slice\(\)](#), [restrict_derivation\(\)](#)

Examples

```
library(tibble)
library(stringr)
advs <- tribble(
  ~USUBJID, ~VSDTC, ~VSTPT,
  "1", "2020-04-16", NA_character_,
  "1", "2020-04-16", "BEFORE TREATMENT"
)

# For the second slice filter is set to TRUE. Thus derive_vars_dtm is called
# with time_imputation = "last" for all observations which do not match for the
# first slice.
slice_derivation(
  advs,
  derivation = derive_vars_dtm,
  args = params(
    dtc = VSDTC,
    new_vars_prefix = "A"
  ),
  derivation_slice(
    filter = str_detect(VSTPT, "PRE|BEFORE"),
    args = params(time_imputation = "first")
  ),
  derivation_slice(
    filter = TRUE,
    args = params(time_imputation = "last")
  )
)
```

tte_source

Create a tte_source Object

Description

The tte_source object is used to define events and possible censorings.

Usage

```
tte_source(dataset_name, filter = NULL, date, censor = 0, set_values_to = NULL)
```

Arguments

dataset_name	The name of the source dataset The name refers to the dataset provided by the source_datasets parameter of derive_param_tte().
filter	An unquoted condition for selecting the observations from dataset which are events or possible censoring time points.
date	A variable providing the date of the event or censoring. A date, or a datetime can be specified. An unquoted symbol is expected. Refer to derive_vars_dt() to impute and derive a date from a date character vector to a date object.
censor	Censoring value CDISC strongly recommends using 0 for events and positive integers for censoring.
set_values_to	A named list returned by exprs() defining the variables to be set for the event or censoring, e.g. exprs(EVENTDESC = "DEATH", SRCDOM = "ADSL", SRCVAR = "DTHDT"). The values must be a symbol, a character string, a numeric value, or NA.

Value

An object of class tte_source

See Also

[derive_param_tte\(\)](#), [censor_source\(\)](#), [event_source\(\)](#)

Source Objects: [basket_select\(\)](#), [censor_source\(\)](#), [date_source\(\)](#), [death_event](#), [dthcaus_source\(\)](#), [event_source\(\)](#), [event\(\)](#), [query\(\)](#)

use_ad_template

Open an ADaM Template Script

Description

Open an ADaM Template Script

Usage

```
use_ad_template(
  adam_name = "adsl",
  save_path = paste0("./", adam_name, ".R"),
  package = "admiral",
  overwrite = FALSE,
  open = interactive()
)
```

Arguments

adam_name	An ADaM dataset name. You can use any of the available dataset name ADAE, ADCM, ADEG, ADEX, ADLB, ADLBHY, ADMH, ADPC, ADPP, ADSL, ADVS, and the dataset name is case-insensitive. The default dataset name is ADSL.
save_path	Path to save the script.
package	The R package in which to look for templates. By default "admiral".
overwrite	Whether to overwrite an existing file named save_path.
open	Whether to open the script right away.

Details

Running without any arguments such as `use_ad_template()` auto-generates `adsl.R` in the current path. Use `list_all_templates()` to discover which templates are available.

Value

No return values, called for side effects

See Also

Utilities used for examples and template scripts: [list_all_templates\(\)](#)

Examples

```
if (interactive()) {  
  use_ad_template("adsl")  
}
```

validate_basket_select

Validate an object is indeed a basket_select object

Description

Validate an object is indeed a `basket_select` object

Usage

```
validate_basket_select(obj)
```

Arguments

obj	An object to be validated.
-----	----------------------------

Value

The original object.

See Also

[basket_select\(\)](#)

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_terms\(\)](#), [assert_valid_queries\(\)](#), [extend_source_datasets\(\)](#), [filter_date_sources\(\)](#), [format.basket_select\(\)](#), [list_tte_source_objects\(\)](#), [params\(\)](#), [validate_query\(\)](#)

validate_query

Validate an object is indeed a query object

Description

Validate an object is indeed a query object

Usage

```
validate_query(obj)
```

Arguments

obj An object to be validated.

Value

The original object.

See Also

[query\(\)](#)

Other Advanced Functions: [assert_db_requirements\(\)](#), [assert_terms\(\)](#), [assert_valid_queries\(\)](#), [extend_source_datasets\(\)](#), [filter_date_sources\(\)](#), [format.basket_select\(\)](#), [list_tte_source_objects\(\)](#), [params\(\)](#), [validate_basket_select\(\)](#)

vars	<i>Create List of Quosures</i>
------	--------------------------------

Description**[Deprecated]**

This function is *deprecated*, please use `exprs()` instead.

Usage

```
vars(...)
```

Arguments

... List of variables

Value

List of expressions

See Also

Other deprecated: [derive_derived_param\(\)](#), [derive_param_first_event\(\)](#), [derive_var_agegr_fda\(\)](#), [derive_var_confirmation_flag\(\)](#), [derive_var_disposition_status\(\)](#), [derive_var_worst_flag\(\)](#), [derive_vars_disposition_reason\(\)](#), [derive_vars_merged_dtm\(\)](#), [derive_vars_merged_dt\(\)](#), [filter_confirmation\(\)](#), [format_eoxxstt_default\(\)](#), [format_reason_default\(\)](#)

yn_to_numeric	<i>Map "Y" and "N" to Numeric Values</i>
---------------	--

Description

Map "Y" and "N" to numeric values.

Usage

```
yn_to_numeric(arg)
```

Arguments

arg Character vector

Value

1 if arg equals "Y", 0 if arg equals "N", `NA_real_` otherwise

See Also

Utilities for Formatting Observations: [convert_blanks_to_na\(\)](#), [convert_na_to_blanks\(\)](#)

Examples

```
yn_to_numeric(c("Y", "N", NA_character_))
```

%>%

Pipe operator

Description

See [magrittr::%>%](#) for more details.

Usage

```
lhs %>% rhs
```

Arguments

lhs	A value or the magrittr placeholder.
rhs	A function call using the magrittr semantics.

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