

# Package ‘sdtm.oak’

September 3, 2024

**Type** Package

**Title** SDTM Data Transformation Engine

**Version** 0.1.0

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**Description** An Electronic Data Capture system (EDC) and Data Standard agnostic solution that enables the pharmaceutical programming community to develop Clinical Data Interchange Standards Consortium (CDISC) Study Data Tabulation Model (SDTM) datasets in R. The reusable algorithms concept in 'sdtm.oak' provides a framework for modular programming and can potentially automate the conversion of raw clinical data to SDTM through standardized SDTM specifications. SDTM is one of the required standards for data submission to the Food and Drug Administration (FDA) in the United States and Pharmaceuticals and Medical Devices Agency (PMDA) in Japan. SDTM standards are implemented following the SDTM Implementation Guide as defined by CDISC <<https://www.cdisc.org/standards/foundational/sdtmig>>.

**Language** en-US

**License** Apache License (>= 2)

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**BugReports** <https://github.com/pharmaverse/sdtm.oak/issues/>

**URL** <https://pharmaverse.github.io/sdtm.oak/>

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.3.2

**Depends** R (>= 4.2)

**Imports** admiraldev (>= 1.1.0), dplyr (>= 1.0.0), purrr (>= 1.0.1), tidyr (>= 1.2.0), rlang (>= 1.0.2), tibble (>= 3.2.0), vctrs (>= 0.5.0), stringr (>= 1.4.0), assertthat, pillar, cli

**Suggests** knitr, htmltools, lifecycle, magrittr, rmarkdown, spelling, testthat (>= 3.1.7), DT, readr

**VignetteBuilder** knitr

**Config/testthat/edition** 3

**Config/testthat/parallel** true

**NeedsCompilation** no

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**Repository** CRAN

**Date/Publication** 2024-09-03 14:40:02 UTC

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---

`assign_datetime`      *Derive an ISO8601 date-time variable*

---

**Description**

`assign_datetime()` maps one or more variables with date/time components in a raw dataset to a target SDTM variable following the ISO8601 format.

**Usage**

```
assign_datetime(  
  tgt_dat = NULL,  
  tgt_var,  
  raw_dat,  
  raw_var,  
  raw_fmt,  
  raw_unk = c("UN", "UNK"),  
  id_vars = oak_id_vars(),  
  .warn = TRUE  
)
```

**Arguments**

<code>tgt_dat</code>	Target dataset: a data frame to be merged against <code>raw_dat</code> by the variables indicated in <code>id_vars</code> . This parameter is optional, see section Value for how the output changes depending on this argument value.
<code>tgt_var</code>	The target SDTM variable: a single string indicating the name of variable to be derived.
<code>raw_dat</code>	The raw dataset (dataframe); must include the variables passed in <code>id_vars</code> and <code>raw_var</code> .
<code>raw_var</code>	The raw variable(s): a character vector indicating the name(s) of the raw variable(s) in <code>raw_dat</code> with date or time components to be parsed into a ISO8601 format variable in <code>tgt_var</code> .
<code>raw_fmt</code>	A date/time parsing format. Either a character vector or a list of character vectors. If a character vector is passed then each element is taken as parsing format for each variable indicated in <code>raw_var</code> . If a list is provided, then each element must be a character vector of formats. The first vector of formats is used for parsing the first variable in <code>raw_var</code> , and so on.
<code>raw_unk</code>	A character vector of string literals to be regarded as missing values during parsing.
<code>id_vars</code>	Key variables to be used in the join between the raw dataset ( <code>raw_dat</code> ) and the target data set ( <code>tgt_dat</code> ).
<code>.warn</code>	Whether to warn about parsing failures.

**Value**

The returned data set depends on the value of `tgt_dat`:

- If no target dataset is supplied, meaning that `tgt_dat` defaults to `NULL`, then the returned data set is `raw_dat`, selected for the variables indicated in `id_vars`, and a new extra column: the derived variable, as indicated in `tgt_var`.
- If the target dataset is provided, then it is merged with the raw data set `raw_dat` by the variables indicated in `id_vars`, with a new column: the derived variable, as indicated in `tgt_var`.

**Examples**

```
# `md1`: an example raw data set.
md1 <-
  tibble::tribble(
    ~oak_id, ~raw_source, ~patient_number, ~MDBDR,      ~MDEDR,      ~MDETM,
    1L,      "MD1",      375,          NA,          NA,          NA,
    2L,      "MD1",      375,          "15-Sep-20", NA,          NA,
    3L,      "MD1",      376,          "17-Feb-21", "17-Feb-21", NA,
    4L,      "MD1",      377,          "4-Oct-20",  NA,          NA,
    5L,      "MD1",      377,          "20-Jan-20", "20-Jan-20", "10:00:00",
    6L,      "MD1",      377,          "UN-UNK-2019", "UN-UNK-2019", NA,
    7L,      "MD1",      377,          "20-UNK-2019", "20-UNK-2019", NA,
    8L,      "MD1",      378,          "UN-UNK-2020", "UN-UNK-2020", NA,
    9L,      "MD1",      378,          "26-Jan-20",  "26-Jan-20",  "07:00:00",
    10L,     "MD1",      378,          "28-Jan-20",  "1-Feb-20",   NA,
    11L,     "MD1",      378,          "12-Feb-20",  "18-Feb-20",  NA,
    12L,     "MD1",      379,          "10-UNK-2020", "20-UNK-2020", NA,
    13L,     "MD1",      379,          NA,           NA,           NA,
    14L,     "MD1",      379,          NA,           "17-Feb-20",  NA
  )

# Using the raw data set `md1`, derive the variable CMSTDTC from MDBDR using
# the parsing format (`raw_fmt`) `d-m-y` (day-month-year), while allowing
# for the presence of special date component values (e.g. `UN` or `UNK`),
# indicating that these values are missing/unknown (unk).
cm1 <-
  assign_datetime(
    tgt_var = "CMSTDTC",
    raw_dat = md1,
    raw_var = "MDBDR",
    raw_fmt = "d-m-y",
    raw_unk = c("UN", "UNK")
  )

cm1

# Inspect parsing failures associated with derivation of CMSTDTC.
problems(cm1$CMSTDTC)

# `cm_inter`: an example target data set.
cm_inter <-
  tibble::tibble(
```

```
oak_id = 1L:14L,
raw_source = "MD1",
patient_number = c(
  375, 375, 376, 377, 377, 377, 377, 378,
  378, 378, 378, 379, 379, 379
),
CMTRT = c(
  "BABY ASPIRIN",
  "CORTISPORIN",
  "ASPIRIN",
  "DIPHENHYDRAMINE HCL",
  "PARCETEMOL",
  "VOMIKIND",
  "ZENFLOX OZ",
  "AMITRYPTYLINE",
  "BENADRYL",
  "DIPHENHYDRAMINE HYDROCHLORIDE",
  "TETRACYCLINE",
  "BENADRYL",
  "SOMINEX",
  "ZQUILL"
),
CMINDC = c(
  "NA",
  "NAUSEA",
  "ANEMIA",
  "NAUSEA",
  "PYREXIA",
  "VOMITINGS",
  "DIARRHHEA",
  "COLD",
  "FEVER",
  "LEG PAIN",
  "FEVER",
  "COLD",
  "COLD",
  "PAIN"
)
)
)

# Same derivation as above but now involving the merging with the target
# data set `cm_inter`.
cm2 <-
  assign_datetime(
    tgt_dat = cm_inter,
    tgt_var = "CMSTDTC",
    raw_dat = md1,
    raw_var = "MDBDR",
    raw_fmt = "d-m-y"
  )

cm2
```

```

# Inspect parsing failures associated with derivation of CMSTDTC.
problems(cm2$CMSTDTC)

# Derive CMSTDTC using both MDEDR and MDETM variables.
# Note that the format `d-m-y` is used for parsing MDEDR and `H:M:S` for
# MDETM (correspondence is by positional matching).
cm3 <-
  assign_datetime(
    tgt_var = "CMSTDTC",
    raw_dat = md1,
    raw_var = c("MDEDR", "MDETM"),
    raw_fmt = c("d-m-y", "H:M:S"),
    raw_unk = c("UN", "UNK")
  )

cm3

# Inspect parsing failures associated with derivation of CMSTDTC.
problems(cm3$CMSTDTC)

```

---

 assign\_no\_ct

*Derive an SDTM variable*


---

### Description

- `assign_no_ct()` maps a variable in a raw dataset to a target SDTM variable that has no terminology restrictions.
- `assign_ct()` maps a variable in a raw dataset to a target SDTM variable following controlled terminology recoding.

### Usage

```

assign_no_ct(
  tgt_dat = NULL,
  tgt_var,
  raw_dat,
  raw_var,
  id_vars = oak_id_vars()
)

assign_ct(
  tgt_dat = NULL,
  tgt_var,
  raw_dat,
  raw_var,
  ct_spec,
  ct_clst,
  id_vars = oak_id_vars()
)

```

**Arguments**

tgt_dat	Target dataset: a data frame to be merged against raw_dat by the variables indicated in id_vars. This parameter is optional, see section Value for how the output changes depending on this argument value.
tgt_var	The target SDTM variable: a single string indicating the name of variable to be derived.
raw_dat	The raw dataset (dataframe); must include the variables passed in id_vars and raw_var.
raw_var	The raw variable: a single string indicating the name of the raw variable in raw_dat.
id_vars	Key variables to be used in the join between the raw dataset (raw_dat) and the target data set (raw_dat).
ct_spec	Study controlled terminology specification: a dataframe with a minimal set of columns, see <code>ct_spec_vars()</code> for details.
ct_clst	A codelist code indicating which subset of the controlled terminology to apply in the derivation.

**Value**

The returned data set depends on the value of tgt\_dat:

- If no target dataset is supplied, meaning that tgt\_dat defaults to NULL, then the returned data set is raw\_dat, selected for the variables indicated in id\_vars, and a new extra column: the derived variable, as indicated in tgt\_var.
- If the target dataset is provided, then it is merged with the raw data set raw\_dat by the variables indicated in id\_vars, with a new column: the derived variable, as indicated in tgt\_var.

**Examples**

```
md1 <-
  tibble::tibble(
    oak_id = 1:14,
    raw_source = "MD1",
    patient_number = 101:114,
    MDIND = c(
      "NAUSEA", "NAUSEA", "ANEMIA", "NAUSEA", "PYREXIA",
      "VOMITINGS", "DIARRHHEA", "COLD",
      "FEVER", "LEG PAIN", "FEVER", "COLD", "COLD", "PAIN"
    )
  )

assign_no_ct(
  tgt_var = "CMINDC",
  raw_dat = md1,
  raw_var = "MDIND"
)

cm_inter <-
```

```
tibble::tibble(
  oak_id = 1:14,
  raw_source = "MD1",
  patient_number = 101:114,
  CMTRT = c(
    "BABY ASPIRIN",
    "CORTISPORIN",
    "ASPIRIN",
    "DIPHENHYDRAMINE HCL",
    "PARCETEMOL",
    "VOMIKIND",
    "ZENFLOX OZ",
    "AMITRYPTYLIN",
    "BENADRYL",
    "DIPHENHYDRAMINE HYDROCHLORIDE",
    "TETRACYCLINE",
    "BENADRYL",
    "SOMINEX",
    "ZQUILL"
  ),
  CMROUTE = c(
    "ORAL",
    "ORAL",
    NA,
    "ORAL",
    "ORAL",
    "ORAL",
    "INTRAMUSCULAR",
    "INTRA-ARTERIAL",
    NA,
    "NON-STANDARD",
    "RANDOM_VALUE",
    "INTRA-ARTICULAR",
    "TRANSDERMAL",
    "OPHTHALMIC"
  )
)

# Controlled terminology specification
(ct_spec <- read_ct_spec_example("ct-01-cm"))

assign_ct(
  tgt_dat = cm_inter,
  tgt_var = "CMINDC",
  raw_dat = md1,
  raw_var = "MDIND",
  ct_spec = ct_spec,
  ct_clst = "C66729"
)
```



---

condition_add	<i>Add filtering tags to a data set</i>
---------------	---

---

### Description

condition\_add() tags records in a data set, indicating which rows match the specified conditions, resulting in a conditioned data frame. Learn how to integrate conditioned data frames in your SDTM domain derivation in vignette("cnd\_df").

### Usage

```
condition_add(dat, ..., .na = NA, .dat2 = rlang::env())
```

### Arguments

dat	A data frame.
...	Conditions to filter the data frame.
.na	Return value to be used when the conditions evaluate to NA.
.dat2	An optional environment to look for variables involved in logical expression passed in ... A data frame or a list can also be passed that will be coerced to an environment internally.

### Value

A conditioned data frame, meaning a tibble with an additional class cnd\_df and a logical vector attribute indicating matching rows.

### Examples

```
(df <- tibble::tibble(x = 1L:3L, y = letters[x]))  
  
# Mark rows for which `x` greater than `1`  
(cnd_df <- condition_add(dat = df, x > 1L))
```

---

create_iso8601	<i>Convert date or time collected values to ISO 8601</i>
----------------	--

---

### Description

create\_iso8601() converts vectors of dates, times or date-times to **ISO 8601** format. Learn more in vignette("iso\_8601").

**Usage**

```
create_iso8601(
  ...,
  .format,
  .fmt_c = fmt_cmp(),
  .na = NULL,
  .cutoff_2000 = 68L,
  .check_format = FALSE,
  .warn = TRUE
)
```

**Arguments**

...	Character vectors of dates, times or date-times' components.
.format	Parsing format(s). Either a character vector or a list of character vectors. If a character vector is passed then each element is taken as parsing format for each vector passed in ... If a list is provided, then each element must be a character vector of formats. The first vector of formats is used for parsing the first vector passed in ..., and so on.
.fmt_c	A list of regexps to use when parsing .format. Use <a href="#">fmt_cmp()</a> to create such an object to pass as argument to this parameter.
.na	A character vector of string literals to be regarded as missing values during parsing.
.cutoff_2000	An integer value. Two-digit years smaller or equal to .cutoff_2000 are parsed as though starting with 20, otherwise parsed as though starting with 19.
.check_format	Whether to check the formats passed in .format, meaning to check against a selection of validated formats in <a href="#">dtc_formats</a> ; or to have a more permissible interpretation of the formats.
.warn	Whether to warn about parsing failures.

**Value**

A vector of dates, times or date-times in **ISO 8601** format

**Examples**

```
# Converting dates
create_iso8601(c("2020-01-01", "20200102"), .format = "y-m-d")
create_iso8601(c("2020-01-01", "20200102"), .format = "ymd")
create_iso8601(c("2020-01-01", "20200102"), .format = list(c("y-m-d", "ymd")))

# Two-digit years are supported
create_iso8601(c("20-01-01", "200101"), .format = list(c("y-m-d", "ymd")))

# `.cutoff_2000` sets the cutoff for two-digit to four-digit year conversion
# Default is at 68.
create_iso8601(c("67-01-01", "68-01-01", "69-01-01"), .format = "y-m-d")
```

```

# Change it to 80.
create_iso8601(c("79-01-01", "80-01-01", "81-01-01"), .format = "y-m-d", .cutoff_2000 = 80)

# Converting times
create_iso8601("15:10", .format = "HH:MM")
create_iso8601("2:10", .format = "HH:MM")
create_iso8601("2:1", .format = "HH:MM")
create_iso8601("02:01:56", .format = "HH:MM:SS")
create_iso8601("020156.5", .format = "HHMMSS")

# Converting date-times
create_iso8601("12 NOV 202015:15", .format = "dd mmm yyyyHH:MM")

# Indicate allowed missing values to make the parsing pass
create_iso8601("U DEC 201914:00", .format = "dd mmm yyyyHH:MM")
create_iso8601("U DEC 201914:00", .format = "dd mmm yyyyHH:MM", .na = "U")

create_iso8601("NOV 2020", .format = "m y")
create_iso8601(c("MAR 2019", "MaR 2020", "mar 2021"), .format = "m y")

create_iso8601("2019-04-041045-", .format = "yyyy-mm-ddHHMM-")

create_iso8601("20200507null", .format = "ymd(HH:MM:SS)")
create_iso8601("20200507null", .format = "ymd((HH:MM:SS)|null)")

# Fractional seconds
create_iso8601("2019-120602:20:13.1230001", .format = "y-mdH:M:S")

# Use different reserved characters in the format specification
# Here we change "H" to "x" and "M" to "w", for hour and minute, respectively.
create_iso8601("14H00M", .format = "HHMM")
create_iso8601("14H00M", .format = "xHwM", .fmt_c = fmt_cmp(hour = "x", min = "w"))

# Alternative formats with unknown values
datetimes <- c("UN UNK 201914:00", "UN JAN 2021")
format <- list(c("dd mmm yyyy", "dd mmm yyyyHH:MM"))
create_iso8601(datetimes, .format = format, .na = c("UN", "UNK"))

# Dates and times may come in many format variations
fmt <- "dd MMM yyyy HH nn ss"
fmt_cmp <- fmt_cmp(mon = "MMM", min = "nn", sec = "ss")
create_iso8601("05 feb 1985 12 55 02", .format = fmt, .fmt_c = fmt_cmp)

```

---

ctl\_new\_rowid\_pillar.cnd\_df

*Conditioned tibble pillar print method*

---

## Description

Conditioned tibble pillar print method

**Usage**

```
## S3 method for class 'cnd_df'
ctl_new_rowid_pillar(controller, x, width, ...)
```

**Arguments**

controller	The object of class "tbl" currently printed.
x	A simple (one-dimensional) vector.
width	The available width, can be a vector for multiple tiers.
...	These dots are for future extensions and must be empty.

**Value**

A character vector to print the tibble which is a conditioned dataframe.

**See Also**

[tbl\\_sum.cnd\\_df\(\)](#).

---

ct\_map

*Recode according to controlled terminology*

---

**Description**

[ct\\_map\(\)](#) recodes a vector following a controlled terminology.

**Usage**

```
ct_map(
  x,
  ct_spec = NULL,
  ct_clst = NULL,
  from = ct_spec_vars("from"),
  to = ct_spec_vars("to")
)
```

**Arguments**

x	A character vector of terms to be recoded following a controlled terminology.
ct_spec	A <a href="#">tibble</a> providing a controlled terminology specification.
ct_clst	A character vector indicating a set of possible controlled terminology codelists codes to be used for recoding. By default (NULL) all codelists available in ct_spec are used.
from	A character vector of column names indicating the variables containing values to be matched against for terminology recoding.
to	A single string indicating the column whose values are to be recoded into.

**Value**

A character vector of terminology recoded values from `x`. If no match is found in the controlled terminology spec provided in `ct_spec`, then `x` values are returned in uppercase. If `ct_spec` is not provided `x` is returned unchanged.

**Examples**

```
# A few example terms.
terms <-
  c(
    "/day",
    "Yes",
    "Unknown",
    "Prior",
    "Every 2 hours",
    "Percentage",
    "International Unit"
  )

# Load a controlled terminology example
(ct_spec <- read_ct_spec_example("ct-01-cm"))

# Use all possible matching terms in the controlled terminology.
ct_map(x = terms, ct_spec = ct_spec)

# Note that if the controlled terminology mapping is restricted to a codelist
# code, e.g. C71113, then only `"/day"` and `"Every 2 hours"` get mapped to
# `"QD"` and `"Q2H"`, respectively; remaining terms won't match given the
# codelist code restriction, and will be mapped to an uppercase version of
# the original terms.
ct_map(x = terms, ct_spec = ct_spec, ct_clst = "C71113")
```

---

 ct\_spec\_example

*Find the path to an example controlled terminology file*


---

**Description**

`ct_spec_example()` resolves the local path to an example controlled terminology file.

**Usage**

```
ct_spec_example(example)
```

**Arguments**

`example` A string with either the basename, file name, or relative path to a controlled terminology file bundled with `{stdm.oak}`, see examples.

**Value**

The local path to an example file if example is supplied, or a character vector of example file names.

**Examples**

```
# Get the local path to controlled terminology example file 01
# Using the basename only:
ct_spec_example("ct-01-cm")

# Using the file name:
ct_spec_example("ct-01-cm.csv")

# Using the relative path:
ct_spec_example("ct/ct-01-cm.csv")

# If no example is provided it returns a vector of possible choices.
ct_spec_example()
```

---

 derive\_bflf

---

*Derive Baseline Flag or Last Observation Before Exposure Flag*


---

**Description**

Derive the baseline flag variable (--BLFL) or the last observation before exposure flag (--LOBXFL), from the observation date/time (--DTC), and a DM domain reference date/time.

**Usage**

```
derive_bflf(
  sdtm_in,
  dm_domain,
  tgt_var,
  ref_var,
  baseline_visits = character(),
  baseline_timepoints = character()
)
```

**Arguments**

sdtm_in	Input SDTM domain.
dm_domain	DM domain with the reference variable ref_var
tgt_var	Name of variable to be derived (--BLFL or --LOBXFL where -- is domain).
ref_var	vector of a date/time from the Demographics (DM) dataset, which serves as a point of comparison for other observations in the study. Common choices for this reference variable include "RFSTDTC" (the date/time of the first study treatment) or "RFXSTDTC" (the date/time of the first exposure to the study drug).

**baseline\_visits**

A character vector specifying the baseline visits within the study. These visits are identified as critical points for data collection at the start of the study, before any intervention is applied. This allows the function to assign the baseline flag if the `--DTC` matches to the reference date.

**baseline\_timepoints**

A character vector of timepoints values in `--TPT` that specifies the specific timepoints during the baseline visits when key assessments or measurements were taken. This allows the function to assign the baseline flag if the `--DTC` matches to the reference date.

**Details**

The derivation is as follows:

- Remove records where the result (`--ORRES`) is missing. Also, exclude records with results labeled as "ND" (No Data) or "NOT DONE" in the `--ORRES` column, which indicate that the measurement or observation was not completed.
- Remove records where the status (`--STAT`) indicates the observation or test was not performed, marked as "NOT DONE".
- Divide the date and time column (`--DTC`) and the reference date/time variable (`ref_var`) into separate date and time components. Ignore any seconds recorded in the time component, focusing only on hours and minutes for further calculations.
- Set partial or missing dates to NA.
- Set partial or missing times to NA.
- Filter on rows that have domain and reference dates not equal to NA. (Ref to as **X**)
- Filter **X** on rows with domain date (`--DTC`) prior to (less than) reference date. (Ref to as **A**)
- Filter **X** on rows with domain date (`--DTC`) equal to reference date but domain and reference times not equal to NA and domain time prior to (less than) reference time. (Ref to as **B**)
- Filter **X** on rows with domain date (`--DTC`) equal to reference date but domain and/or reference time equal to NA and:
  - VISIT is in baseline visits list (if it exists) and
  - xxTPT is in baseline timepoints list (if it exists). (Ref to as **C**)
- Combine the rows from **A**, **B**, and **C** to get a data frame of pre-reference date observations. Sort the rows by `USUBJID`, `--STAT`, and `--ORRES`.
- Group by `USUBJID` and `--TESTCD` and filter on the rows that have maximum value from `--DTC`. Keep only the oak id variables and `--TESTCD` (because these are the unique values). Remove any duplicate rows. Assign the baseline flag variable, `--BLFL`, the last observation before exposure flag (`--LOBXFL`) variable to these rows.
- Join the baseline flag onto the input dataset based on oak id vars

**Value**

Modified input data frame with baseline flag variable `--BLFL` or last observation before exposure flag `--LOBXFL` added.

**Examples**

```
dm <- tibble::tribble(
  ~USUBJID, ~RFSTDTC, ~RFXSTDTC,
  "test_study-375", "2020-09-28T10:10", "2020-09-28T10:10",
  "test_study-376", "2020-09-21T11:00", "2020-09-21T11:00",
  "test_study-377", NA, NA,
  "test_study-378", "2020-01-20T10:00", "2020-01-20T10:00",
  "test_study-379", NA, NA,
)
```

```
dm
```

```
sdtm_in <-
  tibble::tribble(
    ~DOMAIN,
    ~oak_id,
    ~raw_source,
    ~patient_number,
    ~USUBJID,
    ~VSDTC,
    ~VSTESTCD,
    ~VSORRES,
    ~VSSTAT,
    ~VISIT,
    "VS",
    1L,
    "VTLS1",
    375L,
    "test_study-375",
    "2020-09-01T13:31",
    "DIABP",
    "90",
    NA,
    "SCREENING",
    "VS",
    2L,
    "VTLS1",
    375L,
    "test_study-375",
    "2020-10-01T11:20",
    "DIABP",
    "90",
    NA,
    "SCREENING",
    "VS",
    1L,
    "VTLS1",
    375L,
    "test_study-375",
    "2020-09-28T10:10",
    "PULSE",
    "ND",
```



NA,  
"SCREENING",  
"VS",  
2L,  
"VTLS1",  
375L,  
"test\_study-375",  
"2020-10-01T13:31",  
"PULSE",  
"85",  
NA,  
"SCREENING",  
"VS",  
1L,  
"VTLS2",  
375L,  
"test\_study-375",  
"2020-09-28T10:10",  
"SYSBP",  
"120",  
NA,  
"SCREENING",  
"VS",  
2L,  
"VTLS2",  
375L,  
"test\_study-375",  
"2020-09-28T10:05",  
"SYSBP",  
"120",  
NA,  
"SCREENING",  
"VS",  
1L,  
"VTLS1",  
376L,  
"test\_study-376",  
"2020-09-20",  
"DIABP",  
"75",  
NA,  
"SCREENING",  
"VS",  
1L,  
"VTLS1",  
376L,  
"test\_study-376",  
"2020-09-20",  
"PULSE",  
NA,  
"NOT DONE",  
"SCREENING",  
"VS",

```

    2L,
    "VTLS1",
    376L,
    "test_study-376",
    "2020-09-20",
    "PULSE",
    "110",
    NA,
    "SCREENING",
    "VS",
    2L,
    "VTLS1",
    378L,
    "test_study-378",
    "2020-01-20T10:00",
    "PULSE",
    "110",
    NA,
    "SCREENING",
    "VS",
    3L,
    "VTLS1",
    378L,
    "test_study-378",
    "2020-01-21T11:00",
    "PULSE",
    "105",
    NA,
    "SCREENING"
  )

sdtm_in

observed_output <- derive_b1f1(
  sdtm_in = sdtm_in,
  dm_domain = dm,
  tgt_var = "VSL0BXFL",
  ref_var = "RFXSTDTC",
  baseline_visits = c("SCREENING")
)
observed_output

```

---

 derive\_seq

---

*Derive the sequence number (--SEQ) variable*


---

### Description

`derive_seq()` creates a new identifier variable: the sequence number (--SEQ).

This function adds a newly derived variable to `tgt_dat`, namely the sequence number (`--SEQ`) whose name is the one provided in `tgt_var`. An integer sequence is generated that uniquely identifies each record within the domain.

Prior to the derivation of `tgt_var`, the data frame `tgt_dat` is sorted according to grouping variables indicated in `rec_vars`.

### Usage

```
derive_seq(
  tgt_dat,
  tgt_var,
  rec_vars,
  sbj_vars = sdtm.oak::sbj_vars(),
  start_at = 1L
)
```

### Arguments

<code>tgt_dat</code>	The target dataset, a data frame.
<code>tgt_var</code>	The target SDTM variable: a single string indicating the name of the sequence number ( <code>--SEQ</code> ) variable, e.g. "DSSEQ". Note that supplying a name not ending in "SEQ" will raise a warning.
<code>rec_vars</code>	A character vector of record-level identifier variables.
<code>sbj_vars</code>	A character vector of subject-level identifier variables.
<code>start_at</code>	The sequence numbering starts at this value (default is 1).

### Value

Returns the data frame supplied in `tgt_dat` with the newly derived variable, i.e. the sequence number (`--SEQ`), whose name is that passed in `tgt_var`. This variable is of type integer.

### Examples

```
# A VS raw data set example
(vs <- read_domain_example("vs"))

# Derivation of VSSEQ
rec_vars <- c("STUDYID", "USUBJID", "VSTESTCD", "VSDTC", "VSTPTNUM")
derive_seq(tgt_dat = vs, tgt_var = "VSSEQ", rec_vars = rec_vars)

# An APSC raw data set example
(apsc <- read_domain_example("apsc"))

# Derivation of APSEQ
derive_seq(
  tgt_dat = apsc,
  tgt_var = "APSEQ",
  rec_vars = c("STUDYID", "RSUBJID", "SCTESTCD"),
  sbj_vars = c("STUDYID", "RSUBJID")
)
```

---

derive_study_day	derive_study_day performs study day calculation
------------------	---

---

### Description

This function takes the an input data frame and a reference data frame (which is DM domain in most cases), and calculate the study day from reference date and target date. In case of unexpected conditions like reference date is not unique for each patient, or reference and input dates are not actual dates, NA will be returned for those records.

### Usage

```
derive_study_day(
  sdtm_in,
  dm_domain,
  tgdt,
  refdt,
  study_day_var,
  merge_key = "USUBJID"
)
```

### Arguments

sdtm_in	Input data frame that contains the target date.
dm_domain	Reference date frame that contains the reference date.
tgdt	Target date from sdtm_in that will be used to calculate the study day.
refdt	Reference date from dm_domain that will be used as reference to calculate the study day.
study_day_var	New study day variable name in the output. For example, AESTDY for AE domain and CMSTDY for CM domain.
merge_key	Character to represent the merging key between sdtm_in and dm_domain.

### Value

Data frame that takes all columns from sdtm\_in and a new variable to represent the calculated study day.

### Examples

```
ae <- data.frame(
  USUBJID = c("study123-123", "study123-124", "study123-125"),
  AESTDTC = c("2012-01-01", "2012-04-14", "2012-04-14")
)
dm <- data.frame(
  USUBJID = c("study123-123", "study123-124", "study123-125"),
  RFSTDTC = c("2012-02-01", "2012-04-14", NA)
)
```

```
)  
ae$AESTDTC <- as.Date(ae$AESTDTC)  
dm$RFSTDTC <- as.Date(dm$RFSTDTC)  
derive_study_day(ae, dm, "AESTDTC", "RFSTDTC", "AESTDY")
```

---

domain\_example

*Find the path to an example SDTM domain file*

---

## Description

`domain_example()` resolves the local path to a SDTM domain example file. The domain examples files were imported from [pharmaversesdtm](#). See Details section for available datasets.

## Usage

```
domain_example(example)
```

## Arguments

`example` A string with either the basename, file name, or relative path to a SDTM domain example file bundled with `{sdm.oak}`, e.g. "cm" (Concomitant Medication) or "ae" (Adverse Events).

## Details

Datasets were obtained from [pharmaversesdtm](#) but are originally sourced from the [CDISC pilot project](#) or have been constructed ad-hoc by the [admiral](#) team. These datasets are bundled with `{sdm.oak}`, thus obviating a dependence on `{pharmaversesdtm}`.

### Example SDTM domains:

- "ae": Adverse Events (AE) data set.
- "apsc": Associated Persons Subject Characteristics (APSC) data set.
- "cm": Concomitant Medications (CM) data set.
- "vs": Vital Signs (VS) data set.

## Value

The local path to an example file if `example` is supplied, or a character vector of example file names.

## Source

See <https://cran.r-project.org/package=pharmaversesdtm>.

## See Also

[read\\_domain\\_example\(\)](#)

**Examples**

```
# If no example is provided it returns a vector of possible choices.
domain_example()

# Get the local path to the Concomitant Medication dataset file.
domain_example("cm")

# Local path to the Adverse Events dataset file.
domain_example("ae")
```

---

dtc_formats	<i>Date/time collection formats</i>
-------------	-------------------------------------

---

**Description**

Date/time collection formats

**Usage**

```
dtc_formats
```

**Format**

A [tibble](#) of 20 formats with three variables:

fmt Format string.

type Whether a date, time or date-time.

description Description of which date-time components are parsed.

**Examples**

```
dtc_formats
```

---

fmt_cmp	<i>Regexps for date/time format components</i>
---------	--

---

**Description**

[fmt\\_cmp\(\)](#) creates a character vector of patterns to match individual format date/time components.

**Usage**

```

fmt_cmp(
  sec = "S+",
  min = "M+",
  hour = "H+",
  mday = "d+",
  mon = "m+",
  year = "y+"
)

```

**Arguments**

sec	A string pattern for matching the second format component.
min	A string pattern for matching the minute format component.
hour	A string pattern for matching the hour format component.
mday	A string pattern for matching the month day format component.
mon	A string pattern for matching the month format component.
year	A string pattern for matching the year format component.

**Value**

A named character vector of date/time format patterns. This a vector of six elements, one for each date/time component.

**Examples**

```

# Regexp to parse format components
fmt_cmp()

fmt_cmp(year = "yyyy")

```

---

generate\_oak\_id\_vars *A function to generate oak\_id\_vars*

---

**Description**

A function to generate oak\_id\_vars

**Usage**

```
generate_oak_id_vars(raw_dat, pat_var, raw_src)
```

**Arguments**

raw_dat	The raw dataset (dataframe)
pat_var	Variable that holds the patient number
raw_src	Name of the raw source

**Value**

dataframe

**Examples**

```
raw_dataset <-
  tibble::tribble(
    ~patnum, ~MDRAW,
    101L, "BABY ASPIRIN",
    102L, "CORTISPORIN",
    103L, NA_character_,
    104L, "DIPHENHYDRAMINE HCL"
  )

# Generate oak_id_vars
generate_oak_id_vars(
  raw_dat = raw_dataset,
  pat_var = "patnum",
  raw_src = "Concomitant Medication"
)
```

---

harcode

*Derive an SDTM variable with a hardcoded value*

---

**Description**

- `harcode_no_ct()` maps a hardcoded value to a target SDTM variable that has no terminology restrictions.
- `harcode_ct()` maps a hardcoded value to a target SDTM variable with controlled terminology recoding.

**Usage**

```
harcode_no_ct(
  tgt_dat = NULL,
  tgt_val,
  raw_dat,
  raw_var,
  tgt_var,
  id_vars = oak_id_vars()
)
```



```

hardcode_ct(
  tgt_dat = NULL,
  tgt_val,
  raw_dat,
  raw_var,
  tgt_var,
  ct_spec,
  ct_clst,
  id_vars = oak_id_vars()
)

```

### Arguments

<code>tgt_dat</code>	Target dataset: a data frame to be merged against <code>raw_dat</code> by the variables indicated in <code>id_vars</code> . This parameter is optional, see section Value for how the output changes depending on this argument value.
<code>tgt_val</code>	The target SDTM value to be hardcoded into the variable indicated in <code>tgt_var</code> .
<code>raw_dat</code>	The raw dataset (dataframe); must include the variables passed in <code>id_vars</code> and <code>raw_var</code> .
<code>raw_var</code>	The raw variable: a single string indicating the name of the raw variable in <code>raw_dat</code> .
<code>tgt_var</code>	The target SDTM variable: a single string indicating the name of variable to be derived.
<code>id_vars</code>	Key variables to be used in the join between the raw dataset ( <code>raw_dat</code> ) and the target data set ( <code>raw_dat</code> ).
<code>ct_spec</code>	Study controlled terminology specification: a dataframe with a minimal set of columns, see <code>ct_spec_vars()</code> for details. This parameter is optional, if left as NULL no controlled terminology recoding is applied.
<code>ct_clst</code>	A codelist code indicating which subset of the controlled terminology to apply in the derivation. This parameter is optional, if left as NULL, all possible recodings in <code>ct_spec</code> are attempted.

### Value

The returned data set depends on the value of `tgt_dat`:

- If no target dataset is supplied, meaning that `tgt_dat` defaults to NULL, then the returned data set is `raw_dat`, selected for the variables indicated in `id_vars`, and a new extra column: the derived variable, as indicated in `tgt_var`.
- If the target dataset is provided, then it is merged with the raw data set `raw_dat` by the variables indicated in `id_vars`, with a new column: the derived variable, as indicated in `tgt_var`.

### Examples

```

md1 <-
  tibble::tribble(

```

```

    ~oak_id, ~raw_source, ~patient_number, ~MDRAW,
  1L,      "MD1",      101L,      "BABY ASPIRIN",
  2L,      "MD1",      102L,      "CORTISPORIN",
  3L,      "MD1",      103L,      NA_character_,
  4L,      "MD1",      104L,      "DIPHENHYDRAMINE HCL"
)

# Derive a new variable `CMCAT` by overwriting `MDRAW` with the
# hardcoded value "GENERAL CONCOMITANT MEDICATIONS".
hardcode_no_ct(
  tgt_val = "GENERAL CONCOMITANT MEDICATIONS",
  raw_dat = md1,
  raw_var = "MDRAW",
  tgt_var = "CMCAT"
)

cm_inter <-
  tibble::tribble(
    ~oak_id, ~raw_source, ~patient_number, ~CMTRT,      ~CMINDC,
    1L,      "MD1",      101L,      "BABY ASPIRIN",      NA,
    2L,      "MD1",      102L,      "CORTISPORIN",      "NAUSEA",
    3L,      "MD1",      103L,      "ASPIRIN",          "ANEMIA",
    4L,      "MD1",      104L,      "DIPHENHYDRAMINE HCL", "NAUSEA",
    5L,      "MD1",      105L,      "PARACETAMOL",      "PYREXIA"
  )

# Derive a new variable `CMCAT` by overwriting `MDRAW` with the
# hardcoded value "GENERAL CONCOMITANT MEDICATIONS" with a prior join to
# `target_dataset`.
hardcode_no_ct(
  tgt_dat = cm_inter,
  tgt_val = "GENERAL CONCOMITANT MEDICATIONS",
  raw_dat = md1,
  raw_var = "MDRAW",
  tgt_var = "CMCAT"
)

# Controlled terminology specification
(ct_spec <- read_ct_spec_example("ct-01-cm"))

# Hardcoding of `CMCAT` with the value `"GENERAL CONCOMITANT MEDICATIONS"`
# involving terminology recoding. `NA` values in `MDRAW` are preserved in
# `CMCAT`.
hardcode_ct(
  tgt_dat = cm_inter,
  tgt_var = "CMCAT",
  raw_dat = md1,
  raw_var = "MDRAW",
  tgt_val = "GENERAL CONCOMITANT MEDICATIONS",
  ct_spec = ct_spec,
  ct_clst = "C66729"
)

```

---

mutate.cnd_df	<i>Mutate method for conditioned data frames</i>
---------------	--

---

### Description

`mutate.cnd_df()` is an S3 method to be dispatched by `mutate` generic on conditioned data frames. This function implements a conditional mutate by only changing rows for which the condition stored in the conditioned data frame is TRUE.

### Usage

```
## S3 method for class 'cnd_df'
mutate(
  .data,
  ...,
  .by = NULL,
  .keep = c("all", "used", "unused", "none"),
  .before = NULL,
  .after = NULL
)
```

### Arguments

<code>.data</code>	A conditioned data frame.
<code>...</code>	<p><code>&lt;data-masking&gt;</code> Name-value pairs. The name gives the name of the column in the output.</p> <p>The value can be:</p> <ul style="list-style-type: none"> <li>• A vector of length 1, which will be recycled to the correct length.</li> <li>• A vector the same length as the current group (or the whole data frame if ungrouped).</li> <li>• NULL, to remove the column.</li> <li>• A data frame or tibble, to create multiple columns in the output.</li> </ul>
<code>.by</code>	Not used when <code>.data</code> is a conditioned data frame.
<code>.keep</code>	<p>Control which columns from <code>.data</code> are retained in the output. Grouping columns and columns created by <code>...</code> are always kept.</p> <ul style="list-style-type: none"> <li>• "all" retains all columns from <code>.data</code>. This is the default.</li> <li>• "used" retains only the columns used in <code>...</code> to create new columns. This is useful for checking your work, as it displays inputs and outputs side-by-side.</li> <li>• "unused" retains only the columns <i>not</i> used in <code>...</code> to create new columns. This is useful if you generate new columns, but no longer need the columns used to generate them.</li> <li>• "none" doesn't retain any extra columns from <code>.data</code>. Only the grouping variables and columns created by <code>...</code> are kept.</li> </ul>
<code>.before</code>	Not used, use <code>.after</code> instead.
<code>.after</code>	Control where new columns should appear, i.e. after which columns.

**Value**

A conditioned data frame, meaning a tibble with mutated values.

---

oak_id_vars	<i>Raw dataset keys</i>
-------------	-------------------------

---

**Description**

`oak_id_vars()` is a helper function providing the variable (column) names to be regarded as keys in `tibbles` representing raw datasets. By default, the set of names is `oak_id`, `raw_source`, and `patient_number`. Extra variable names may be indicated and passed in `extra_vars` which are appended to the default names.

**Usage**

```
oak_id_vars(extra_vars = NULL)
```

**Arguments**

`extra_vars` A character vector of extra column names to be appended to the default names: `oak_id`, `raw_source`, and `patient_number`.

**Value**

A character vector of column names to be regarded as keys in raw datasets.

---

problems	<i>Retrieve date/time parsing problems</i>
----------	--

---

**Description**

`problems()` is a companion helper function to `create_iso8601()`. It retrieves ISO 8601 parsing problems from an object of class `iso8601`, which is `create_iso8601()`'s return value and that might contain a `problems` attribute in case of parsing failures. `problems()` is a helper function that provides easy access to these parsing problems.

**Usage**

```
problems(x = .Last.value)
```

**Arguments**

`x` An object of class `iso8601`, as typically obtained from a call to `create_iso8601()`. The argument can also be left empty, in that case `problems()` will use the last returned value, making it convenient to use immediately after `create_iso8601()`.

**Value**

If there are no parsing problems in `x`, then the returned value is `NULL`; otherwise, a [tibble](#) of parsing failures is returned. Each row corresponds to a parsing problem. There will be a first column named `..i` indicating the position(s) in the inputs to the `create_iso8601()` call that resulted in failures; remaining columns correspond to the original input values passed on to `create_iso8601()`, with columns being automatically named `..var1`, `..var2`, and so on, if the inputs to `create_iso8601()` were unnamed, otherwise, the original variable names are used instead.

**Examples**

```

dates <-
  c(
    "2020-01-01",
    "2020-02-11",
    "2020-01-06",
    "2020-0921",
    "2020/10/30",
    "2020-12-05",
    "20231225"
  )

# By inspecting the problematic dates it can be understood that
# the `.format` parameter needs to be updated to include other variations.
iso8601_dttm <- create_iso8601(dates, .format = "y-m-d")
problems(iso8601_dttm)

# Including more parsing formats addresses the previous problems
formats <- c("y-m-d", "y-md", "y/m/d", "ymd")
iso8601_dttm2 <- create_iso8601(dates, .format = list(formats))

# So now `problems()` returns `NULL` because there are no more parsing issues.
problems(iso8601_dttm2)

# If you pass named arguments when calling `create_iso8601()` then they will
# be used to create the problems object.
iso8601_dttm3 <- create_iso8601(date = dates, .format = "y-m-d")
problems(iso8601_dttm3)

```

---

read\_ct\_spec

*Read in a controlled terminology*


---

**Description**

`read_ct_spec()` imports a controlled terminology specification data set as a [tibble](#).

**Usage**

```
read_ct_spec(file = cli::cli_abort("`file` must be specified"))
```

## Arguments

- `file` A path to a file containing a controlled terminology specification data set. The following are expected of this file:
- The file is expected to be a CSV file;
  - The file is expected to contain a first row of column names;
  - This minimal set of variables is expected: `codelist_code`, `collected_value`, `term_synonyms`, and `term_value`.

## Value

A [tibble](#) with a controlled terminology specification.

## Examples

```
# Get the local path to one of the controlled terminology example files.
path <- ct_spec_example("ct-01-cm")

# Import it to R.
read_ct_spec(file = path)
```

---

`read_ct_spec_example` *Read an example controlled terminology specification*

---

## Description

`read_ct_spec_example()` imports one of the bundled controlled terminology specification data sets as a [tibble](#) into R.

## Usage

```
read_ct_spec_example(example)
```

## Arguments

- `example` The file name of a controlled terminology data set bundled with `{stdm.oak}`, run `read_ct_spec_example()` for available example files.

## Value

A [tibble](#) with a controlled terminology specification data set, or a character vector of example file names.

## Examples

```
# Leave the `example` parameter as missing for available example files.
read_ct_spec_example()

# Read an example controlled terminology spec file.
read_ct_spec_example("ct-01-cm.csv")

# You may omit the file extension.
read_ct_spec_example("ct-01-cm")
```

---

read\_domain\_example    *Read an example SDTM domain*

---

## Description

[read\\_domain\\_example\(\)](#) imports one of the bundled SDTM domain examples as a [tibble](#) into R. See [domain\\_example\(\)](#) for possible choices.

## Usage

```
read_domain_example(example)
```

## Arguments

**example**            The name of SDTM domain example, e.g. "cm" (Concomitant Medication) or "ae" (Adverse Events). Run [read\\_domain\\_example\(\)](#) for available example files.

## Value

A [tibble](#) with an SDTM domain dataset, or a character vector of example file names.

## See Also

[domain\\_example\(\)](#)

## Examples

```
# Leave the `example` parameter as missing for available example files.
read_domain_example()

# Read the example Concomitant Medication domain.
read_domain_example("cm")

# Read the example Adverse Events domain.
read_domain_example("ae")
```

---

sbj_vars	<i>Subject-level key variables</i>
----------	------------------------------------

---

**Description**

`sbj_vars()` returns the set of variable names that uniquely define a subject.

**Usage**

```
sbj_vars()
```

**Value**

A character vector of variable names.

**Examples**

```
sbj_vars()
```

---

tbl_sum.cnd_df	<i>Conditioned tibble header print method</i>
----------------	---

---

**Description**

Conditioned tibble header print method. This S3 method adds an extra line in the header of a tibble that indicates the tibble is a conditioned tibble (`# Cond. tbl:`) followed by the tally of the conditioning vector: number of TRUE, FALSE and NA values: e.g., 1/1/1.

**Usage**

```
## S3 method for class 'cnd_df'
tbl_sum(x, ...)
```

**Arguments**

<code>x</code>	A conditioned tibble of class <code>cnd_df</code> .
<code>...</code>	Additional arguments passed to the default print method.

**Value**

A character vector with header values of the conditioned data frame.

**See Also**

`ctl_new_rowid_pillar.cnd_df()`.



**Examples**

```
df <- data.frame(x = c(1L, NA_integer_, 3L))
(cnd_df <- condition_add(dat = df, x >= 2L))
pillar::tbl_sum(cnd_df)
```

%.&gt;%

*Explicit Dot Pipe***Description****[Experimental]**

This operator pipes an object forward into a function or call expression using an explicit placement of the dot (.) placeholder. Unlike magrittr's `%>%` operator, `%.>%` does not automatically place the left-hand side (lhs) as the first argument in the right-hand side (rhs) call. This operator provides a simpler alternative to the use of braces with magrittr, while achieving similar behavior.

**Usage**

```
lhs %.>% rhs
```

**Arguments**

lhs	A value to be piped forward.
rhs	A function call that utilizes the dot (.) placeholder to specify where lhs should be placed.

**Details**

The `%.>%` operator is used to pipe the lhs value into the rhs function call. Within the rhs expression, the placeholder `.` represents the position where lhs will be inserted. This provides more control over where the lhs value appears in the rhs function call, compared to the magrittr pipe operator which always places lhs as the first argument of rhs.

Unlike magrittr's pipe, which may require the use of braces to fully control the placement of lhs in nested function calls, `%.>%` simplifies this by directly allowing multiple usages of the dot placeholder without requiring braces. For example, the following expression using magrittr's pipe and braces:

```
library(magrittr)

1:10 %>% { c(min(.), max(.)) }
```

can be written as:

```
1:10 %.>% c(min(.), max(.))
```

without needing additional braces.

**Downside:**

The disadvantage of `%.>%` is that you always need to use the dot placeholder, even when piping to the first argument of the right-hand side (rhs).

**Value**

No Return Value.

**Examples**

```
# Equivalent to `subset(head(iris), 1:nrow(head(iris)) %% 2 == 0)`  
head(iris) %.>% subset(., 1:nrow(.) %% 2 == 0)
```

```
# Equivalent to `c(min(1:10), max(1:10))`  
1:10 %.>% c(min(.), max(.))
```

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