

Package ‘teal.data’

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Description Provides a 'teal_data' class as a unified data model for 'teal' applications focusing on reproducibility and relational data.

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cdisc_data	<i>Data input for teal app</i>
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Description

[Stable]

Function is a wrapper around `teal_data()` and guesses `join_keys` for given datasets whose names match ADAM datasets names.

Usage

```
cdisc_data(
  ...,
  join_keys = teal.data::default_cdisc_join_keys[names(rlang::list2(...))],
  code = character(0),
  check
)
```

Arguments

<code>...</code>	any number of objects (presumably data objects) provided as name = value pairs.
<code>join_keys</code>	(<code>join_keys</code>) object or a single (<code>join_key_set</code>) object (optional) object with datasets column names used for joining. If empty then it would be automatically derived basing on intersection of datasets primary keys. For ADAM datasets it would be automatically derived.
<code>code</code>	(character, language) optional code to reproduce the datasets provided in <code>...</code> . Note this code is not executed and the <code>teal_data</code> may not be reproducible
<code>check</code>	(logical) [Deprecated] Use <code>verify()</code> to verify code reproducibility .

Details

This function checks if there were keys added to all data sets.

Value

A `teal_data` object.

Examples

```
data <- cdisc_data(
  join_keys = join_keys(
    join_key("ADSL", "ADTTE", c("STUDYID" = "STUDYID", "USUBJID" = "USUBJID"))
  )
)

data <- within(data, {
```

```
ADSL <- example_cdisc_data("ADSL")
ADTTE <- example_cdisc_data("ADTTE")
})
```

col_labels

Variable labels

Description

Get or set variable labels in a `data.frame`.

Usage

```
col_labels(x, fill = FALSE)
```

```
col_labels(x) <- value
```

```
col_relabel(x, ...)
```

```
get_labels(...)
```

Arguments

<code>x</code>	(<code>data.frame</code> or <code>DataFrame</code>) data object
<code>fill</code>	(<code>logical(1)</code>) specifying what to return if variable has no label
<code>value</code>	(<code>character</code>) vector of variable labels of length equal to number of columns in <code>x</code> ; if named, names must match variable names in <code>x</code> and will be used as key to set labels; use <code>NA</code> to remove label from variable
<code>...</code>	name-value pairs, where name corresponds to a variable name in <code>x</code> and value is the new variable label

Details

Variable labels can be stored as a label attribute set on individual variables. These functions get or set this attribute, either on all (`col_labels`) or some variables (`col_relabel`).

[Deprecated]

In previous versions of `teal.data` labels were managed with `get_labels()`. This function is deprecated as of `0.4.0`, use `col_labels` instead.

Value

For `col_labels`, named character vector of variable labels, the names being the corresponding variable names. If the label attribute is missing, the vector elements will be the variable names themselves if `fill = TRUE` and `NA` if `fill = FALSE`.

For `col_labels<-` and `col_relabel`, copy of `x` with variable labels modified.

Source

These functions were taken from `formatters` package, to reduce the complexity of the dependency tree and rewritten.

Examples

```
x <- iris
col_labels(x)
col_labels(x) <- paste("label for", names(iris))
col_labels(x)
y <- col_relabel(x, Sepal.Length = "Sepal Length of iris flower")
col_labels(y)
```

datanames	<i>Names of data sets in teal_data object</i>
-----------	---

Description

Get or set the value of the datanames slot.

Usage

```
datanames(x)

datanames(x) <- value
```

Arguments

x	(teal_data) object to access or modify
value	(character) new value for @datanames; all elements must be names of variables existing in @env

Details

The @datanames slot in a teal_data object specifies which of the variables stored in its environment (the @env slot) are data sets to be taken into consideration. The contents of @datanames can be specified upon creation and default to all variables in @env. Variables created later, which may well be data sets, are not automatically considered such. Use this function to update the slot.

Value

The contents of @datanames or teal_data object with updated @datanames.

Examples

```
td <- teal_data(iris = iris)
td <- within(td, mtcars <- mtcars)
datanames(td)

datanames(td) <- c("iris", "mtcars")
datanames(td)
```

default_cdisc_join_keys

List containing default joining keys for CDISC datasets

Description

This data object is created at loading time from `cdisc_datasets/cdisc_datasets.yaml`.

Usage

```
default_cdisc_join_keys
```

Format

An object of class `join_keys` (inherits from `list`) of length 19.

Source

internal

example_cdisc_data *Generate sample CDISC datasets*

Description

Retrieves example CDISC datasets for use in examples and testing.

Usage

```
example_cdisc_data(
  dataname = c("ADSL", "ADAE", "ADLB", "ADCM", "ADEX", "ADRS", "ADTR", "ADTTE", "ADVS")
)
```

Arguments

dataname (character(1)) name of a CDISC dataset

Details

This function returns a dummy dataset and should only be used within `teal.data`. Note that the datasets are not created and maintained in `teal.data`, they are retrieved its dependencies.

Value

A CDISC dataset as a `data.frame`.

get_code,teal_data-method

Get code from teal_data object

Description

Retrieve code from `teal_data` object.

Usage

```
## S4 method for signature 'teal_data'
get_code(object, deparse = TRUE, datanames = NULL, ...)
```

Arguments

<code>object</code>	(<code>teal_data</code>)
<code>deparse</code>	(logical) flag specifying whether to return code as character (<code>deparse = TRUE</code>) or as expression (<code>deparse = FALSE</code>).
<code>datanames</code>	[Experimental] (character) vector of dataset names to return the code for. For more details see the "Extracting dataset-specific code" section.
<code>...</code>	Parameters passed to internal methods. Currently, the only supported parameter is <code>check_names</code> (logical(1)) flag, which is <code>TRUE</code> by default. Function warns about missing objects, if they do not exist in code but are passed in <code>datanames</code> . To remove the warning, set <code>check_names = FALSE</code> .

Details

Retrieve code stored in `@code`, which (in principle) can be used to recreate all objects found in `@env`. Use `datanames` to limit the code to one or more of the datasets enumerated in `@datanames`. If the code has not passed verification (with `verify()`), a warning will be prepended.

Value

Either a character string or an expression. If `datanames` is used to request a specific dataset, only code that *creates* that dataset (not code that uses it) is returned. Otherwise, all contents of `@code`.

Extracting dataset-specific code

When `datanames` is specified, the code returned will be limited to the lines needed to *create* the requested datasets. The code stored in the `@code` slot is analyzed statically to determine which lines the datasets of interest depend upon. The analysis works well when objects are created with standard infix assignment operators (see `?assignOps`) but it can fail in some situations.

Consider the following examples:

Case 1: Usual assignments.

```
data <- teal_data() |>
  within({
    foo <- function(x) {
      x + 1
    }
    x <- 0
    y <- foo(x)
  })
get_code(data, datanames = "y")
```

`x` has no dependencies, so `get_code(data, datanames = "x")` will return only the second call. `y` depends on `x` and `foo`, so `get_code(data, datanames = "y")` will contain all three calls.

Case 2: Some objects are created by a function's side effects.

```
data <- teal_data() |>
  within({
    foo <- function() {
      x <<- x + 1
    }
    x <- 0
    foo()
    y <- x
  })
get_code(data, datanames = "y")
```

Here, `y` depends on `x` but `x` is modified by `foo` as a side effect (not by reassignment) and so `get_code(data, datanames = "y")` will not return the `foo()` call.

To overcome this limitation, code dependencies can be specified manually. Lines where side effects occur can be flagged by adding `"# @linksto <object name>"` at the end.

Note that `within` evaluates code passed to `expr` as is and comments are ignored. In order to include comments in code one must use the `eval_code` function instead.

```
data <- teal_data() |>
  eval_code("
    foo <- function() {
      x <<- x + 1
    }
    x <- 0
```



```

    foo() # @linksto x
    y <- x
  ")
get_code(data, datanames = "y")

```

Now the `foo()` call will be properly included in the code required to recreate `y`.

Note that two functions that create objects as side effects, `assign` and `data`, are handled automatically.

Here are known cases where manual tagging is necessary:

- non-standard assignment operators, *e.g.* `%<>%`
- objects used as conditions in `if` statements: `if (<condition>)`
- objects used to iterate over in `for` loops: `for(i in <sequence>)`
- creating and evaluating language objects, *e.g.* `eval(<call>)`

Examples

```

tdata1 <- teal_data()
tdata1 <- within(tdata1, {
  a <- 1
  b <- a^5
  c <- list(x = 2)
})
get_code(tdata1)
get_code(tdata1, datanames = "a")
get_code(tdata1, datanames = "b")

tdata2 <- teal_data(x1 = iris, code = "x1 <- iris")
get_code(tdata2)
get_code(verify(tdata2))

```

join_key

Create a relationship between a pair of datasets

Description

[Stable]

Create a relationship between two datasets, `dataset_1` and `dataset_2`. By default, this function establishes a directed relationship with `dataset_1` as the parent. If `dataset_2` is not specified, the function creates a primary key for `dataset_1`.

Usage

```
join_key(dataset_1, dataset_2 = dataset_1, keys, directed = TRUE)
```

Arguments

dataset_1, dataset_2	(character(1)) Dataset names. When dataset_2 is omitted, a primary key for dataset_1 is created.
keys	(optionally named character) Column mapping between the datasets, where names(keys) maps columns in dataset_1 corresponding to columns of dataset_2 given by the elements of keys. <ul style="list-style-type: none"> • If unnamed, the same column names are used for both datasets. • If any element of the keys vector is empty with a non-empty name, then the name is used for both datasets.
directed	(logical(1)) Flag that indicates whether it should create a parent-child relationship between the datasets. <ul style="list-style-type: none"> • TRUE (default) dataset_1 is the parent of dataset_2; • FALSE when the relationship is undirected.

Value

object of class `join_key_set` to be passed into `join_keys` function.

See Also

[join_keys\(\)](#), [parents\(\)](#)

Examples

```
join_key("d1", "d2", c("A"))
join_key("d1", "d2", c("A" = "B"))
join_key("d1", "d2", c("A" = "B", "C"))
```

join_keys

Manage relationships between datasets using join_keys

Description

Facilitates the creation and retrieval of relationships between datasets. `join_keys` class extends `list` and contains keys connecting pairs of datasets. Each element of the list contains keys for specific dataset. Each dataset can have a relationship with itself (primary key) and with other datasets.

Note that `join_keys` list is symmetrical and assumes a default direction, that is: when keys are set between `ds1` and `ds2`, it defines `ds1` as the parent in a parent-child relationship and the mapping is automatically mirrored between `ds2` and `ds1`.

Usage

```

## Constructor, getter and setter
join_keys(...)

## Default S3 method:
join_keys(...)

## S3 method for class 'join_keys'
join_keys(...)

## S3 method for class 'teal_data'
join_keys(...)

## S3 method for class 'join_keys'
x[i, j]

## S3 replacement method for class 'join_keys'
x[i, j, directed = TRUE] <- value

## S3 method for class 'join_keys'
c(...)

## S3 method for class 'join_key_set'
c(...)

join_keys(x) <- value

## S3 replacement method for class 'join_keys'
join_keys(x) <- value

## S3 replacement method for class 'teal_data'
join_keys(x) <- value

## S3 method for class 'join_keys'
format(x, ...)

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

```

Arguments

```

...          (optional)
             • either teal_data or join_keys object to extract join_keys
             • or any number of join_key_set objects to create join_keys
             • or nothing to create an empty join_keys
x            (join_keys) empty object to set the new relationship pairs. x is typically an ob-
             ject of join_keys class. When called with the join_keys(x) or join_keys(x)
             <- value then it can also take a supported class (teal_data, join_keys)

```

<code>i, j</code>	indices specifying elements to extract or replace. Index should be a character vector, but it can also take numeric, logical, NULL or missing.
<code>directed</code>	(logical(1)) Flag that indicates whether it should create a parent-child relationship between the datasets. <ul style="list-style-type: none"> • TRUE (default) dataset_1 is the parent of dataset_2; • FALSE when the relationship is undirected.
<code>value</code>	For <code>x[i, j, directed = TRUE]</code> <- value (named/unnamed character) Column mapping between datasets. For <code>join_keys(x) <- value</code> : (join_key_set or list of join_key_set) relationship pairs to add to join_keys list. <code>[i, j, directed = TRUE]</code> : <code>R:i,%20j,%20directed%20=%20TRUE)</code>

Value

join_keys object.

Methods (by class)

- `join_keys()`: Returns an empty join_keys object when called without arguments.
- `join_keys(join_keys)`: Returns itself.
- `join_keys(teal_data)`: Returns the join_keys object contained in teal_data object.
- `join_keys(...)`: Creates a new object with one or more join_key_set parameters.

Functions

- `x[datanames]`: Returns a subset of the join_keys object for given datanames, including parent datanames and symmetric mirror keys between datanames in the result.
- `x[i, j]`: Returns join keys between datasets i and j, including implicit keys inferred from their relationship with a parent.
- `x[i, j] <- value`: Assignment of a key to pair (i, j).
- `x[i] <- value`: This (without j parameter) **is not** a supported operation for join_keys.
- `join_keys(x)[i, j] <- value`: Assignment to join_keys object stored in x, such as a teal_data object or join_keys object itself.
- `join_keys(x) <- value`: Assignment of the join_keys in object with value. value needs to be an object of class join_keys or join_key_set.

See Also

[join_key\(\)](#) for creating join_key_set, [parents\(\)](#) for parent operations, [teal_data\(\)](#) for teal_data constructor and [default_cdisc_join_keys](#) for default CDISC keys.

Examples

```

# Creating a new join keys ----

jk <- join_keys(
  join_key("ds1", "ds1", "pk1"),
  join_key("ds2", "ds2", "pk2"),
  join_key("ds3", "ds3", "pk3"),
  join_key("ds1", "ds2", c(pk1 = "pk2")),
  join_key("ds1", "ds3", c(pk1 = "pk3"))
)

jk

# Getter for join_keys ---

jk["ds1", "ds2"]

# Subsetting join_keys ----

jk["ds1"]
jk[1:2]
jk[c("ds1", "ds2")]

# Setting a new primary key ---

jk["ds4", "ds4"] <- "pk4"
jk["ds5", "ds5"] <- "pk5"

# Setting a single relationship pair ---

jk["ds1", "ds4"] <- c("pk1" = "pk4")

# Removing a key ---

jk["ds5", "ds5"] <- NULL
# Merging multiple `join_keys` objects ---

jk_merged <- c(
  jk,
  join_keys(
    join_key("ds4", keys = c("pk4", "pk4_2")),
    join_key("ds3", "ds4", c(pk3 = "pk4_2"))
  )
)
# note: merge can be performed with both join_keys and join_key_set

jk_merged <- c(
  jk_merged,
  join_key("ds5", keys = "pk5"),
  join_key("ds1", "ds5", c(pk1 = "pk5"))
)

```

```

)

# Assigning keys via join_keys(x)[i, j] <- value ----

obj <- join_keys()
# or
obj <- teal_data()

join_keys(obj)["ds1", "ds1"] <- "pk1"
join_keys(obj)["ds2", "ds2"] <- "pk2"
join_keys(obj)["ds3", "ds3"] <- "pk3"
join_keys(obj)["ds1", "ds2"] <- c(pk1 = "pk2")
join_keys(obj)["ds1", "ds3"] <- c(pk1 = "pk3")

identical(jk, join_keys(obj))

# Setter for join_keys within teal_data ----

td <- teal_data()
join_keys(td) <- jk

join_keys(td)["ds1", "ds2"] <- "new_key"
join_keys(td) <- c(join_keys(td), join_keys(join_key("ds3", "ds2", "key3")))
join_keys(td)

```

names<-join_keys *The names of a join_keys object*

Description

The names of a join_keys object

Usage

```
## S3 replacement method for class 'join_keys'
names(x) <- value
```

Arguments

x an R object.

value a character vector of up to the same length as x, or NULL.

parents

Get and set parents in join_keys object

Description

parents() facilitates the creation of dependencies between datasets by assigning a parent-child relationship.

Usage

```
parents(x)

## S3 method for class 'join_keys'
parents(x)

## S3 method for class 'teal_data'
parents(x)

parents(x) <- value

## S3 replacement method for class 'join_keys'
parents(x) <- value

## S3 replacement method for class 'teal_data'
parents(x) <- value

parent(x, dataset_name)
```

Arguments

x	(join_keys or teal_data) object that contains "parents" information to retrieve or manipulate.
value	(named list) of character vectors.
dataset_name	(character(1)) Name of dataset to query on their parent.

Details

Each element is defined by a list element, where list("child" = "parent").

Value

a list of character representing the parents.

For parent(x, dataset_name) returns NULL if parent does not exist.

Methods (by class)

- `parents(join_keys)`: Retrieves parents of `join_keys` object.
- `parents(teal_data)`: Retrieves parents of `join_keys` inside `teal_data` object.

Functions

- `parents(x) <- value`: Assignment of parents in `join_keys` object.
- `parents(join_keys) <- value`: Assignment of parents of `join_keys` object.
- `parents(teal_data) <- value`: Assignment of parents of `join_keys` inside `teal_data` object.
- `parent()`: Getter for individual parent.

See Also

[join_keys\(\)](#)

Examples

```
# Get parents of join_keys ---

jk <- default_cdisc_join_keys["ADEX"]
parents(jk)

# Get parents of join_keys inside teal_data object ---

td <- teal_data(
  ADSL = rADSL,
  ADTTE = rADTTE,
  ADRS = rADRS,
  join_keys = default_cdisc_join_keys[c("ADSL", "ADTTE", "ADRS")]
)
parents(td)

# Assignment of parents ---

jk <- join_keys(
  join_key("ds1", "ds2", "id"),
  join_key("ds5", "ds6", "id"),
  join_key("ds7", "ds6", "id")
)

parents(jk) <- list(ds2 = "ds1")

# Setting individual parent-child relationship

parents(jk)["ds6"] <- "ds5"
parents(jk)["ds7"] <- "ds6"

# Assignment of parents of join_keys inside teal_data object ---
```



```
parents(td) <- list("ADTTE" = "ADSL") # replace existing
parents(td)["ADRS"] <- "ADSL" # add new parent

# Get individual parent ---

parent(jk, "ds2")
parent(td, "ADTTE")
```

show,teal_data-method *Show teal_data object*

Description

Prints teal_data object.

Usage

```
## S4 method for signature 'teal_data'
show(object)
```

Arguments

object (teal_data)

Value

Input teal_data object.

Examples

```
teal_data()
teal_data(x = iris, code = "x = iris")
verify(teal_data(x = iris, code = "x = iris"))
```

TealData

Deprecated TealData class and related functions

Description

[Deprecated]

The TealData class and associated functions have been deprecated. Use `teal_data()` instead. See the [Migration guide](#) for details.

Usage

```
as_cdisc(...)  
callable_code(...)  
callable_function(...)  
code_dataset_connector(...)  
code_cdisc_dataset_connector(...)  
csv_dataset_connector(...)  
csv_cdisc_dataset_connector(...)  
python_code(...)  
python_dataset_connector(...)  
python_cdisc_dataset_connector(...)  
cdisc_data_connector(...)  
cdisc_dataset(...)  
cdisc_dataset_connector(...)  
cdisc_dataset_connector_file(...)  
cdisc_dataset_file(...)  
dataset(...)  
dataset_connector(...)  
dataset_connector_file(...)  
dataset_file(...)  
data_connection(...)  
fun_dataset_connector(...)  
fun_cdisc_dataset_connector(...)  
relational_data_connector(...)  
mae_dataset(...)
```

```
get_attrs(...)  
get_dataset_label(...)  
get_dataset(...)  
get_datasets(...)  
get_dataname(...)  
get_key_duplicates(...)  
get_keys(...)  
get_raw_data(...)  
is_pulled(...)  
load_dataset(...)  
load_datasets(...)  
mutate_data(...)  
mutate_dataset(...)  
set_args(...)  
rds_dataset_connector(...)  
rds_cdisc_dataset_connector(...)  
script_dataset_connector(...)  
script_cdisc_dataset_connector(...)  
set_keys(...)  
read_script(...)  
to_relational_data(...)  
validate_metadata(...)  
get_cdisc_keys(...)  
cdisc_data_file(...)
```

```
teal_data_file(...)
get_join_keys(...)
get_join_keys(...) <- value
```

Arguments

... any argument supported in TealData related functions.
 value value to assign

Value

nothing

See Also

[cdisc_data\(\)](#) , [join_keys\(\)](#)

 teal_data

Comprehensive data integration function for teal applications

Description

[Stable]

Universal function to pass data to teal application.

Usage

```
teal_data(..., join_keys = teal.data::join_keys(), code = character(0), check)
```

Arguments

... any number of objects (presumably data objects) provided as name = value pairs.
 join_keys (join_keys) object or a single (join_key_set) object.
 (optional) object with dataset column relationships used for joining. If empty then no joins between pairs of objects.
 code (character, language) optional code to reproduce the datasets provided in ...
 Note this code is not executed and the teal_data may not be reproducible
 check (logical) **[Deprecated]** Use [verify\(\)](#) to verify code reproducibility .

Value

A teal_data object.

Examples

```
teal_data(x1 = iris, x2 = mtcars)
```

verify

Verify code reproducibility

Description

Checks whether code in `teal_data` object reproduces the stored objects.

Usage

```
verify(x)
```

Arguments

`x` teal_data object

Details

If objects created by code in the `@code` slot of `x` are `all_equal` to the contents of the `@env` slot, the function updates the `@verified` slot to `TRUE` in the returned `teal_data` object. Once verified, the slot will always be set to `TRUE`. If the `@code` fails to recreate objects in `teal_data@env`, an error is raised.

Value

Input `teal_data` object or error.

Examples

```
tdata1 <- teal_data()
tdata1 <- within(tdata1, {
  a <- 1
  b <- a^5
  c <- list(x = 2)
})
verify(tdata1)

tdata2 <- teal_data(x1 = iris, code = "x1 <- iris")
verify(tdata2)
verify(tdata2)@verified
tdata2@verified

tdata3 <- teal_data()
tdata3 <- within(tdata3, {
  stop("error")
})
```

```
try(verify(tdata3)) # fails

a <- 1
b <- a + 2
c <- list(x = 2)
d <- 5
tdata4 <- teal_data(
  a = a, b = b, c = c, d = d,
  code = "a <- 1
         b <- a
         c <- list(x = 2)
         e <- 1"
)
tdata4
try(verify(tdata4)) # fails
```

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