

Package ‘reply’

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Type Package

Title Patches to Use 'dplyr' on Remote Data Sources

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URL <https://github.com/WinVector/reply/>,
<https://winvector.github.io/reply/>

BugReports <https://github.com/WinVector/reply/issues>

Description Patches to use 'dplyr' on remote data sources ('SQL' databases, 'Spark' 2.0.0 and above) in a reliable ``generic" fashion (generic meaning user code works similarly on all such sources, without needing per-source adaption). Due to the fluctuating nature of 'dplyr'/'dbplyr'/'rlang' APIs this package is going into maintenance mode. Most of the 'reply' functions are already done better by one of the non-monolithic replacement packages: 'wrapr', 'seplyr', 'rquery', or 'cdata'.

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LazyData TRUE

Depends R (>= 3.4.0), wrapr (>= 1.8.5)

Imports dplyr (>= 0.7.0), rlang (>= 0.2.0), DBI

RoxygenNote 6.1.1

Suggests testthat, knitr, rmarkdown, sparklyr, igraph, DiagrammeR, RSQLite

VignetteBuilder knitr

ByteCompile true

NeedsCompilation no

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<code>addConstantColumn</code>	<i>Add constant to a table.</i>
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Description

Work around different treatment of character types across remote data sources when adding a constant column to a table. Deals with issues such as Postgresql requiring a character-cast and MySQL not allowing such.

Usage

```
addConstantColumn(d, colName, val, ...,
  tempNameGenerator = mk_tmp_name_source("replyr_addConstantColumn"))
```

Arguments

<code>d</code>	data.frame like object to add column to.
<code>colName</code>	character, name of column to add.
<code>val</code>	scalar, value to add.
<code>...</code>	force later arguments to be bound by name.
<code>tempNameGenerator</code>	temp name generator produced by <code>wrapr::mk_tmp_name_source</code> , used to record <code>dplyr::compute()</code> effects.

Value

table with new column added.

Examples

```
d <- data.frame(x= c(1:3))
addConstantColumn(d, 'newCol', 'newVal')
```

buildJoinPlan	<i>Build a join plan</i>
---------------	--------------------------

Description

Please see `vignette('DependencySorting', package = 'replyr')` and `vignette('joinController', package = 'replyr')` for more details.

Usage

```
buildJoinPlan(tDesc, ..., check = TRUE)
```

Arguments

tDesc	description of tables from tableDescription (and likely altered by user). Note: no column names must intersect with names of the form <code>table_CLEANEDTABNAME_present</code> .
...	force later arguments to bind by name.
check	logical, if TRUE check the join plan for consistency.

Value

detailed column join plan (appropriate for editing)

See Also

[tableDescription](#), [inspectDescrAndJoinPlan](#), [makeJoinDiagramSpec](#), [executeLeftJoinPlan](#)

Examples

```
d <- data.frame(id=1:3, weight= c(200, 140, 98))
tDesc <- rbind(tableDescription('d1', d),
              tableDescription('d2', d))
tDesc$keys[[1]] <- list(PrimaryKey= 'id')
tDesc$keys[[2]] <- list(PrimaryKey= 'id')
buildJoinPlan(tDesc)
```

`dplyr_src_to_db_handle`

Obsolete with dplyr 0.7.0 and forward

Description

Obsolete with dplyr 0.7.0 and forward

Usage

`dplyr_src_to_db_handle(dplyr_src)`

Arguments

`dplyr_src` remote data handle

Value

`dplyr_src`

`example_employeeAndDate`

build some example tables

Description

build some example tables

Usage

`example_employeeAndDate(con)`

Arguments

`con` db connection

Value

example tables

executeLeftJoinPlan *Execute an ordered sequence of left joins.*

Description

Please see vignette('DependencySorting', package = 'replyr') and vignette('joinController', package= 're' for more details.

Usage

```
executeLeftJoinPlan(tDesc, columnJoinPlan, ..., checkColumns = FALSE,
  computeFn = function(x, name) { dplyr::compute(x, name = name) },
  eagerCompute = TRUE, checkColClasses = FALSE, verbose = FALSE,
  dryRun = FALSE,
  tempNameGenerator = mk_tmp_name_source("executeLeftJoinPlan"))
```

Arguments

tDesc	description of tables, either a data.frame from tableDescription , or a list mapping from names to handles/frames. Only used to map table names to data.
columnJoinPlan	columns to join, from buildJoinPlan (and likely altered by user). Note: no column names must intersect with names of the form table_CLEANEDTABNAME_present.
...	force later arguments to bind by name.
checkColumns	logical if TRUE confirm column names before starting joins.
computeFn	function to call to try and materialize intermediate results.
eagerCompute	logical if TRUE materialize intermediate results with computeFn.
checkColClasses	logical if true check for exact class name matches
verbose	logical if TRUE print more.
dryRun	logical if TRUE do not perform joins, only print steps.
tempNameGenerator	temp name generator produced by wrapr::mk_tmp_name_source, used to record dplyr::compute() effects.

Value

joined table

See Also

[tableDescription](#), [buildJoinPlan](#), [inspectDescrAndJoinPlan](#), [makeJoinDiagramSpec](#)

Examples

```

# example data
meas1 <- data.frame(id= c(1,2),
                    weight= c(200, 120),
                    height= c(60, 14))
meas2 <- data.frame(pid= c(2,3),
                    weight= c(105, 110),
                    width= 1)
# get the initial description of table defs
tDesc <- rbind(tableDescription('meas1', meas1),
              tableDescription('meas2', meas2))
# declare keys (and give them consistent names)
tDesc$keys[[1]] <- list(PatientID= 'id')
tDesc$keys[[2]] <- list(PatientID= 'pid')
# build the column join plan
columnJoinPlan <- buildJoinPlan(tDesc)
# decide we don't want the width column
columnJoinPlan$want[columnJoinPlan$resultColumn=='width'] <- FALSE
# double check our plan
if(!is.null(inspectDescrAndJoinPlan(tDesc, columnJoinPlan,
                                   checkColClasses= TRUE))) {
  stop("bad join plan")
}
# execute the left joins
executeLeftJoinPlan(tDesc, columnJoinPlan,
                   checkColClasses= TRUE,
                   verbose= TRUE)

# also good
executeLeftJoinPlan(list('meas1'=meas1, 'meas2'=meas2),
                   columnJoinPlan,
                   checkColClasses= TRUE,
                   verbose= TRUE)

```

expandColumn

Expand a column of vectors into one row per value of each vector.

Description

Similar to `tidyr::unnest` but lands rowids and value ids, and can work on remote data sources. Fairly expensive per-row operation, not suitable for big data.

Usage

```

expandColumn(data, colName, ..., rowidSource = NULL, rowidDest = NULL,
            idxDest = NULL,
            tempNameGenerator = mk_tmp_name_source("replyr_expandColumn"))

```

Arguments

<code>data</code>	data.frame to work with.
<code>colName</code>	character name of column to expand.
<code>...</code>	force later arguments to be bound by name
<code>rowidSource</code>	optional character name of column to take row indices from (rowidDest must be NULL to use this).
<code>rowidDest</code>	optional character name of column to write row indices to (must not be an existing column name, rowidSource must be NULL to use this).
<code>idxDest</code>	optional character name of column to write value indices to (must not be an existing column name).
<code>tempNameGenerator</code>	temp name generator produced by <code>wrpr::mk_tmp_name_source</code> , used to record <code>dplyr::compute()</code> effects.

Value

expanded data frame where each value of `colName` column is in a new row.

Examples

```
d <- data.frame(name= c('a','b'))
d$value <- list(c('x','y'),'z')
expandColumn(d, 'value',
             rowidDest= 'origRowId',
             idxDest= 'valueIndex')
```

gapply

grouped ordered apply

Description

Partitions from by values in grouping column, applies a generic transform to each group and then binds the groups back together. Only advised for a moderate number of groups and better if grouping column is an index. This is powerful enough to implement "The Split-Apply-Combine Strategy for Data Analysis" <https://www.jstatsoft.org/article/view/v040i01>

Usage

```
gapply(df, gcolumn, f, ..., ocolumn = NULL, decreasing = FALSE,
       partitionMethod = "split", bindrows = TRUE, maxgroups = 100,
       eagerCompute = FALSE, restoreGroup = FALSE,
       tempNameGenerator = mk_tmp_name_source("replyr_gapply"))
```


Arguments

df	remote dplyr data item
gcolumn	grouping column
f	transform function or pipeline
...	force later values to be bound by name
ocolumn	ordering column (optional)
decreasing	logical, if TRUE sort in decreasing order by ocolumn
partitionMethod	method to partition the data, one of 'group_by' (depends on f being dplyr compatible), 'split' (only works over local data frames), or 'extract'
bindrows	logical, if TRUE bind the rows back into a data item, else return split list
maxgroups	maximum number of groups to work over (intentionally not enforced if partitionMethod=='group_by')
eagerCompute	logical, if TRUE call compute on split results
restoreGroup	logical, if TRUE restore group column after apply when partitionMethod %in% c('extract', 'split')
tempNameGenerator	temp name generator produced by wrapr::mk_tmp_name_source, used to record dplyr::compute() effects.

Details

Note this is a fairly expensive operator, so it only makes sense to use in situations where f itself is fairly complicated and/or expensive.

Value

transformed frame

Examples

```
d <- data.frame(
  group = c(1, 1, 2, 2, 2),
  order = c(.1, .2, .3, .4, .5),
  values = c(10, 20, 2, 4, 8)
)

# User supplied window functions. They depend on known column names and
# the data back-end matching function names (as cumsum).
cumulative_sum <- function(d) {
  dplyr::mutate(d, cv = cumsum(values))
}
rank_in_group <- function(d) {
  d %>%
    dplyr::mutate(., constcol = 1) %>%
    dplyr::mutate(., rank = cumsum(constcol)) %>%
    dplyr::select(., -constcol)
}
```

```

for (partitionMethod in c('group_by', 'split', 'extract')) {
  print(partitionMethod)
  print('cumulative sum example')
  print(
    gapply(
      d,
      'group',
      cumulative_sum,
      ocolumn = 'order',
      partitionMethod = partitionMethod
    )
  )
  print('ranking example')
  print(
    gapply(
      d,
      'group',
      rank_in_group,
      ocolumn = 'order',
      partitionMethod = partitionMethod
    )
  )
  print('ranking example (decreasing)')
  print(
    gapply(
      d,
      'group',
      rank_in_group,
      ocolumn = 'order',
      decreasing = TRUE,
      partitionMethod = partitionMethod
    )
  )
}

```

inspectDescrAndJoinPlan

check that a join plan is consistent with table descriptions

Description

Please see vignette('DependencySorting', package = 'replyr') and vignette('joinController', package = 'replyr') for more details.

Usage

```
inspectDescrAndJoinPlan(tDesc, columnJoinPlan, ...,
  checkColClasses = FALSE)
```

Arguments

tDesc description of tables, from [tableDescription](#) (and likely altered by user).

columnJoinPlan columns to join, from [buildJoinPlan](#) (and likely altered by user). Note: no column names must intersect with names of the form table_CLEANEDTABNAME_present.

... force later arguments to bind by name.

checkColClasses logical if true check for exact class name matches

Value

NULL if okay, else a string

See Also

[tableDescription](#), [buildJoinPlan](#), [makeJoinDiagramSpec](#), [executeLeftJoinPlan](#)

Examples

```
# example data
d1 <- data.frame(id= 1:3,
                 weight= c(200, 140, 98),
                 height= c(60, 24, 12))
d2 <- data.frame(pid= 2:3,
                 weight= c(130, 110),
                 width= 1)
# get the initial description of table defs
tDesc <- rbind(tableDescription('d1', d1),
              tableDescription('d2', d2))
# declare keys (and give them consistent names)
tDesc$keys[[1]] <- list(PrimaryKey= 'id')
tDesc$keys[[2]] <- list(PrimaryKey= 'pid')
# build the join plan
columnJoinPlan <- buildJoinPlan(tDesc)
# confirm the plan
inspectDescrAndJoinPlan(tDesc, columnJoinPlan,
                       checkColClasses= TRUE)
# damage the plan
columnJoinPlan$sourceColumn[columnJoinPlan$sourceColumn=='width'] <- 'wd'
# find a problem
inspectDescrAndJoinPlan(tDesc, columnJoinPlan,
                       checkColClasses= TRUE)
```

keysAreUnique *Check uniqueness of rows with respect to keys.*

Description

Can be an expensive operation.

Usage

```
keysAreUnique(tDesc)
```

Arguments

tDesc description of tables, from [tableDescription](#) (and likely altered by user).

Value

logical TRUE if keys are unique

See Also

[tableDescription](#)

Examples

```
d <- data.frame(x=c(1,1,2,2,3,3), y=c(1,2,1,2,1,2))
tDesc1 <- tableDescription('d1', d)
tDesc2 <- tableDescription('d2', d)
tDesc <- rbind(tDesc1, tDesc2)
tDesc$keys[[2]] <- c(x='x')
keysAreUnique(tDesc)
```

key_inspector_all_cols *Return all columns as guess at preferred primary keys.*

Description

Return all columns as guess at preferred primary keys.

Usage

```
key_inspector_all_cols(handle)
```

Arguments

handle data handle

Value

map of keys to keys

See Also

tableDescription

Examples

```
d <- data.frame(x=1:3, y=NA)
key_inspector_all_cols(d)
```

key_inspector_postgresql

Return all primary key columns as guess at preferred primary keys for a PostgreSQL handle.

Description

Return all primary key columns as guess at preferred primary keys for a PostgreSQL handle.

Usage

```
key_inspector_postgresql(handle)
```

Arguments

handle data handle

Value

map of keys to keys

See Also

tableDescription

`key_inspector_sqlite` *Return all primary key columns as guess at preferred primary keys for a SQLite handle.*

Description

Return all primary key columns as guess at preferred primary keys for a SQLite handle.

Usage

```
key_inspector_sqlite(handle)
```

Arguments

`handle` data handle

Value

map of keys to keys

See Also

`tableDescription`

`makeJoinDiagramSpec` *Build a drawable specification of the join diagram*

Description

Please see `vignette('DependencySorting', package = 'replyr')` and `vignette('joinController', package = 'replyr')` for more details.

Usage

```
makeJoinDiagramSpec(columnJoinPlan, ..., groupByKeys = TRUE,
  graphOpts = NULL)
```

Arguments

`columnJoinPlan` join plan
`...` force later arguments to bind by name
`groupByKeys` logical if true build key-equivalent sub-graphs
`graphOpts` options for `graphViz`

Value

grViz diagram spec

See Also

[tableDescription](#), [buildJoinPlan](#), [executeLeftJoinPlan](#)

Examples

```
if (requireNamespace("RSQLite", quietly = TRUE)) {
  # note: employeeanddate is likely built as a cross-product
  #   join of an employee table and set of dates of interest
  #   before getting to the join controller step. We call
  #   such a table "row control" or "experimental design."
  my_db <- DBI::dbConnect(RSQLite::SQLite(), ":memory:")
  RSQLite::initExtension(my_db)
  tDesc <- example_employeeAndDate(my_db)
  # fix order by hand, please see replyr::topoSortTables for
  # how to automate this.
  ord <- match(c('employeeanddate', 'orgtable', 'activity', 'revenue'),
              tDesc$tableName)
  tDesc <- tDesc[ord, , drop=FALSE]
  columnJoinPlan <- buildJoinPlan(tDesc, check= FALSE)
  # unify keys
  columnJoinPlan$resultColumn[columnJoinPlan$resultColumn=='id'] <- 'eid'
  # look at plan defects
  print(paste('problems:',
              inspectDescrAndJoinPlan(tDesc, columnJoinPlan)))
  diagramSpec <- makeJoinDiagramSpec(columnJoinPlan)
  # to render as JavaScript:
  #   DiagrammeR::grViz(diagramSpec)
  DBI::dbDisconnect(my_db)
  my_db <- NULL
}
```

Description

Methods to reliably use dplyr on remote data sources in R (SQL databases, Spark 2.0.0 and above) in a generic fashion.

Details

replyr is going into maintenance mode. It has been hard to track shifting dplyr/dbplyr/rlang APIs and data structures post dplyr 0.5. Most of what it does is now done better in one of the newer non-monolithic packages:

- Programming and meta-programming tools: wrapr <https://CRAN.R-project.org/package=wrapr>.
- Adapting dplyr to standard evaluation interfaces: seplyr <https://CRAN.R-project.org/package=seplyr>.
- Big data data manipulation: rquery <https://CRAN.R-project.org/package=rquery> and cdata <https://CRAN.R-project.org/package=cdata>.

replyr helps with the following:

- Summarizing remote data (via replyr_summarize).
- Facilitating writing "source generic" code that works similarly on multiple 'dplyr' data sources.
- Providing big data versions of functions for splitting data, binding rows, pivoting, adding row-ids, ranking, and completing experimental designs.
- Packaging common data manipulation tasks into operators such as the `gapply` function.
- Providing support code for common SparklyR tasks, such as tracking temporary handle IDs.

replyr is in maintenance mode. Better version of the functionality have been ported to the following packages: wrapr, cdata, rquery, and seplyr.

To learn more about replyr, please start with the vignette: `vignette('replyr', 'replyr')`

replyr_add_ids

Add unique ids to rows. Note: re-arranges rows in many cases.

Description

Add unique ids to rows. Note: re-arranges rows in many cases.

Usage

```
replyr_add_ids(df, idColName, env = parent.frame(),
  local_short_cut = TRUE)
```

Arguments

df	data.frame object to work with
idColName	name of column to add
env	environment to evaluate in (not used).
local_short_cut	logical, if TRUE use base R on local data.

Examples

```
replyr_add_ids(data.frame(x=c('a','b')), 'id', local_short_cut = FALSE)
```

```
replyr_apply_f_mapped Apply a function to a re-mapped data frame.
```

Description

Apply a function to a re-mapped data frame.

Usage

```
replyr_apply_f_mapped(d, f, nmap, ..., restrictMapIn = FALSE,
  rmap = replyr::replyr_reverseMap(nmap), restrictMapOut = FALSE)
```

Arguments

d	data.frame to work on
f	function to apply.
nmap	named list mapping with keys specifying new column names, and values as original column names.
...	force later arguments to bind by name
restrictMapIn	logical if TRUE restrict columns when mapping in.
rmap	reverse map (for after f is applied).
restrictMapOut	logical if TRUE restrict columns when mapping out.

See Also

[let](#), [replyr_reverseMap](#), [replyr_mapRestrictCols](#)

Examples

```
# an external function with hard-coded column names
DecreaseRankColumnByOne <- function(d) {
  d$RankColumn <- d$RankColumn - 1
  d
}

# our example data, with different column names
d <- data.frame(Sepal_Length=c(5.8,5.7),
  Sepal_Width=c(4.0,4.4),
  Species='setosa',rank=c(1,2))

print(d)
```

```
# map our data to expected column names so we can use function
nmap <- c(GroupColumn='Species',
          ValueColumn='Sepal_Length',
          RankColumn='rank')
print(nmap)

dF <- replyr_apply_f_mapped(d, DecreaseRankColumnByOne, nmap)
print(dF)
```

replyr_arrange *arrange by a single column*

Description

arrange by a single column

Usage

```
replyr_arrange(.data, colname, descending = FALSE)
```

Arguments

.data	data object to work on
colname	character column name
descending	logical if true sort descending (else sort ascending)

Examples

```
d <- data.frame(Sepal_Length= c(5.8,5.7),
               Sepal_Width= c(4.0,4.4))
replyr_arrange(d, 'Sepal_Length', descending= TRUE)
```

replayr_bind_rows	<i>Bind a list of items by rows (can't use dplyr::bind_rows or dplyr::combine on remote sources). Columns are intersected.</i>
-------------------	--

Description

Bind a list of items by rows (can't use dplyr::bind_rows or dplyr::combine on remote sources). Columns are intersected.

Usage

```
replayr_bind_rows(lst, ..., useDplyrLocal = TRUE, useSparkRbind = TRUE,
  useUnionALL = TRUE,
  tempNameGenerator = mk_tmp_name_source("replayr_bind_rows"))
```

Arguments

lst	list of items to combine, must be all in same dplyr data service
...	force other arguments to be used by name
useDplyrLocal	logical if TRUE use dplyr for local data.
useSparkRbind	logical if TRUE try to use rbind on Sparklyr data
useUnionALL	logical if TRUE try to use union all binding
tempNameGenerator	temp name generator produced by wrapr::mk_tmp_name_source, used to record dplyr::compute() effects.

Value

single data item

Examples

```
if (requireNamespace("RSQLite", quietly = TRUE)) {
  my_db <- DBI::dbConnect(RSQLite::SQLite(), ":memory:")
  # my_db <- sparklyr::spark_connect(master = "local")
  d <- replayr_copy_to(my_db, data.frame(x = 1:2), 'd',
    temporary = TRUE)
  # dplyr::bind_rows(list(d, d))
  # # Argument 1 must be a data frame or a named atomic vector,
  # # not a tbl_dbi/tbl_sql/tbl_lazy/tbl
  print(replayr_bind_rows(list(d, d)))
  DBI::dbDisconnect(my_db)
}
```

replyr_check_ranks *confirm data has good ranked groups*

Description

confirm data has good ranked groups

Usage

```
replyr_check_ranks(x, GroupColumnName, ValueColumnName, RankColumnName,
  ..., decreasing = FALSE,
  tempNameGenerator = mk_tmp_name_source("replyr_check_ranks"))
```

Arguments

x data item to work with

GroupColumnName column to group by

ValueColumnName column determining order

RankColumnName column having proposed rank (function of order)

... force later arguments to bind by name

decreasing if true make order decreasing instead of increasing.

tempNameGenerator temp name generator produced by wrapr::mk_tmp_name_source, used to record dplyr::compute() effects.

Value

summary of quality of ranking.

Examples

```
d <- data.frame(Sepal_Length=c(5.8,5.7),Sepal_Width=c(4.0,4.4),
  Species='setosa',rank=c(1,2))
replyr_check_ranks(d, 'Species', 'Sepal_Length', 'rank', decreasing=TRUE)
```

replyr_coalesce	<i>Augment a data frame by adding additional rows.</i>
-----------------	--

Description

Note: do not count on order of resulting data. Also only added rows are altered by the fill instructions.

Usage

```
replyr_coalesce(data, support, ..., fills = NULL, newRowColumn = NULL,
  copy = TRUE,
  tempNameGenerator = mk_tmp_name_source("replyr_coalesce"))
```

Arguments

data	data.frame data to augment
support	data.frame rows of unique key-values into data
...	not used, force later arguments to bind by name
fills	list default values to fill in columns
newRowColumn	character if not null name to use for new row indicator
copy	logical if TRUE copy support to data's source
tempNameGenerator	temp name generator produced by wrapr::mk_tmp_name_source, used to record dplyr::compute() effects.

Value

augmented data

Examples

```
# single column key example
data <- data.frame(year = c(2005,2007,2010),
  count = c(6,1,NA),
  name = c('a','b','c'),
  stringsAsFactors = FALSE)
support <- data.frame(year=2005:2010)
filled <- replyr_coalesce(data, support,
  fills=list(count=0))
filled <- filled[order(filled$year), ]
filled

# complex key example
data <- data.frame(year = c(2005,2007,2010),
```

```
count = c(6,1,NA),
name = c('a', 'b', 'c'),
stringsAsFactors = FALSE)
support <- expand.grid(year=2005:2010,
name= c('a', 'b', 'c', 'd'),
stringsAsFactors = FALSE)
filled <- replyr_coalesce(data, support,
fills=list(count=0))
filled <- filled[order(filled$year, filled$name), ]
filled
```

replyr_colClasses *Get column classes.*

Description

Get column classes.

Usage

```
replyr_colClasses(x)
```

Arguments

x tbl or item that can be coerced into such.

Value

list of column classes.

Examples

```
d <- data.frame(x=c(1,2))
replyr_colClasses(d)
```

replyr_copy_from	<i>Bring remote data back as a local data frame tbl.</i>
------------------	--

Description

Bring remote data back as a local data frame tbl.

Usage

```
replyr_copy_from(d, maxrow = 1e+06)
```

Arguments

d	remote dplyr data item
maxrow	max rows to allow (stop otherwise, set to NULL to allow any size).

Value

local tbl.

Examples

```
if (requireNamespace("RSQLite", quietly = TRUE)) {
  my_db <- DBI::dbConnect(RSQLite::SQLite(), ":memory:")
  RSQLite::initExtension(my_db)
  d <- replyr_copy_to(my_db, data.frame(x=c(1,2)), 'd')
  d2 <- replyr_copy_from(d)
  print(d2)
  DBI::dbDisconnect(my_db)
}
```

replyr_copy_to	<i>Copy data to remote service.</i>
----------------	-------------------------------------

Description

Copy data to remote service.

Usage

```
replyr_copy_to(dest, df, name = paste(deparse(substitute(df)), collapse =
  " "), ..., rowNumberColumn = NULL, temporary = FALSE,
  overwrite = TRUE, maxrow = 1e+06)
```

Arguments

dest	remote data source
df	local data frame
name	name for new remote table
...	force later values to be bound by name
rowNumberColumn	if not null name to add row numbers to
temporary	logical, if TRUE try to create a temporary table
overwrite	logical, if TRUE try to overwrite
maxrow	max rows to allow in a remote to remote copy.

Value

remote handle

Examples

```
if (requireNamespace("RSQLite", quietly = TRUE)) {
  my_db <- DBI::dbConnect(RSQLite::SQLite(), ":memory:")
  RSQLite::initExtension(my_db)
  d <- replyr_copy_to(my_db, data.frame(x=c(1,2)), 'd')
  print(d)
  DBI::dbDisconnect(my_db)
}
```

replyr_dim	<i>Compute dimensions of a data.frame (work around https://github.com/rstudio/sparklyr/issues/976).</i>
------------	---

Description

Compute dimensions of a data.frame (work around <https://github.com/rstudio/sparklyr/issues/976>).

Usage

```
replyr_dim(x)
```

Arguments

x	tbl or item that can be coerced into such.
---	--

Value

dimensions (including rows)

Examples

```
d <- data.frame(x=c(1,2))
replyr_dim(d)
```

<code>replyr_filter</code>	<i>Filter a tbl on a column having values in a given set.</i>
----------------------------	---

Description

Filter a tbl on a column having values in a given set.

Usage

```
replyr_filter(x, cname, values, ..., verbose = TRUE,
             tempNameGenerator = mk_tmp_name_source("replyr_filter"))
```

Arguments

<code>x</code>	tbl or item that can be coerced into such.
<code>cname</code>	name of the column to test values of.
<code>values</code>	set of values to check set membership of.
<code>...</code>	force later arguments to bind by name.
<code>verbose</code>	logical if TRUE echo warnings
<code>tempNameGenerator</code>	temp name generator produced by <code>wrpr::mk_tmp_name_source</code> , used to record <code>dplyr::compute()</code> effects.

Value

new tbl with only rows where `cname` value is in `values` set.

Examples

```
values <- c('a','c')
d <- data.frame(x=c('a','a','b','b','c','c'),y=1:6,
               stringsAsFactors=FALSE)
replyr_filter(d,'x',values)
```

replyr_get_src	<i>Get the "remote data source" where a data.frame like object lives.</i>
----------------	---

Description

Get the "remote data source" where a data.frame like object lives.

Usage

```
replyr_get_src(df)
```

Arguments

df	data.frame style object
----	-------------------------

Value

source (string if data.frame, tbl, or data.table, NULL if unknown, remote source otherwise)

Examples

```
replyr_get_src(data.frame(x=1:2))
```

replyr_group_by	<i>group_by columns</i>
-----------------	-------------------------

Description

See also: <https://gist.github.com/skranz/9681509>

Usage

```
replyr_group_by(.data, colnames)
```

Arguments

.data	data.frame
colnames	character vector of column names to group by.

Value

.data grouped by columns named in colnames

See Also

[group_by](#), [group_by_at](#)

Examples

```
d <- data.frame(Sepal_Length= c(5.8,5.7),
                Sepal_Width= c(4.0,4.4),
                Species= 'setosa')
replyr_group_by(d, 'Species')
```

replyr_hasrows	<i>Check if a table has rows.</i>
----------------	-----------------------------------

Description

Check if a table has rows.

Usage

```
replyr_hasrows(d)
```

Arguments

d tbl or item that can be coerced into such.

Value

number of rows

Examples

```
d <- data.frame(x=c(1,2))
replyr_hasrows(d)
```

replyr_has_table	<i>check for a table</i>
------------------	--------------------------

Description

Work around connection v.s. handle issues <https://github.com/tidyverse/dplyr/issues/2849>

Usage

```
replyr_has_table(con, name)
```

Arguments

con	connection
name	character name to check for

Value

TRUE if table present

Examples

```
if (requireNamespace("RSQLite", quietly = TRUE)) {
  my_db <- DBI::dbConnect(RSQLite::SQLite(), ":memory:")
  RSQLite::initExtension(my_db)
  d <- replyr_copy_to(my_db, data.frame(x=c(1,2)), 'd')
  print(d)
  print(replyr_has_table(my_db, 'd'))
  DBI::dbDisconnect(my_db)
}
```

replyr_inTest	<i>Product a column noting if another columns values are in a given set.</i>
---------------	--

Description

Product a column noting if another columns values are in a given set.

Usage

```
replyr_inTest(x, cname, values, nname, ...,
  tempNameGenerator = mk_tmp_name_source("replyr_inTest"),
  verbose = TRUE)
```

Arguments

x	tbl or item that can be coerced into such.
cname	name of the column to test values of.
values	set of values to check set membership of.
nname	name for new column
...	force later parameters to bind by name
tempNameGenerator	temp name generator produced by wrapr::mk_tmp_name_source, used to record dplyr::compute() effects.
verbose	logical if TRUE echo warnings

Value

table with membership indications.

Examples

```
values <- c('a', 'c')
d <- data.frame(x=c('a', 'a', 'b', NA, 'c', 'c'), y=1:6,
               stringsAsFactors=FALSE)
replyr_inTest(d, 'x', values, 'match')
```

replyr_is_local_data *Test if data is local.*

Description

Test if data is local.

Usage

```
replyr_is_local_data(d)
```

Arguments

d data frame

Value

TRUE if local data (data.frame, tbl/tibble)

Examples

```
replyr_is_local_data(data.frame(x=1:3))
```

replayr_is_MySQL_data *Test if data is MySQL.*

Description

Test if data is MySQL.

Usage

```
replayr_is_MySQL_data(d)
```

Arguments

d data frame

Value

TRUE if Spark data

Examples

```
replayr_is_MySQL_data(data.frame(x=1:3))
```

replayr_is_Spark_data *Test if data is Spark.*

Description

Test if data is Spark.

Usage

```
replayr_is_Spark_data(d)
```

Arguments

d data frame

Value

TRUE if Spark data

Examples

```
replayr_is_Spark_data(data.frame(x=1:3))
```

replyr_list_tables *list tables*

Description

Work around connection v.s. handle issues <https://github.com/tidyverse/dplyr/issues/2849>

Usage

```
replyr_list_tables(con)
```

Arguments

con connection

Value

list of tables names

Examples

```
if (requireNamespace("RSQLite", quietly = TRUE)) {
  my_db <- DBI::dbConnect(RSQLite::SQLite(), ":memory:")
  RSQLite::initExtension(my_db)
  d <- replyr_copy_to(my_db, data.frame(x=c(1,2)), 'd',
                    overwrite=TRUE, temporary=TRUE)

  print(d)
  print(replyr_list_tables(my_db))
  DBI::dbDisconnect(my_db)
}
```

replyr_mapRestrictCols

Map names of columns to known values and drop other columns.

Description

Restrict a data item's column names and re-name them in bulk.

Usage

```
replyr_mapRestrictCols(x, nmap, ..., restrict = FALSE, reverse = FALSE)
```

Arguments

x	data item to work on
nmap	named list mapping with keys specifying new column names, and values as original column names.
...	force later arguments to bind by name
restrict	logical if TRUE restrict to columns mentioned in nmap.
reverse	logical if TRUE apply the inverse of nmap instead of nmap.

Details

Something like `replyr::replyr_mapRestrictCols` is only useful to get control of a function that is not parameterized (in the sense it has hard-coded column names inside its implementation that don't match column names in our data).

Value

data item with columns renamed (and possibly restricted).

See Also

[let](#), [replyr_reverseMap](#), [replyr_apply_f_mapped](#)

Examples

```
# an external function with hard-coded column names
DecreaseRankColumnByOne <- function(d) {
  d$RankColumn <- d$RankColumn - 1
  d
}

# our example data, with different column names
d <- data.frame(Sepal_Length=c(5.8,5.7),
                Sepal_Width=c(4.0,4.4),
                Species='setosa',rank=c(1,2))
print(d)

# map our data to expected column names so we can use function
nmap <- c(GroupColumn='Species',
          ValueColumn='Sepal_Length',
          RankColumn='rank')
print(nmap)
dm <- replyr_mapRestrictCols(d,nmap)
print(dm)

# can now apply code that expects hard-coded names.
dm <- DecreaseRankColumnByOne(dm)

# map back to our original column names (for the columns we retained)
```



```
# Note: can only map back columns that were retained in first mapping.
replyr_mapRestrictCols(dm, nmap, reverse=TRUE)
```

replyr_ncol	<i>Compute number of columns of a data.frame (work around https://github.com/rstudio/sparklyr/issues/976).</i>
-------------	--

Description

Compute number of columns of a data.frame (work around <https://github.com/rstudio/sparklyr/issues/976>).

Usage

```
replyr_ncol(x)
```

Arguments

x tbl or item that can be coerced into such.

Value

number of columns

Examples

```
d <- data.frame(x=c(1,2))
replyr_ncol(d)
```

replyr_nrow	<i>Compute number of rows of a tbl.</i>
-------------	---

Description

Number of row in a table. This function is not "group aware" it returns the total number of rows, not rows per dplyr group. Also replyr_nrow depends on data being returned to count, so some corner cases (such as zero columns) will count as zero rows. In particular work around dplyr issue 2871 <https://github.com/tidyverse/dplyr/issues/2871>.

Usage

```
replyr_nrow(x)
```

Arguments

x tbl or item that can be coerced into such.

Value

number of rows

Examples

```
d <- data.frame(x=c(1,2))
replyr_nrow(d)
```

replyr_quantile	<i>Compute quantiles on remote column (NA's filtered out) using binary search.</i>
-----------------	--

Description

NA's filtered out and does not break ties the same as stats::quantile.

Usage

```
replyr_quantile(x, cname, probs = seq(0, 1, 0.25), ...,
  tempNameGenerator = mk_tmp_name_source("replyr_quantile"))
```

Arguments

x tbl or item that can be coerced into such.

cname column name to compute over

probs numeric vector of probabilities with values in [0,1].

... force later arguments to be bound by name.

tempNameGenerator
temp name generator produced by wrapr::mk_tmp_name_source, used to record
dplyr::compute() effects.

Examples

```
d <- data.frame(xvals=rev(1:1000))
replyr_quantile(d, 'xvals')
```

replyr_quantilec	<i>Compute quantiles on remote column (NA's filtered out) using cumsum.</i>
------------------	---

Description

NA's filtered out and does not break ties the same as stats::quantile.

Usage

```
replyr_quantilec(x, cname, probs = seq(0, 1, 0.25), ...,
  tempNameGenerator = mk_tmp_name_source("replyr_quantilec"))
```

Arguments

x	tbl or item that can be coerced into such.
cname	column name to compute over (not 'n' or 'csum')
probs	numeric vector of probabilities with values in [0,1].
...	force later arguments to bind by name.
tempNameGenerator	temp name generator produced by wrapr::mk_tmp_name_source, used to record dplyr::compute() effects.

Examples

```
d <- data.frame(xvals=rev(1:1000))
replyr_quantilec(d, 'xvals')
```

replyr_rename	<i>Rename a column</i>
---------------	------------------------

Description

Rename a column

Usage

```
replyr_rename(.data, ..., newName, oldName)
```

Arguments

.data	data object to work on
...	force later arguments to bind by name
newName	character new column name
oldName	character old column name

Examples

```
d <- data.frame(Sepal_Length= c(5.8,5.7),
               Sepal_Width= c(4.0,4.4),
               Species= 'setosa', rank=c(1,2))
replyr_rename(d, newName = 'family', oldName = 'Species')
```

replyr_reverseMap	<i>Reverse a name assignment map (which are written NEW-NAME=OLDNAME).</i>
-------------------	--

Description

Reverse a name assignment map (which are written NEWNAME=OLDNAME).

Usage

```
replyr_reverseMap(nmap)
```

Arguments

nmap	named list mapping with keys specifying new column names, and values as original column names.
------	--

Value

inverse map

See Also

[let](#), [replyr_apply_f_mapped](#), [replyr_mapRestrictCols](#)

Examples

```
mp <- c(A='x', B='y')
print(mp)
replyr_reverseMap(mp)
```

replyr_select	<i>select columns</i>
---------------	-----------------------

Description

select columns

Usage

```
replyr_select(.data, colnames)
```

Arguments

.data	data object to work on
colnames	character column names

Examples

```
d <- data.frame(Sepal_Length= c(5.8,5.7),
                Sepal_Width= c(4.0,4.4),
                Species= 'setosa', rank=c(1,2))
replyr_select(d, c('Sepal_Length', 'Species'))
```

replyr_split	<i>split a data item by values in a column.</i>
--------------	---

Description

Partitions from by values in grouping column, and returns list. Only advised for a moderate number of groups and better if grouping column is an index. This plus lapply and replyr::bind_rows is powerful enough to implement "The Split-Apply-Combine Strategy for Data Analysis" <https://www.jstatsoft.org/article/view/v040>

Usage

```
replyr_split(df, gcolumn, ..., ocolumn = NULL, decreasing = FALSE,
             partitionMethod = "extract", maxgroups = 100, eagerCompute = FALSE)
```

Arguments

df	remote dplyr data item
gcolumn	grouping column
...	force later values to be bound by name
ocolumn	ordering column (optional)
decreasing	if TRUE sort in decreasing order by ocolumn
partitionMethod	method to partition the data, one of 'split' (only works over local data frames), or 'extract'
maxgroups	maximum number of groups to work over
eagerCompute	if TRUE call compute on split results

Value

list of data items

Examples

```
d <- data.frame(group=c(1,1,2,2,2),
                order=c(.1,.2,.3,.4,.5),
                values=c(10,20,2,4,8))
dSplit <- replyr_split(d, 'group', partitionMethod='extract')
dApp <- lapply(dSplit, function(di) data.frame(as.list(colMeans(di))))
replyr_bind_rows(dApp)
```

replyr_summary

Compute usable summary of columns of tbl.

Description

Compute per-column summaries and return as a data.frame. Warning: can be an expensive operation.

Usage

```
replyr_summary(x, ..., countUniqueNum = FALSE,
               countUniqueNonNum = FALSE, cols = NULL, compute = TRUE)
```

Arguments

x tbl or item that can be coerced into such.
... force additional arguments to be bound by name.
countUniqueNum logical, if true include unique non-NA counts for numeric cols.
countUniqueNonNum logical, if true include unique non-NA counts for non-numeric cols.
cols if not NULL set of columns to restrict to.
compute logical if TRUE call compute before working

Details

Can be slow compared to `dplyr::summarize_all()` (but serves a different purpose). Also, for numeric columns includes NaN in nna count (as is typical for R, e.g., `is.na(NaN)`). And note: `replayr_summary()` currently skips "raw" columns.

Value

summary of columns.

See Also

[rsummary](#)

Examples

```

d <- data.frame(p= c(TRUE, FALSE, NA),
               r= I(list(1,2,3)),
               s= NA,
               t= as.raw(3:5),
               w= 1:3,
               x= c(NA,2,3),
               y= factor(c(3,5,NA)),
               z= c('a',NA,'z'),
               stringsAsFactors=FALSE)
# sc <- sparklyr::spark_connect(version='2.2.0',
#                               master = "local")
# dS <- replayr_copy_to(sc, dplyr::select(d, -r, -t), 'dS',
#                      temporary=TRUE, overwrite=TRUE)
# replayr_summary(dS)
# sparklyr::spark_disconnect(sc)
if (requireNamespace("RSQLite", quietly = TRUE)) {
  my_db <- DBI::dbConnect(RSQLite::SQLite(), ":memory:")
  RSQLite::initExtension(my_db)
  dM <- replayr_copy_to(my_db, dplyr::select(d, -r, -t), 'dM',
                      temporary=TRUE, overwrite=TRUE)
  print(replayr_summary(dM))
  DBI::dbDisconnect(my_db)
}

```

```
d$q <- list(1,2,3)
replyr_summary(d)
```

replyr_testCols	<i>Run test on columns.</i>
-----------------	-----------------------------

Description

Applies user function to head of each column. Good for determining things such as column class.

Usage

```
replyr_testCols(x, f, n = 6L)
```

Arguments

x	tbl or item that can be coerced into such.
f	test function (returning logical, not depending on data length).
n	number of rows to use in calculation.

Value

logical vector of results.

Examples

```
d <- data.frame(x=c(1,2),y=c('a','b'))
replyr_testCols(d,is.numeric)
```

replyr_union_all	<i>Union two tables.</i>
------------------	--------------------------

Description

Spark 2* union_all has issues (<https://github.com/WinVector/replyr/blob/master/issues/UnionIssue.md>), and exposed union_all semantics differ from data-source back-end to back-end. This is an attempt to provide a join-based replacement.

Usage

```
replyr_union_all(tabA, tabB, ..., useDplyrLocal = TRUE,
  useSparkRbind = TRUE,
  tempNameGenerator = mk_tmp_name_source("replyr_union_all"))
```


Arguments

tabA	not-NULL table with at least 1 row.
tabB	not-NULL table with at least 1 row on same data source as tabA and common columns.
...	force later arguments to be bound by name.
useDplyrLocal	logical if TRUE use dplyr::bind_rows for local data.
useSparkRbind	logical if TRUE try to use rbind on Sparklyr data
tempNameGenerator	temp name generator produced by wrapr::mk_tmp_name_source, used to record dplyr::compute() effects.

Value

table with all rows of tabA and tabB (union_all).

Examples

```
d1 <- data.frame(x = c('a','b'), y = 1, stringsAsFactors= FALSE)
d2 <- data.frame(x = 'c', z = 1, stringsAsFactors= FALSE)
replayr_union_all(d1, d2, useDplyrLocal= FALSE)
```

replayr_uniqueValues *Compute number of unique values for each level in a column.*

Description

Compute number of unique values for each level in a column.

Usage

```
replayr_uniqueValues(x, cname)
```

Arguments

x	tbl or item that can be coerced into such.
cname	name of columns to examine, must not be equal to 'replayr_private_value_n'.

Value

unique values for the column.

Examples

```
d <- data.frame(x=c(1,2,3,3))
replayr_uniqueValues(d, 'x')
```

tableDescription	<i>Build a nice description of a table.</i>
------------------	---

Description

Please see <http://www.win-vector.com/blog/2017/05/managing-spark-data-handles-in-r/> for details. Note: one usually needs to alter the keys column which is just populated with all columns.

Usage

```
tableDescription(tableName, handle, ...,  
  keyInspector = key_inspector_all_cols)
```

Arguments

tableName	name of table to add to join plan.
handle	table or table handle to add to join plan (can already be in the plan).
...	force later arguments to bind by name.
keyInspector	function that determines preferred primary key set for table.

Details

Please see vignette('DependencySorting', package = 'replayr') and vignette('joinController', package = 'replayr') for more details.

Value

table describing the data.

See Also

[buildJoinPlan](#), [keysAreUnique](#), [makeJoinDiagramSpec](#), [executeLeftJoinPlan](#)

Examples

```
d <- data.frame(x=1:3, y=NA)  
tableDescription('d', d)
```

topoSortTables	<i>Topologically sort join plan so values are available before uses.</i>
----------------	--

Description

Depends on igraph package. Please see vignette('DependencySorting', package = 'replryr') and vignette('joinController', package = 'replryr') for more details.

Usage

```
topoSortTables(columnJoinPlan, leftTableName, ...)
```

Arguments

```
columnJoinPlan  join plan
leftTableName   which table is left
...             force later arguments to bind by name
```

Value

list with dependencyGraph and sorted columnJoinPlan

Examples

```
if (requireNamespace("RSQLite", quietly = TRUE)) {
  # note: employeeanddate is likely built as a cross-product
  #       join of an employee table and set of dates of interest
  #       before getting to the join controller step. We call
  #       such a table "row control" or "experimental design."
  my_db <- DBI::dbConnect(RSQLite::SQLite(), ":memory:")
  RSQLite::initExtension(my_db)
  tDesc <- example_employeeAndDate(my_db)
  columnJoinPlan <- buildJoinPlan(tDesc, check = FALSE)
  # unify keys
  columnJoinPlan$resultColumn[columnJoinPlan$resultColumn == 'id'] <- 'eid'
  # look at plan defects
  print(paste('problems:',
             inspectDescrAndJoinPlan(tDesc, columnJoinPlan)))
  # fix plan
  if (requireNamespace('igraph', quietly = TRUE)) {
    sorted <- topoSortTables(columnJoinPlan, 'employeeanddate')
    print(paste('problems:',
               inspectDescrAndJoinPlan(tDesc, sorted$columnJoinPlan)))
    # plot(sorted$dependencyGraph)
  }
  DBI::dbDisconnect(my_db)
  my_db <- NULL
}
```

`%land%`*Land a value to variable from a pipeline.*

Description

`%land%` and `%->%` ("writearrow") copy a pipeline value to a variable on the right hand side. `%land_%` and `%->_%` copy a pipeline value to a variable named by the value referenced by its right hand side argument.

Usage

```
value %land% name
```

```
value %->% name
```

```
value %->_% name
```

```
value %land_% name
```

Arguments

value	value to write
name	variable to write to

Details

Technically these operators are not "-> assignment", so they might not be specifically prohibited in an oppugnant reading of some style guides.

Value

value

Examples

```
sin(7) %->% z1
sin(7) %->_% 'z2'
varname <- 'z3'
sin(7) %->_% varname
```

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