

Package ‘joinEasy’

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Type Package

Title Performs Join Operations

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Description

Provides succinct solutions for simplifying the process of various join operations in R.

Depends sqldf

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R topics documented:

anti_join	1
cross_join	2
full_outer_exclusive_join	3
inner_join	3
joinEasy	4
left_outer_join	5
right_outer_join	6
semi_join	6

anti_join

Anti Join

Description

This function returns the set of all tuples that are in neither the intersection of the two input relations nor in the right dataframe.

Usage

```
{  
  # This function can be used without providing a primary key value:  
  anti_join(left, right)  
  # Or with a primary key value:  
  anti_join(left, right, key)  
}
```

Arguments

<code>left</code>	A dataframe which will be used as the left-side table in the operation.
<code>right</code>	A dataframe which will be used as the right-side table in the operation.
<code>key</code>	The primary key value. If not provided, the function will use the dataframes' row numbers as the primary key.

Value

<code>rtn</code>	The resulting dataframe.
------------------	--------------------------

Examples

```
# Example usage:
left <- data.frame(id = 100:106, letter = c('a', 'b', 'c', 'd', 'e', 'f', 'g'))
random_i <- sample(seq_len(nrow(left)))
cutpoint <- round(nrow(left) / 2)
right <- left[random_i[seq_len(cutpoint)],]
output <- anti_join(left, right, key = "id")
output
# right for comparison:
right
```

`cross_join`*Cross Join*

Description

This function returns the cartesian product of two relations.

Usage

```
cross_join(left, right)
```

Arguments

<code>left</code>	A dataframe which will be used as the left-side table in the operation.
<code>right</code>	A dataframe which will be used as the right-side table in the operation.

Value

<code>rtn</code>	The resulting dataframe.
------------------	--------------------------

Examples

```
# Example usage:
left <- data.frame(id = 100:106, letter = c('a', 'b', 'c', 'd', 'e', 'f', 'g'))
random_i <- sample(seq_len(nrow(left)))
cutpoint <- round(nrow(left) / 2)
right <- left[random_i[seq_len(cutpoint)],]
output <- cross_join(left, right)
output
```

```
full_outer_exclusive_join
```

Full Outer (INNER Exclusive) Join

Description

The ‘full_outer_exclusive_join’ function is not yet implemented. Once released, this function will return all rows from both left and right which are unique to the dataframes.

Usage

```
{
  # This function will be usable without providing a primary key value:
  full_outer_exclusive_join(left, right)
  # Or with a primary key value:
  full_outer_exclusive_join(left, right, key)
}
```

Arguments

<code>left</code>	A dataframe which will be used as the left-side table in the operation.
<code>right</code>	A dataframe which will be used as the right-side table in the operation.
<code>key</code>	The primary key value. If not provided, the function will use the dataframes’ row numbers as the primary key.

Value

<code>rtn</code>	The resulting dataframe.
------------------	--------------------------

Examples

```
# Planned example usage:
left <- data.frame(id = 100:106, letter = c('a', 'b', 'c', 'd', 'e', 'f', 'g'))
random_i <- sample(seq_len(nrow(left)))
cutpoint <- round(nrow(left) / 2)
right <- left[random_i[seq_len(cutpoint)],]
output <- full_outer_exclusive_join(left, right, key = "id")
output
```

```
inner_join
```

Inner Join

Description

This function returns only those rows which are common to both dataframes.

Usage

```
{
  # This function can be used without providing a primary key value:
  inner_join(left, right)
  # Or with a primary key value:
  inner_join(left, right, key)
}
```

Arguments

left	A dataframe which will be used as the left-side table in the operation.
right	A dataframe which will be used as the right-side table in the operation.
key	The primary key value. If not provided, the function will use the dataframes' row numbers as the primary key.

Value

rtn	The resulting dataframe.
------------	--------------------------

Examples

```
# Example usage:
left <- data.frame(id = 100:106, letter = c('a', 'b', 'c', 'd', 'e', 'f', 'g'))
random_i <- sample(seq_len(nrow(left)))
cutpoint <- round(nrow(left) / 2)
right <- left[random_i[seq_len(cutpoint)],]
output <- inner_join(left, right, key = "id")
output
```

joinEasy

Performs Join Operations

Description

Provides succinct solutions for simplifying the process of various join operations in R.

Details

The DESCRIPTION file: This package was not yet installed at build time.

Index: This package was not yet installed at build time.

anti_join **cross_join**

Author(s)

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References

www.mtitek.com/tutorials/oracle/sql-join.php Provides visual depictions of all operations present in this package

See Also

#~~ Optional links to other man pages, e.g. ~~ #~~ <pkg> ~~

Examples

```
# simple examples of the most important functions
```

left_outer_join	<i>Left Outer Join</i>
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Description

This function returns only those rows which are either common to both dataframes or only present in the left dataframe.

Usage

```
{
  # This function can be used without providing a primary key value:
  left_outer_join(left, right)
  # Or with a primary key value:
  left_outer_join(left, right, key)
}
```

Arguments

left	A dataframe which will be used as the left-side table in the operation.
right	A dataframe which will be used as the right-side table in the operation.
key	The primary key value. If not provided, the function will use the dataframes' row numbers as the primary key.

Value

rtn	The resulting dataframe.
------------	--------------------------

Examples

```
# Example usage:
left <- data.frame(id = 100:106, letter = c('a', 'b', 'c', 'd', 'e', 'f', 'g'))
random_i <- sample(seq_len(nrow(left)))
cutpoint <- round(nrow(left) / 2)
right <- left[random_i[seq_len(cutpoint)],]
output <- left_outer_join(left, right, key = "id")
output
```

<code>right_outer_join</code>	<i>Right Outer Join</i>
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Description

The ‘`right_outer_join`’ function is not yet implemented. Once released, this function will return only those rows which are either common to both dataframes or only present in the right dataframe.

Usage

```
{
  # This function will be usable without providing a primary key value:
  right_outer_join(left, right)
  # Or with a primary key value:
  right_outer_join(left, right, key)
}
```

Arguments

<code>left</code>	A dataframe which will be used as the left-side table in the operation.
<code>right</code>	A dataframe which will be used as the right-side table in the operation.
<code>key</code>	The primary key value. If not provided, the function will use the dataframes’ row numbers as the primary key.

Value

<code>rtn</code>	The resulting dataframe.
------------------	--------------------------

Examples

```
# Planned example usage:
left <- data.frame(id = 100:106, letter = c('a', 'b', 'c', 'd', 'e', 'f', 'g'))
random_i <- sample(seq_len(nrow(left)))
cutpoint <- round(nrow(left) / 2)
right <- left[random_i[seq_len(cutpoint)],]
output <- right_outer_join(left, right, key = "id")
output
```

<code>semi_join</code>	<i>Semi Join</i>
------------------------	------------------

Description

This function returns a similar result to that of the `inner_join`, except that this one never duplicates rows.

Usage

```
{  
  # This function can be used without providing a primary key value:  
  semi_join(left, right)  
  # Or with a primary key value:  
  semi_join(left, right, key)  
}
```

Arguments

left	A dataframe which will be used as the left-side table in the operation.
right	A dataframe which will be used as the right-side table in the operation.
key	The primary key value. If not provided, the function will use the dataframes' row numbers as the primary key.

Value

rtn	The resulting dataframe.
------------	--------------------------

Examples

```
# Example usage:  
left <- data.frame(id = 100:106, letter = c('a', 'b', 'c', 'd', 'e', 'f', 'g'))  
random_i <- sample(seq_len(nrow(left)))  
cutpoint <- round(nrow(left) / 2)  
right <- left[random_i[seq_len(cutpoint)],]  
output <- semi_join(left, right, key = "id")  
output
```