

# Package ‘biblio’

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**Encoding** UTF-8

**Title** Interacting with BibTeX Databases

**Depends** R(>= 4.1)

**Imports** methods, stringr, tools, utils, yamlme

**Suggests** covr, devtools, rmarkdown, testthat

**LazyData** true

**SystemRequirements** pandoc (>= 1.14) - <http://pandoc.org>

**Description** Reading and writing BibTeX files using data frames in R sessions.

**License** GPL (>= 2)

**URL** <https://kamapu.github.io/biblio/>

**BugReports** <https://github.com/kamapu/biblio/issues>

**RoxygenNote** 7.3.1

**NeedsCompilation** no

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## R topics documented:

as	2
bib2bibentry	2
compare_df	3
comp_df-class	4
detect_keys	4
lib_df-class	5
print	5
read_bib	6

reflist . . . . .	7
synopsis . . . . .	8
update_data . . . . .	9
write_bib . . . . .	10

<b>Index</b>	<b>12</b>
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as	<i>Coerce 'lib_df' objects</i>
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### Description

Coercion 'lib\_df' objects.

### Arguments

x                    An object to be coerced.

### Examples

```
## Read installed library
Refs <- read_bib(x = file.path(path.package("biblio"),
                              "LuebertPliscoeff.bib"))

# Convert lib_df to data frame
as(Refs[1:5, ], "data.frame")

# Convert lib_df to bibentry
as(Refs[1:5, ], "bibentry")
```

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bib2bibentry	<i>Convert lib_df to bibentry</i>
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### Description

Conversion method for lib\_df objects into bibentry.

### Usage

```
bib2bibentry(x, ...)
```

```
## S3 method for class 'lib_df'
bib2bibentry(x, ...)
```

### Arguments

x                    A lib\_df object to be converted.  
 ...                 Further arguments passed among methods (not yet in use).

**Value**

An bibentry object.

**Examples**

```
## Read installed electronic library
Bib <- read_bib(x = file.path(path.package("biblio"),
                             "LuebertPliscoff.bib"))

## Convert the first five entries
bib2bibentry(Bib[1:5, ])
```

---

compare\_df

*Compare data frames and libraries*

---

**Description**

Report on differences between two versions of the same data frame or electronic library. When used for data frames, you need to indicate the variable containing IDs for each entry, while applied to [lib\\_df](#) objects, the variable 'bibtexkey' will be considered as ID per default.

The output printed in the console will advice about added and deleted entries in 'y' as well as any change in the entries common to both versions.

**Usage**

```
compare_df(x, y, key, ...)
```

## S4 method for signature 'data.frame,data.frame,character'

```
compare_df(x, y, key, ...)
```

## S4 method for signature 'lib\_df,lib\_df,missing'

```
compare_df(x, y, key, ...)
```

**Arguments**

x                   The (old) reference data frame.  
y                   The updated (new) data frame.  
key                  A character value with the name of the variable used as primary key in the tables.  
...                  Further arguments passed among methods.

**Value**

A S3 object of class [comp\\_df](#), which can be printed in the console by [print\(\)](#).

**See Also**

[update\(\)](#), [lib\\_df](#), [comp\\_df](#)

**Examples**

```
# Partially matching libraries
Refs1 <- synopsis[1:10, ]
Refs2 <- synopsis[6:15, ]

# some modification in second library
Refs2[3, "title"] <- "New Title"

# compare libraries
compare_df(Refs1, Refs2)
```

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comp_df-class	<i>Compared libraries</i>
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**Description**

An S3 class for compared data frames. A list containing added, deleted entries on the regarding a key column and cells that are modified.

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detect_keys	<i>Detect bibtexkeys used in an r-markdown document</i>
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**Description**

This function screens a character vector (usually an imported r-markdown document) for the use of citations by bibtexkeys (@bibtexkey), retrieving the detected key with its occurrence in the vector, assuming each element as a line of the original document.

This function is based on `bbt_detect_citations()` from the package `rbbt`.

**Usage**

```
detect_keys(x, ...)
```

```
## S3 method for class 'character'
detect_keys(x, ...)
```

```
## S3 method for class 'rmd_doc'
detect_keys(x, ...)
```

**Arguments**

x	A character vector, a file imported by <code>readLines()</code> or an object imported by <code>read_rmd()</code> . If the character vector is the name of a Rmd or a Quarto document, <code>readLines()</code> will be internally called to read it.
...	Further arguments passed among methods. In character-method they are passed to <code>readLines()</code> .

**Value**

A data frame with two columns, bibtexkey for the found keys and line with the line number of the occurrence of the key in the document.

**Examples**

```
## Screen for citations in installed document
cited_refs <- detect_keys(file.path(path.package("biblio"), "document.Rmd"))
cited_refs
```

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lib_df-class	<i>Electronic library</i>
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**Description**

An S3 class for library entries. This class inherits properties from data frames.

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print	<i>Print content of lib_df objects</i>
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---

**Description**

A method for a brief overview on the content of a [lib\\_df](#) or a [comp\\_df](#) object.

**Usage**

```
## S3 method for class 'lib_df'
print(x, maxsum = 4, ...)

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

**Arguments**

x	An object of class 'lib_df'.
maxsum	An integer value indicating the number of entries to be displayed in the printed output.
...	Further arguments passed among methods.

**Value**

A print in the console.

**Author(s)**

Miguel Alvarez

## Examples

```
Refs <- read_bib(x = file.path(
  path.package("biblio"),
  "LuebertPliscoeff.bib"
))

print(Refs, maxsum = 10)
```

---

read\_bib

*Read BibTeX Databases*

---

## Description

Reading BibTeX databases and importing into R as a data frame. All the fields will be inserted as character values.

## Usage

```
read_bib(x, ...)
```

## Arguments

**x** A single character value with the path to a BibTeX file. Alternatively it can be a character vector containing the lines of a BibTeX library, for instance after using [readLines\(\)](#).

**...** Further arguments passed to [readLines\(\)](#).

## Value

An object of class [lib\\_df](#).

## Examples

```
Refs <- read_bib(x = file.path(
  path.package("biblio"),
  "LuebertPliscoeff.bib"
))
Refs
```

---

 reflist

*Write a Reference List in rmarkdown*


---

### Description

A fast way to produce a reference list in an r-markdown document from a `lib_df` object.

This function may or may not produce intermediate files (bib and Rmd) and the result can be assigned to an object for further edition (see `yamlme::update()`).

A html file will be written by `write_rmd()` and `render_rmd()` in the working directory and displayed by `browseURL()`.

### Usage

```
reflist(x, ...)

## S3 method for class 'character'
reflist(
  x,
  output_file,
  delete_source = TRUE,
  title = "Automatic Reference List",
  output = "html_document",
  nocite = "'@*'",
  urlcolor = "blue",
  encoding = "UTF-8",
  ...
)

## S3 method for class 'lib_df'
reflist(x, filename, delete_source = TRUE, ...)
```

### Arguments

<code>x</code>	A <code>lib_df</code> object to produce the reference list. In the character method, a character value indicating the path of a bibtex file (passed to <code>read_bib()</code> ).
<code>...</code>	Further arguments passed to the yaml header in the intermediate Rmarkdown document.
<code>output_file</code>	A character value with the name for the written Rmarkdown file.
<code>delete_source</code>	A logical value indicating whether written bib file should be deleted after rendering html or not.
<code>title, output, nocite, urlcolor</code>	Arguments used for the yaml-header in r-markdown and passed to <code>write_rmd()</code> . They can be cancelled using the value NULL (not recommended for nocite).
<code>encoding</code>	A character value indicating the encoding string. It is passed to <code>write_bib()</code> .
<code>filename</code>	A character value with the name for the written Rmd file, without file extension.

**Value**

An invisible object of class `rmd_doc`. A Rmd file will be written by `write_rmd()` as well.

**Examples**

```
## Not run:  
relist(synopsis)  
  
## End(Not run)
```

---

synopsis

*References by Lueber and Plischoff (2018)*

---

**Description**

Example of an object formatted as `lib_df`. This library is published with the references of the book **Bioclimatic and vegetational synopsis of Chile** by **Luebert and Plischoff (2017)**.

**Usage**

```
synopsis
```

**Format**

An object of class `lib_df` (inherits from `data.frame`) with 1701 rows and 23 columns.

**Source**

[doi:10.5281/zenodo.60800](https://doi.org/10.5281/zenodo.60800)

**Examples**

```
data(synopsis)  
  
## Import from installed bibtex file  
synopsis <- read_bib(x = file.path(  
  path.package("biblio"),  
  "LuebertPlischoff.bib"  
))
```

---

update_data	<i>Update data frames</i>
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---

## Description

This function compares two versions of the same data frame and detect changes as additions, deleted entries or updates (modified entries).

A method to compare [lib\\_df](#) objects is also provided as well as a replace method.

## Usage

```
update_data(object, revision, key, ...)

## S4 method for signature 'data.frame,data.frame,character'
update_data(
  object,
  revision,
  key,
  add = FALSE,
  delete = FALSE,
  update = FALSE,
  ...
)

## S4 method for signature 'lib_df,lib_df,missing'
update_data(object, revision, key, ...)

update_data(object, key, ...) <- value

## S4 replacement method for signature 'data.frame,character,data.frame'
update_data(object, key, ...) <- value

## S4 replacement method for signature 'lib_df,missing,lib_df'
update_data(object, key, ...) <- value
```

## Arguments

object	A data frame or a <a href="#">lib_df</a> object representing the original version.
revision	The updated version of 'object' to be compared.
key	A character value indicating the column used as identifier. This variable have to be in both versions otherwise this function will retrieve an error.
...	Further arguments passed among methods.
delete, add, update	Logical value indicating whether the action should be carried out. If all are 'FALSE', this function will just report differences as done by <a href="#">compare_df</a> .
value	The updated version of 'object' in the replace methods.

**Value**

Either an invisible output with a print in the console or an updated object of class `lib_df`.

**Examples**

```
# Adding an ID to data set iris
iris2 <- iris
iris2$id <- 1:nrow(iris2)

# rows to add using mean values per species
iris_mod <- aggregate(cbind(
  Sepal.Length, Sepal.Width, Petal.Length,
  Petal.Width
) ~ Species, data = iris2, FUN = mean)
iris_mod$id <- (1:nrow(iris_mod)) + nrow(iris2)
iris_mod <- do.call(rbind, list(iris2, iris_mod[, colnames(iris2)]))

# delete some entries
iris_mod <- iris_mod[-c(15, 75, 105, 145), ]

# modify entries
iris_mod$Petal.Length[c(20, 30)] <- 0
iris_mod$Petal.Width[c(20, 50)] <- 0

# just a comparison
update_data(iris2, iris_mod, key = "id")

# do update
iris2 <- update_data(iris2, iris_mod,
  key = "id", delete = TRUE, add = TRUE,
  update = TRUE
)
```

---

write\_bib

*Write BibTeX Files*


---

**Description**

BibTeX databases can be created from data frames, interacting with Postgres databases.

**Usage**

```
write_bib(x, ...)

## S3 method for class 'lib_df'
write_bib(x, filename, encoding = "UTF-8", ...)
```

**Arguments**

- x            A data frame with bibliographic entries.
- ...        Further arguments passed to `file()`.
- filename    A character value with the path and the name of the file to be written.
- encoding    Character value with the encoding (passed to `file()`).

**Value**

A bibtex file.

# Index

- \* **datasets**
  - synopsis, 8
- as, 2
- bib2bibentry, 2
- bib2bibentry, lib\_df-method
  - (bib2bibentry), 2
- bib2bibentry.lib\_df (bib2bibentry), 2
- browseURL(), 7
  
- coerce, bibentry, lib\_df-method (as), 2
- coerce, lib\_df, bibentry-method (as), 2
- coerce, lib\_df, data.frame-method (as), 2
- comp\_df, 3, 5
- comp\_df-class, 4
- compare\_df, 3, 9
- compare\_df, data.frame, data.frame, character-method
  - (compare\_df), 3
- compare\_df, lib\_df, lib\_df, missing-method
  - (compare\_df), 3
  
- detect\_keys, 4
- detect\_keys, character-method
  - (detect\_keys), 4
- detect\_keys, rmd\_doc-method
  - (detect\_keys), 4
- detect\_keys.character (detect\_keys), 4
- detect\_keys.rmd\_doc (detect\_keys), 4
  
- file(), 11
  
- lib\_df, 3, 5, 6, 8–10
- lib\_df-class, 5
  
- print, 5
- print(), 3
- print, comp\_df-method (print), 5
- print, lib\_df-method (print), 5
- print.comp\_df (print), 5
- print.lib\_df (print), 5
  
- read\_bib, 6
- read\_bib(), 7
- read\_rmd(), 4
- readLines(), 4, 6
- reflist, 7
- reflist, character-method (reflist), 7
- reflist, lib\_df-method (reflist), 7
- reflist.character (reflist), 7
- reflist.lib\_df (reflist), 7
- render\_rmd(), 7
  
- synopsis, 8
  
- update(), 3
- update\_data, 9
- update\_data, data.frame, data.frame, character-method
  - (update\_data), 9
- update\_data, lib\_df, lib\_df, character-method
  - (update\_data), 9
- update\_data, lib\_df, lib\_df, missing-method
  - (update\_data), 9
- update\_data<- (update\_data), 9
- update\_data<- , data.frame, character, data.frame-method
  - (update\_data), 9
- update\_data<- , lib\_df, missing, lib\_df-method
  - (update\_data), 9
  
- write\_bib, 10
- write\_bib(), 7
- write\_rmd(), 7, 8
  
- yamlme:::update(), 7