

# Package: InsectLabelR (via r-universe)

June 2, 2026

**Title** Create Labels for Insect in Collection

**Version** 1.0.4

**Description** Streamlines the creation of high-quality labels for insect pinning. By taking a dataset as input, the package allow to generate printable labels in 'LaTeX' and PDF format, helping researchers and entomologists maintain accurate and standardized specimen records. Requires a compatible installation of 'pdflatex' (e.g. <https://www.tug.org/texlive/>). For enhanced accessibility, the package includes a user-friendly 'shiny' application (accessible online <https://nicolas-moiroux.shinyapps.io/InsectLabelR/>), which provides a graphical interface for generating labels without requiring programming expertise.

**License** GPL (>= 3)

**Encoding** UTF-8

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**Suggests** testthat (>= 3.0.0), shinyjs, readxl, readr, readODS, DT, editbl (>= 1.1.0)

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**Depends** R (>= 4.0)

**LazyData** true

**URL** <https://github.com/Nmoiroux/InsectLabelR>

**BugReports** <https://github.com/Nmoiroux/InsectLabelR/issues>

**SystemRequirements** pdflatex

**Config/pak/sysreqs** cmake make libicu-dev libuv1-dev zlib1g-dev

**Repository** <https://r-multiverse.r-universe.dev>

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create_pdf	<i>Generate a Complete 'LaTeX' Document and Compile it into a PDF</i>
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## Description

This function generates a 'LaTeX' document by sequentially adding a header, printing labels for each row of a data table, appending a footer, and then compiling the 'LaTeX' code into a PDF document. The 'LaTeX' code is adapted from work by Samuel Brown (see <https://github.com/sdjbrown/publicFiles/blob/master/labels.tex> and <http://the-praise-of-insects.blogspot.com/2010/03/latex-insect-labels.html>). Ensure you have a compatible installation of 'pdflatex' (see <https://www.latex-project.org/get/>).

## Usage

```
create_pdf(
  file_out,
  ind_list,
  print_info,
  lab_width = 15,
  lab_height = 9,
  font_size = 4,
  n_col = 8,
  col_N_name = NA,
  hl_col = "orange",
  compile = TRUE
)
```

**Arguments**

file_out	A character string specifying the name of returned 'LaTeX' and PDF files.
ind_list	A data frame containing individual data. Each row represents data for one individual to be printed in the document.
print_info	A data frame specifying the parameters for printing, including field names, formatting options, what data to print and, on how many labels.
lab_width	An integer specifying the width (in mm) for the labels (default is 15 mm).
lab_height	An integer specifying the height (in mm) for the labels (default is 9).
font_size	A real (one digit) specifying the size of the font for the label (default is 4)
n_col	An integer specifying the number of label columns in the 'LaTeX' document (default is 8 columns).
col_N_name	A character string specifying the column name to be used for labels duplication (default is NA).
hl_col	A character string specifying the highlight color for specific elements (default is "orange").
compile	Boolean (TRUE/FALSE). Does the 'LaTeX' document should be compiled into a PDF document ? (default to TRUE)

**Details**

This function first calls `print_header` to write the beginning of the 'LaTeX' document. Then, it iterates over each row of `ind_list`, calling `print_line` to generate labels according to the provided `print_info`. After all labels are printed, it appends the 'LaTeX' document footer using `print_bottom`. Finally, it compiles the 'LaTeX' document into a PDF using `pdflatex`.

**Value**

The function creates a 'LaTeX' file, compiles it into a PDF, and saves the outputs to the specified location. It does not return any value in R.

**Examples**

```
outfile <- file.path(tempdir(), "labels.tex")

# This example only generates the LaTeX file (no pdflatex required)
create_pdf(
  file_out = outfile,
  ind_list = mosquito_collection,
  print_info = print_parameters,
  compile = FALSE
)

## Not run:
# Full PDF generation requires pdflatex installed
create_pdf(
  file_out = outfile,
  ind_list = mosquito_collection,
```

```
  print_info = print_parameters
)

## End(Not run)
```

---

InsectLabelR_App	<i>Launch the InsectLabelR 'shiny' application</i>
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### Description

This function launches an interactive 'shiny' application allowing users to use InsectLabelR with either example datasets included in the package or user-provided data. It provides a graphical interface for generating labels without requiring programming expertise.

The application is available online at <https://nicolas-moiroux.shinyapps.io/InsectLabelR/>.

### Usage

```
InsectLabelR_App()
```

### Value

A 'shiny' **application** object.

### Examples

```
if(interactive()){
  InsectLabelR_App()
}
```

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mosquito_collection	<i>Example mosquito collection dataset for label generation</i>
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### Description

A toy dataset representing mosquito specimens collected in southern France and intended for pinning and label printing with `create_pdf()`.

**Format**

A data frame with 24 rows and 12 variables:

**rec\_date** Collection date (character). Example: "06/2024".

**rec\_place** Locality name, optionally including administrative unit.

**rec\_cntry** Country of collection.

**Y** Latitude in degrees-minutes-seconds (DMS) format (character).

**X** Longitude in degrees-minutes-seconds (DMS) format (character).

**rec\_name** Collector name.

**id\_name** Identifier (person who identified the specimen).

**sex** Biological sex of the specimen (e.g., "Male", "Female").

**genus** Genus name (e.g., *Culiseta*, *Culex*).

**subgenus** Subgenus name (may be empty).

**species** Species epithet.

**N** Number of specimens sharing identical metadata.

**id** Unique specimen identifier (integer).

**Details**

The dataset follows the standard input structure expected by **InsectLabelR**, with one row per specimen (or per group of identical specimens when  $N > 1$ ). It includes collection metadata, taxonomic identification, sex, geographic coordinates (in DMS format), and a unique specimen identifier.

This dataset can be used to test label layout, formatting, and PDF generation without requiring external files.

The dataset contains specimens of *Culiseta (Cal.) longiareolata* and *Culex pipiens*.

**Source**

Simulated dataset for demonstration purposes.

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print\_bottom

*Append the Closing Section to a 'LaTeX' Document*

---

**Description**

This function writes the closing commands to a 'LaTeX' document. Specifically, it appends the end of a multicols\* environment and the document environment.

**Usage**

```
print_bottom(file_out)
```

**Arguments**

`file_out` A character string specifying the path to the output file where the 'LaTeX' closing commands will be appended.

**Details**

The function appends 'LaTeX' commands for ending a multiple-column layout (using the `multicols*` environment) and the document. It ensures the proper closure of a 'LaTeX' document that was generated by the preceding steps.

**Value**

This function appends 'LaTeX' code to the file specified in `file_out`. It does not return any output in R.

---

`print_header` *Create 'LaTeX' Header for Mosquito Labels Document*

---

**Description**

This function generates a 'LaTeX' document header for mosquito labels and writes it to the specified output file. The header includes various 'LaTeX' packages, document settings, metadata, and the front cover of the document. The 'LaTeX' code is adapted from work by Samuel Brown (see <https://github.com/sdjbrown/publicFiles/blob/master/labels.tex> and <http://the-praise-of-insects.blogspot.com/2010/03/latex-insect-labels.html>).

**Usage**

```
print_header(
  file_out,
  lab_width = 15,
  lab_height = 9,
  font_size = 4,
  n_col = 8
)
```

**Arguments**

`file_out` A character string specifying the name of the output file to write the 'LaTeX' header to.

`lab_width` An integer specifying the width (in mm) for the labels (default is 15 mm).

`lab_height` An integer specifying the height (in mm) for the labels (default is 9).

`font_size` A real (one digit) specifying the size of the font for the label (default is 4)

`n_col` An integer specifying the number of columns in the 'LaTeX' document (default is 8 columns).

## Details

This function is primarily used for creating labels in 'LaTeX' for mosquito specimen identification. It sets the document's layout and font sizes, allowing customization of label size and number of columns on the page.

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print_line	<i>Print 'LaTeX' Labels for Each Line of a Data Table</i>
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## Description

This function prints 'LaTeX' labels for each line of a data table based on specified information and user-defined printing parameters. It formats each label according to the provided field names, data, and formatting options. The function generates 'LaTeX' code for printing the labels into an external file. The 'LaTeX' code is based on the work by Samuel Brown (see <https://github.com/sdjbrown/publicFiles/blob/master/labels.tex> and <http://the-praise-of-insects.blogspot.com/2010/03/latex-insect-labels.html>)

## Usage

```
print_line(
  file_out,
  ind_list,
  print_info,
  line_n,
  col_N_name = NA,
  hl_col = "orange"
)
```

## Arguments

file_out	A character string specifying the path to the output file where 'LaTeX' code will be appended.
ind_list	A data frame containing the data for individuals. Each row corresponds to a specific individual.
print_info	A data frame specifying the printing parameters, including which fields to print, formatting options, and field names.
line_n	An integer indicating the row number in ind_list for which the labels should be created.
col_N_name	A character string specifying the name of the column in ind_list that indicates the number of individuals per row. Default is NA indicating 1 row per individual (and no replication of the labels)
hl_col	A character string specifying the color to be used for text highlighting

### Details

The function retrieves data from a specified row in `ind_list` and matches it with the corresponding print parameters in `print_info`. It formats each label using the 'LaTeX' code according to the user-defined options in `print_info`, such as whether to italicize or bracket certain fields, and whether to include field names before the information. The function generates 'LaTeX' code for individual labels and appends it to the specified output file.

### Value

The function appends 'LaTeX' code to the file specified in `file_out`. It does not return anything in R.

---

print\_parameters

*Example print parameter table for label formatting*

---

### Description

A toy parameter table defining how specimen data fields are arranged, formatted, and distributed across labels when using `create_pdf()`.

### Format

A data frame with 13 rows and 10 variables:

**field\_name** Name of the field in the 'mosquito\_collection' data table.

**print** Logical (0/1). Whether the field is printed on labels.

**label\_no** Numeric. Label number on which the field is printed (e.g., 1 = identification label, 2 = locality/date label, 3 = collector/identifier label).

**order\_lab** Numeric. Order of appearance within the label.

**prefix** Character string added before the field value (e.g., "leg.", "det.", "no.").

**print\_opt\_it** Logical (0/1). Whether the field should be printed in italics (typically for taxonomic names).

**print\_opt\_par** Logical (0/1). Whether the field should be printed in parentheses (commonly used for subgenus).

**line\_break** Logical (0/1). Whether a line break follows the field.

**print\_opt\_hl** Logical (0/1). Whether the field should be highlighted.

**print\_sex\_symbol** Logical (0/1). Whether to print a sex symbol instead of the full sex text.

## Details

This dataset illustrates the structure expected by **InsectLabelR** to control label composition (field selection, ordering, formatting options, line breaks, highlighting, and symbol printing).

Each row corresponds to a field present in the specimen data table and specifies whether and how it should be printed.

This example configuration produces three labels per specimen:

1. Taxonomic label (genus, subgenus, species, sex).
2. Collection label (date and locality).
3. Collector/identifier label (collector, determiner, specimen number).

Taxonomic names are printed in italics, the subgenus is enclosed in parentheses, and sex can optionally be rendered as a biological symbol.

## Source

Simulated parameter table for demonstration purposes.

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sex_to_latex	<i>Convert Sex Data to 'LaTeX' Format</i>
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## Description

This function converts a string representing sex data into the appropriate 'LaTeX' code for displaying male or female symbols in small font.

## Usage

```
sex_to_latex(sex_data)
```

## Arguments

sex_data	A character string representing the sex of an individual. It can start with "f" or "F" for female, "m" or "M" for male, or it can be NA.
----------	--

## Details

The function checks the first letter of the input string. If it starts with "f" or "F", it returns the 'LaTeX' code for the female symbol. If it starts with "m" or "M", it returns the 'LaTeX' code for the male symbol. If the input is NA or unrecognized, the function returns NA.

## Value

A character string containing the corresponding 'LaTeX' code: "\smallfemale" for female, "\smallmale" for male, or NA if the input is not recognized or is NA.

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