

# Package ‘licoread’

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**Title** Reads Raw Files from Li-COR Gas Analyzers

**Version** 0.2.0

**Date** 2025-05-08

**Description** Reads raw files from Li-COR gas analyzers and produces a dataframe that can directly be used with 'flexible' <<https://cran.r-project.org/package=flexible>>.

**License** GPL (>= 3)

**Encoding** UTF-8

**RoxygenNote** 7.3.2

**Suggests** knitr, rmarkdown, testthat (>= 3.0.0)

**Config/testthat/edition** 3

**Imports** dplyr, jsonlite, lubridate, purrr, readr, rlang, stringr, tibble, tidyr, yaml

**URL** <https://jogaudard.github.io/licoread/>

**BugReports** <https://github.com/jogaudard/licoread/issues>

**VignetteBuilder** knitr

**NeedsCompilation** no

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**Depends** R (>= 4.1.0)

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data_82z	<i>to read the raw data</i>
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**Description**

to read the raw data

**Usage**

data\_82z(filepath, data\_file, data\_name, gases, filename)

**Arguments**

- filepath            name and path to the 82z archive
- data\_file          name of the file with raw data
- data\_name          vector of colnames
- gases              list of gases
- filename           name of the 82z archive

**Value**

a long df with the actual data contained in the data file

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data_name_82z	<i>create colnames for data tibble</i>
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**Description**

create colnames for data tibble

**Usage**

```
data_name_82z(filepath, data_file)
```

**Arguments**

filepath	name and path to the 82z archive
data_file	name of the file with raw data

**Value**

a character vector or the colnames of the data file

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flexible_81x	<i>makes df from 81x files compatible with flexible</i>
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**Description**

makes df from 81x files compatible with flexible

**Usage**

```
flexible_81x(df, focus_gas, id_cols, datetime_col)
```

**Arguments**

df	input dataframe from <a href="#">licoread</a>
focus_gas	gas to select
id_cols	columns to identify unique fluxes
datetime_col	column containing datetime information

**Value**

a df with the focus gas column renamed as "f\_conc" and f\_fluxid in chronological order of datetime

---

flexible_82z	<i>makes df from 82z files compatible with flexible</i>
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**Description**

makes df from 82z files compatible with flexible

**Usage**

```
flexible_82z(df, focus_gas)
```

**Arguments**

df	input dataframe from <a href="#">licoread</a>
focus_gas	gas to select

**Value**

an unnested df with only the selected gas

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licoread	<i>reads Li-COR files in a given location</i>
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**Description**

reads Li-COR files in a given location

**Usage**

```
licoread(  
  location,  
  file_type = "auto",  
  file_type_list = c("82z", "81x", "auto"),  
  data_file = "data.csv",  
  meta_file = "metadata.json",  
  regex_file = "(\\w*-)*\\w*(?=[. ]82z$)",  
  sample = FALSE  
)
```

**Arguments**

location	location of the files
file_type	type of file (82z or 81x). If "auto" (default), the function will try to detect it by itself.
file_type_list	list of file types
data_file	name of the file with raw data
meta_file	name of the file with meta data
regex_file	regex expression matching the name of the 82z file. Here in case the user has a different than the default and for easier updates.
sample	sample = n randomly selects n files to be imported. This allows for testing the setup before importing a potentially large list of files which will take time and be difficult to handle.

**Value**

a tibble (nested or not depending on raw data) containing all the data from the raw files present at the location provided

**Examples**

```
path_82z <- system.file("extdata/82z", package = "licoread")
licoread(path_82z)
```

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licoread_auto	<i>finds out the file type for licoread</i>
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**Description**

finds out the file type for licoread

**Usage**

```
licoread_auto(file_list)
```

**Arguments**

file_list	list of files found in the location
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**Value**

a single character string indicating the file type

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`licoread_to_fluxible` *selects the focus gas and makes a df to use in [fluxible](#)*

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## Description

selects the focus gas and checks the columns needed for the [fluxible](#) workflow

## Usage

```
licoread_to_fluxible(
  df,
  focus_gas,
  datetime_col,
  id_cols = c("File Name", "Obs#"),
  file_type = "auto",
  file_type_list = c("82z", "81x", "auto")
)
```

## Arguments

<code>df</code>	input dataframe from <a href="#">licoread</a>
<code>focus_gas</code>	gas to select
<code>datetime_col</code>	column containing datetime information if date and time are in two different columns, provide a character vector of the form <code>c("date", "time")</code>
<code>id_cols</code>	columns to identify unique fluxes
<code>file_type</code>	type of file (82z or 81x). If "auto" (default), the function will try to detect it by itself.
<code>file_type_list</code>	list of file types

## Value

an unnested df with only the selected gas

## Examples

```
path_82z <- system.file("extdata/82z", package = "licoread")
gas_df_82z <- licoread(path_82z)
licoread_to_fluxible(gas_df_82z, "LI-7810_CH4_DRY",
  datetime_col = c("LI-8250_DATE", "LI-8250_TIME"))
```

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list_gases	<i>lists gases present in the dataset</i>
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**Description**

prints a character vector of the gases present in the dataset. This is to be used to obtain the exact names of the gases before using [licoread\\_to\\_fluxible](#).

**Usage**

```
list_gases(df, file_type = "auto", file_type_list = c("82z", "81x", "auto"))
```

**Arguments**

df	the dataframe as imported with <a href="#">licoread</a>
file_type	type of file (currently works only for 82z). If "auto" (default), the function will try to detect it by itself.
file_type_list	list of file types

**Value**

a character vector of the gases present in the dataset

**Examples**

```
path_82z <- system.file("extdata/82z", package = "licoread")
gas_df_82z <- licoread(path_82z)
list_gases(gas_df_82z)
```

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metadata_82z	<i>read meta data file inside 82z archive</i>
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**Description**

read meta data file inside 82z archive

**Usage**

```
metadata_82z(filepath, meta_file)
```

**Arguments**

filepath	name and path to the 82z archive
meta_file	name of the file with meta data

**Value**

a tibble with the metadata from one observation

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names_df	<i>to get a vector of names of a df, matching a regex</i>
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**Description**

to get a vector of names of a df, matching a regex

**Usage**

```
names_df(df, regname = "name\\d")
```

**Arguments**

df	the df to get the names from
regname	the regex expression to match

**Value**

a df with the names of the meta df

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oneobs_81x	<i>reading a single measurement from 81x file</i>
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**Description**

reads a single measurement from a licor .81x file

**Usage**

```
oneobs_81x(start, end, all_obs, file)
```

**Arguments**

start	line number at which the measurement starts
end	line number at which the measurement ends
all_obs	list of all the lines from the full file
file	filepath to the 81x files

**Value**

a df with 1 row with the meta data of the measurement and raw data nested



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oneobs_82z	<i>to read one measurement from the 82z archive</i>
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**Description**

to read one measurement from the 82z archive

**Usage**

```
oneobs_82z(filepath, data_file, meta_file, regex_file)
```

**Arguments**

filepath	path to the 82z archive
data_file	name of the file with raw data
meta_file	name of the file with meta data
regex_file	regex expression matching the name of the 82z file. Here in case the user has a different than the default and for easier updates.

**Value**

a tibble with all the data and metadata from one observation (one file)

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read_81x_onefile	<i>reads 81x licor file</i>
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**Description**

reads a .81x file with several measurements

**Usage**

```
read_81x_onefile(file)
```

**Arguments**

file	filepath the the 81x file to read
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**Value**

a nested tibble with the meta data from each measurements as row and the raw data nested

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units_82z	<i>to create a nested tibble with the units of data</i>
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**Description**

to create a nested tibble with the units of data

**Usage**

```
units_82z(filepath, data_file, data_name, filename)
```

**Arguments**

filepath	name and path to the 82z archive
data_file	name of the file with raw data
data_name	vector of colnames
filename	name of the 82z archive

**Value**

a tibble with the units of the variables contained in the raw data

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