

Package ‘licoread’

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Title Reads Raw Files from Li-COR Gas Analyzers

Version 0.1.1

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

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

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

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

Description

makes df from 81x files compatible with flexible

Usage

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

Arguments

df	input dataframe from licoread
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>
--------------	---

Description

makes df from 82z files compatible with flexible

Usage

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

Arguments

df	input dataframe from licoread
focus_gas	gas to select

Value

an unnested df with only the selected gas

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

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)
```

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

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

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 licoread
<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"))
```

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

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

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

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

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

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

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)

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

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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