

Package ‘lightr’

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Title Read Spectrometric Data and Metadata

Version 1.4

Description Parse various reflectance/transmittance/absorbance spectra file formats to extract spectral data and metadata, as described in Gruson, White & Maia (2019) <doi:10.21105/joss.01857>. Among other formats, it can import files from 'Avantes' <<https://www.avantes.com/>>, 'CRAIC' <<https://www.microspectra.com/>>, and 'OceanInsight' (formerly 'OceanOptics') <<https://www.oceaninsight.com/>> brands.

Depends R (>= 3.5.0)

Imports future.apply, progressr, xml2 (>= 1.0.0)

Suggests digest, knitr, pavo, rmarkdown, spelling, testthat (>= 3.0.0), withr

URL <https://docs.ropensci.org/lightr/>,
<https://github.com/ropensci/lightr>

BugReports <https://github.com/ropensci/lightr/issues>

License GPL (>= 2)

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Author Hugo Gruson [cre, aut, cph] (<<https://orcid.org/0000-0002-4094-1476>>),
Rafael Maia [aut, cph] (<<https://orcid.org/0000-0002-7563-9795>>),
Thomas White [aut, cph] (<<https://orcid.org/0000-0002-3976-1734>>),
Kotya Karapetyan [ctb, cph] (Author of the MATLAB script to read
AvaSoft7 binary files (CC-BY))

Maintainer Hugo Gruson <hugo.gruson+R@normalesup.org>

Repository CRAN

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lr_compute_processed *Compute processed spectral data*

Description

Compute processed spectral data, from the raw count/scope data, counts from a dark reference, and from a white reference.

Usage

```
lr_compute_processed(spdata)
```

Arguments

spdata data.frame containing the spectral data with the columns 'scope', 'dark', and 'white'

lr_convert_tocsv *Convert spectral data files to csv files*

Description

Convert spectral data files to csv files

Usage

```
lr_convert_tocsv(  
  where = NULL,  
  ext = "txt",  
  decimal = ".",  
  sep = NULL,  
  subdir = FALSE,  
  cores = NULL,  
  ignore.case = TRUE,  
  overwrite = FALSE  
)
```

Arguments

| | |
|-------------|--|
| where | Folder in which files are located (defaults to current working directory). |
| ext | File extension to be searched for, without the "." (defaults to txt). You can also use a character vector to specify multiple file extensions. |
| decimal | Character to be used to identify decimal plates (defaults to .). |
| sep | Column delimiting characters to be considered in addition to the default (which are: tab, space, and ";") |
| subdir | Should subdirectories within the where folder be included in the search? (defaults to FALSE). |
| cores | deprecated. See future::plan() for more details on how to customise your parallelisation strategy. |
| ignore.case | Should the extension search be case insensitive? (defaults to TRUE) |
| overwrite | logical. Should the function overwrite existing files with the same name? (defaults to FALSE). |

Details

You can customise the type of parallel processing used by this function with the [future::plan\(\)](#) function. This works on all operating systems, as well as high performance computing (HPC) environment. Similarly, you can customise the way progress is shown with the [progressr::handlers\(\)](#) functions (progress bar, acoustic feedback, nothing, etc.)

Value

Convert input files to csv and invisibly return the list of created file paths

Warning

This step loses all metadata associated to the spectra. This metadata is critical to ensure reproducibility. We recommended you use [lr_get_metadata\(\)](#) to extract this information from your raw data.

Ir_get_metadata *Extract metadata from spectra files*

Description

Finds and imports metadata from spectra files in a given location.

Usage

```
Ir_get_metadata(  
  where = getwd(),  
  ext = "ProcSpec",  
  sep = NULL,  
  subdir = FALSE,  
  subdir.names = FALSE,  
  cores = NULL,  
  ignore.case = TRUE  
)
```

Arguments

| | |
|--------------|--|
| where | Folder in which files are located (defaults to current working directory). |
| ext | File extension to be searched for, without the "." (defaults to txt). You can also use a character vector to specify multiple file extensions. |
| sep | Column delimiting characters to be considered in addition to the default (which are: tab, space, and ";") |
| subdir | Should subdirectories within the where folder be included in the search? (defaults to FALSE). |
| subdir.names | Should subdirectory path be included in the name of the spectra? (defaults to FALSE). |
| cores | deprecated. See future::plan() for more details on how to customise your parallelisation strategy. |
| ignore.case | Should the extension search be case insensitive? (defaults to TRUE) |

Details

You can customise the type of parallel processing used by this function with the [future::plan\(\)](#) function. This works on all operating systems, as well as high performance computing (HPC) environment. Similarly, you can customise the way progress is shown with the [progressr::handlers\(\)](#) functions (progress bar, acoustic feedback, nothing, etc.)

Value

A data.frame containing one file per row and the following columns:

- name: File name (without the extension)
- user: Name of the spectrometer operator
- date: Timestamp of the recording (ISO 8601 format)
- spec_model: Model of the spectrometer
- spec_ID: Unique ID of the spectrometer
- white_inttime: Integration time of the white reference (in ms)
- dark_inttime: Integration time of the dark reference (in ms)
- sample_inttime: Integration time of the sample (in ms)
- white_avgs: Number of averaged measurements for the white reference
- dark_avgs: Number of averaged measurements for the dark reference
- sample_avgs: Number of averaged measurements for the sample
- white_boxcar: Boxcar width for the white reference
- dark_boxcar: Boxcar width for the dark reference
- sample_boxcar: Boxcar width for the sample reference

Warning

white_inttime, dark_inttime and sample_inttime should be equal. The normalised data may be inaccurate otherwise.

References

White TE, Dalrymple RL, Noble DWA, O'Hanlon JC, Zurek DB, Umbers KDL. Reproducible research in the study of biological coloration. *Animal Behaviour*. 2015 Aug 1;106:51-7 (doi: [10.1016/j.anbehav.2015.05.007](https://doi.org/10.1016/j.anbehav.2015.05.007)).

Examples

```
lr_get_metadata(system.file("testdata", "procspec_files",  
                           package = "lightr"),  
               ext = "ProcSpec")
```

 lr_get_spec

Extract spectral data from spectra files

Description

Finds and imports reflectance/transmittance/absorbance data from spectra files in a given location.

Usage

```
lr_get_spec(
  where = getwd(),
  ext = "txt",
  lim = c(300, 700),
  decimal = ".",
  sep = NULL,
  subdir = FALSE,
  subdir.names = FALSE,
  cores = NULL,
  ignore.case = TRUE,
  interpolate = TRUE
)
```

Arguments

| | |
|--------------|---|
| where | Folder in which files are located (defaults to current working directory). |
| ext | File extension to be searched for, without the "." (defaults to txt). You can also use a character vector to specify multiple file extensions. |
| lim | A vector with two numbers determining the wavelength limits to be considered (defaults to <code>c(300, 700)</code>). |
| decimal | Character to be used to identify decimal plates (defaults to <code>.</code>). |
| sep | Column delimiting characters to be considered in addition to the default (which are: tab, space, and ";") |
| subdir | Should subdirectories within the where folder be included in the search? (defaults to FALSE). |
| subdir.names | Should subdirectory path be included in the name of the spectra? (defaults to FALSE). |
| cores | deprecated. See <code>future::plan()</code> for more details on how to customise your parallelisation strategy. |
| ignore.case | Should the extension search be case insensitive? (defaults to TRUE) |
| interpolate | Boolean indicated whether spectral data should be interpolated and pruned at every nanometre. Note that this option can only work if all input data samples the same wavelengths. Defaults to TRUE. |

Details

You can customise the type of parallel processing used by this function with the `future::plan()` function. This works on all operating systems, as well as high performance computing (HPC) environment. Similarly, you can customise the way progress is shown with the `progressr::handlers()` functions (progress bar, acoustic feedback, nothing, etc.)

Value

A `data.frame`, containing the wavelengths in the first column and individual imported spectral files in the subsequent columns.

See Also

`pavo::getspec()`

Examples

```
lr_get_spec(system.file("testdata", package = "lightr"), ext = "jdx")
```

| | |
|------------------|--|
| lr_parse_generic | <i>Generic function to parse spectra files that don't have a specific parser</i> |
|------------------|--|

Description

Generic function to parse spectra files that don't have a specific parser

Usage

```
lr_parse_generic(filename, decimal = ".", sep = NULL)
```

Arguments

| | |
|----------|---|
| filename | Path of the file to parse |
| decimal | Character to be used to identify decimal plates (defaults to <code>.</code>). |
| sep | Column delimiting characters to be considered in addition to the default (which are: tab, space, and <code>;</code>) |

Details

'processed' column computed by official software and provided as is.

Value

A named list of two elements:

- data: a dataframe with columns "wl", "dark", "white", "scope" and "processed", in this order.
- metadata: a character vector with metadata including:
 - user: Name of the spectrometer operator
 - date: Timestamp of the recording (ISO 8601 format)
 - spec_model: Model of the spectrometer
 - spec_ID: Unique ID of the spectrometer
 - white_inttime: Integration time of the white reference (in ms)
 - dark_inttime: Integration time of the dark reference (in ms)
 - sample_inttime: Integration time of the sample (in ms)
 - white_avgs: Number of averaged measurements for the white reference
 - dark_avgs: Number of averaged measurements for the dark reference
 - sample_avgs: Number of averaged measurements for the sample
 - white_boxcar: Boxcar width for the white reference
 - dark_boxcar: Boxcar width for the dark reference
 - sample_boxcar: Boxcar width for the sample reference

Examples

```
lr_parse_generic(system.file("testdata", "spec.csv", package = "lightr"),
                 sep = ",")
lr_parse_generic(system.file("testdata", "CRAIC_export.txt",
                             package = "lightr"))
```

lr_parse_jaz

Parse OceanInsight converted file

Description

Parse OceanInsight (formerly OceanOptics) converted file. <https://www.oceaninsight.com/>

Usage

```
lr_parse_jaz(filename)
```

```
lr_parse_jazirrad(filename)
```

Arguments

filename Path of the file to parse

Details

'processed' column computed by official software and provided as is.

Value

A named list of two elements:

- data: a dataframe with columns "wl", "dark", "white", "scope" and "processed", in this order.
- metadata: a character vector with metadata including:
 - user: Name of the spectrometer operator
 - date: Timestamp of the recording (ISO 8601 format)
 - spec_model: Model of the spectrometer
 - spec_ID: Unique ID of the spectrometer
 - white_inttime: Integration time of the white reference (in ms)
 - dark_inttime: Integration time of the dark reference (in ms)
 - sample_inttime: Integration time of the sample (in ms)
 - white_avgs: Number of averaged measurements for the white reference
 - dark_avgs: Number of averaged measurements for the dark reference
 - sample_avgs: Number of averaged measurements for the sample
 - white_boxcar: Boxcar width for the white reference
 - dark_boxcar: Boxcar width for the dark reference
 - sample_boxcar: Boxcar width for the sample reference

Examples

```
lr_parse_jaz(system.file("testdata", "jazspec.jaz", package = "lightr"))
lr_parse_jazirrad(system.file("testdata", "irrad.JazIrrad",
                             package = "lightr"))
```

 lr_parse_jdx

Parse OceanInsight JCAMP-DX (.jdx) file

Description

Parse OceanInsight (formerly OceanOptics) JCAMP-DX (.jdx) file. <https://www.oceaninsight.com/>

Usage

```
lr_parse_jdx(filename)
```

Arguments

filename Path of the file to parse

Details

'processed' column computed by **lightr** with the function `lr_compute_processed()`.

Value

A named list of two elements:

- data: a dataframe with columns "wl", "dark", "white", "scope" and "processed", in this order.
- metadata: a character vector with metadata including:
 - user: Name of the spectrometer operator
 - date: Timestamp of the recording (ISO 8601 format)
 - spec_model: Model of the spectrometer
 - spec_ID: Unique ID of the spectrometer
 - white_inttime: Integration time of the white reference (in ms)
 - dark_inttime: Integration time of the dark reference (in ms)
 - sample_inttime: Integration time of the sample (in ms)
 - white_avgs: Number of averaged measurements for the white reference
 - dark_avgs: Number of averaged measurements for the dark reference
 - sample_avgs: Number of averaged measurements for the sample
 - white_boxcar: Boxcar width for the white reference
 - dark_boxcar: Boxcar width for the dark reference
 - sample_boxcar: Boxcar width for the sample reference

References

McDonald RS, Wilks PA. JCAMP-DX: A Standard Form for Exchange of Infrared Spectra in Computer Readable Form. *Applied Spectroscopy*. 1988;42(1):151-62.

Examples

```
lr_parse_jdx(system.file("testdata", "OceanOptics_period.jdx",  
                        package = "lightr"))
```

| | |
|-------------------|---|
| lr_parse_procspec | <i>Parse OceanInsight ProcSpec file</i> |
|-------------------|---|

Description

Parse OceanInsight (formerly OceanOptics) ProcSpec file. <https://www.oceaninsight.com/>

Usage

```
lr_parse_procspec(filename)
```

Arguments

filename Path of the file to parse

Details

'processed' column computed by official software and provided as is.

Value

A named list of two elements:

- data: a dataframe with columns "wl", "dark", "white", "scope" and "processed", in this order.
- metadata: a character vector with metadata including:
 - user: Name of the spectrometer operator
 - date: Timestamp of the recording (ISO 8601 format)
 - spec_model: Model of the spectrometer
 - spec_ID: Unique ID of the spectrometer
 - white_inttime: Integration time of the white reference (in ms)
 - dark_inttime: Integration time of the dark reference (in ms)
 - sample_inttime: Integration time of the sample (in ms)
 - white_avgs: Number of averaged measurements for the white reference
 - dark_avgs: Number of averaged measurements for the dark reference
 - sample_avgs: Number of averaged measurements for the sample
 - white_boxcar: Boxcar width for the white reference
 - dark_boxcar: Boxcar width for the dark reference
 - sample_boxcar: Boxcar width for the sample reference

References

<https://www.oceaninsight.com/support/faqs/software/>

Examples

```
lr_parse_procspec(system.file("testdata", "procspec_files",  
                             "OceanOptics_Linux.ProcSpec",  
                             package = "lightr"))
```

| | |
|--------------|------------------------------|
| lr_parse_spc | <i>Parse SPC binary file</i> |
|--------------|------------------------------|

Description

Parse SPC binary file. (Used by CRAIC <https://www.microspectra.com/> and OceanInsight <https://www.oceaninsight.com/>)

Usage

```
lr_parse_spc(filename)
```

Arguments

| | |
|----------|---------------------------|
| filename | Path of the file to parse |
|----------|---------------------------|

Details

'processed' column computed by official software and provided as is.

Value

A named list of two elements:

- data: a dataframe with columns "wl", "dark", "white", "scope" and "processed", in this order.
- metadata: a character vector with metadata including:
 - user: Name of the spectrometer operator
 - date: Timestamp of the recording (ISO 8601 format)
 - spec_model: Model of the spectrometer
 - spec_ID: Unique ID of the spectrometer
 - white_inttime: Integration time of the white reference (in ms)
 - dark_inttime: Integration time of the dark reference (in ms)
 - sample_inttime: Integration time of the sample (in ms)
 - white_avgs: Number of averaged measurements for the white reference
 - dark_avgs: Number of averaged measurements for the dark reference
 - sample_avgs: Number of averaged measurements for the sample
 - white_boxcar: Boxcar width for the white reference
 - dark_boxcar: Boxcar width for the dark reference
 - sample_boxcar: Boxcar width for the sample reference

In development

Metadata parsing has not yet been implemented for this file format.

Examples

```
lr_parse_spc(system.file("testdata", "compare", "CRAIC", "CRAIC.spc",
                        package = "lightr"))
```

| | |
|--------------|----------------------------------|
| lr_parse_trm | <i>Parse Avantes binary file</i> |
|--------------|----------------------------------|

Description

Parse Avantes binary file (TRM, ABS, ROH, DRK, REF, RAW8, RFL8 file extensions). <https://www.avantes.com/products/spectrometers/>

Usage

```
lr_parse_trm(filename)
lr_parse_abs(filename)
lr_parse_roh(filename)
lr_parse_rfl8(filename, specnum = 1L)
lr_parse_raw8(filename, specnum = 1L)
```

Arguments

| | |
|----------|---|
| filename | Path of the file to parse |
| specnum | Integer representing the position of the spectrum to read in the file. This option only makes sense for AvaSoft8 files and is ignored in the other cases. |

Details

'processed' column computed by **lightr** with the function [lr_compute_processed\(\)](#).

Value

A named list of two elements:

- data: a dataframe with columns "wl", "dark", "white", "scope" and "processed", in this order.
- metadata: a character vector with metadata including:
 - user: Name of the spectrometer operator
 - date: Timestamp of the recording (ISO 8601 format)
 - spec_model: Model of the spectrometer
 - spec_ID: Unique ID of the spectrometer
 - white_inttime: Integration time of the white reference (in ms)

- dark_inttime: Integration time of the dark reference (in ms)
- sample_inttime: Integration time of the sample (in ms)
- white_avgs: Number of averaged measurements for the white reference
- dark_avgs: Number of averaged measurements for the dark reference
- sample_avgs: Number of averaged measurements for the sample
- white_boxcar: Boxcar width for the white reference
- dark_boxcar: Boxcar width for the dark reference
- sample_boxcar: Boxcar width for the sample reference

Examples

```
lr_parse_trm(system.file("testdata", "avantes_trans.TRM",
                        package = "lightr"))
lr_parse_roh(system.file("testdata", "avantes_reflect.ROH",
                        package = "lightr"))

lr_parse_rfl8(system.file("testdata", "compare", "Avantes",
                        "feather.RFL8",
                        package = "lightr"),
              specnum = 1)
lr_parse_rfl8(system.file("testdata", "compare", "Avantes",
                        "feather.RFL8",
                        package = "lightr"),
              specnum = 2)
```

| | |
|---------------------------|-------------------------------------|
| <code>lr_parse_ttt</code> | <i>Parse Avantes converted file</i> |
|---------------------------|-------------------------------------|

Description

Parse Avantes converted file. <https://www.avantes.com/products/spectrometers/>

Usage

```
lr_parse_ttt(filename)
```

```
lr_parse_trt(filename)
```

Arguments

| | |
|-----------------------|---------------------------|
| <code>filename</code> | Path of the file to parse |
|-----------------------|---------------------------|

Details

'processed' column computed by official software and provided as is.

Value

A named list of two elements:

- data: a dataframe with columns "wl", "dark", "white", "scope" and "processed", in this order.
- metadata: a character vector with metadata including:
 - user: Name of the spectrometer operator
 - date: Timestamp of the recording (ISO 8601 format)
 - spec_model: Model of the spectrometer
 - spec_ID: Unique ID of the spectrometer
 - white_inttime: Integration time of the white reference (in ms)
 - dark_inttime: Integration time of the dark reference (in ms)
 - sample_inttime: Integration time of the sample (in ms)
 - white_avgs: Number of averaged measurements for the white reference
 - dark_avgs: Number of averaged measurements for the dark reference
 - sample_avgs: Number of averaged measurements for the sample
 - white_boxcar: Boxcar width for the white reference
 - dark_boxcar: Boxcar width for the dark reference
 - sample_boxcar: Boxcar width for the sample reference

Examples

```
lr_parse_ttt(system.file("testdata", "avantes_export.ttt",  
                        package = "lightr"))  
lr_parse_trt(system.file("testdata", "avantes_export2.trt",  
                        package = "lightr"))
```

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