

Package ‘lightr’

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Title Read Spectrometric Data and Metadata

Version 2.1.0

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](https://doi.org/10.21105/joss.01857)>. Among other formats, it can import files from 'Avantes' <<https://www.avantes.com/>>, 'CRAIC' <<https://www.microspectra.com/>>, and 'OceanOptics'/'OceanInsight' <<https://www.oceanoptics.com/>> brands.

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Suggests digest, future, knitr, lintr, 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>

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lr_convert_tocsv	<i>Convert spectral data files to csv files</i>
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Description

Convert spectral data files to csv files

Usage

```
lr_convert_tocsv(  
  where = NULL,  
  ext = "txt",  
  subdir = FALSE,  
  ignore.case = TRUE,  
  overwrite = FALSE,  
  metadata = TRUE,  
  parser = NULL,  
  ...  
)
```

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. |
| subdir | Should subdirectories within the where folder be included in the search? (defaults to FALSE). |

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).
metadata	logical (defaults to TRUE). Should metadata be exported as well? They will be exported in csv files with the <code>_metadata.csv</code> suffix.
parser	Optional function to specify the parser to be used. By default, the parser is automatically selected based on the file extension. This argument can be used to force the use of a specific parser, or to pass a custom parser. See details for more information on custom parsers. Please note that this argument can produce easily produce invalid results if used incorrectly and only advanced users should use it, with extra caution.
...	Arguments passed to individual parsers.

Details

Parallel processing:

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.)

Custom parsers:

To create a custom parser to pass to the `parser` argument, you need to create with the following signature:

- input: a file path (string) as a first argument, and optionally, any additional arguments needed to parse the file. These arguments can be passed to the `lr_get_spec()` function via the `...` argument.
- output: a named list of two elements, as defined in `lr_parse_generic()`.

Value

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

Warning

When `metadata = TRUE`, if **either** the data **or** metadata export fails, nothing will be returned for this file.

lr_get_metadata	<i>Extract metadata from spectra files</i>
-----------------	--

Description

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

Usage

```
lr_get_metadata(
  where = getwd(),
  ext = "ProcSpec",
  subdir = FALSE,
  subdir.names = FALSE,
  ignore.case = TRUE,
  parser = NULL,
  ...
)
```

Arguments

<code>where</code>	Folder in which files are located (defaults to current working directory).
<code>ext</code>	File extension to be searched for, without the "." (defaults to txt). You can also use a character vector to specify multiple file extensions.
<code>subdir</code>	Should subdirectories within the <code>where</code> folder be included in the search? (defaults to FALSE).
<code>subdir.names</code>	Should subdirectory path be included in the name of the spectra? (defaults to FALSE).
<code>ignore.case</code>	Should the extension search be case insensitive? (defaults to TRUE)
<code>parser</code>	Optional function to specify the parser to be used. By default, the parser is automatically selected based on the file extension. This argument can be used to force the use of a specific parser, or to pass a custom parser. See details for more information on custom parsers. Please note that this argument can produce easily produce invalid results if used incorrectly and only advanced users should use it, with extra caution.
<code>...</code>	Arguments passed to individual parsers.

Details**Parallel processing:**

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.)

Custom parsers:

To create a custom parser to pass to the `parser` argument, you need to create with the following signature:

- input: a file path (string) as a first argument, and optionally, any additional arguments needed to parse the file. These arguments can be passed to the `lr_get_spec()` function via the `...` argument.
- output: a named list of two elements, as defined in `lr_parse_generic()`.

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
- datetime: 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),
  subdir = FALSE,
  subdir.names = FALSE,
  ignore.case = TRUE,
  interpolate = TRUE,
  parser = NULL,
  ...
)
```

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 c(300, 700)).
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).
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.
parser	Optional function to specify the parser to be used. By default, the parser is automatically selected based on the file extension. This argument can be used to force the use of a specific parser, or to pass a custom parser. See details for more information on custom parsers. Please note that this argument can produce easily produce invalid results if used incorrectly and only advanced users should use it, with extra caution.
...	Arguments passed to individual parsers.

Details

Parallel processing:

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.)

Custom parsers:

To create a custom parser to pass to the `parser` argument, you need to create with the following signature:

- input: a file path (string) as a first argument, and optionally, any additional arguments needed to parse the file. These arguments can be passed to the `lr_get_spec()` function via the `...` argument.
- output: a named list of two elements, as defined in `lr_parse_generic()`.

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

```
spcs <- lr_get_spec(system.file("testdata", package = "lightr"), ext = "jdx")
head(spcs)
```

lr_parse_csv	<i>Parse csv files</i>
--------------	------------------------

Description

Parse csv files

Usage

```
lr_parse_csv(filename, decimal = ".", sep = ",", ...)
```

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>)
...	ignored

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
 - datetime: Timestamp of the recording in format '%Y-%m-%d %H:%M:%S' and UTC timezone. If timezone is missing in source file, UTC time will be assumed (for reproducibility purposes across computers with different localtimes).
 - 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

```
res_csv <- lr_parse_csv(
  system.file("testdata", "spec.csv", package = "lightr")
)
head(res_csv$data)
# No metadata is extracted with this parser
res_csv$metadata
```

lr_parse_generic

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

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 .).
sep	Column delimiting characters to be considered in addition to the default (which are: tab, space, and ";")
...	ignored

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
 - datetime: Timestamp of the recording in format '%Y-%m-%d %H:%M:%S' and UTC timezone. If timezone is missing in source file, UTC time will be assumed (for reproducibility purposes across computers with different localtimes).
 - 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

```
res_csv <- lr_parse_generic(
  system.file("testdata", "spec.csv", package = "lightr"),
  sep = ", "
)
head(res_csv$data)
# No metadata is extracted with this parser
res_csv$metadata

res_craic <- lr_parse_generic(
  system.file("testdata", "CRAIC_export.txt", package = "lightr")
)
head(res_craic$data)
```

```
# No metadata is extracted with this parser
res_craic$metadata
```

lr_parse_jaz	<i>Parse OceanOptics converted file</i>
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Description

Parse OceanOptics/OceanInsight converted file. <https://www.oceanoptics.com/>

Usage

```
lr_parse_jaz(filename, ...)

lr_parse_jazirrad(filename, ...)

lr_parse_oceanoptics_jaz(filename, ...)

lr_parse_oceanoptics_jazirrad(filename, ...)
```

Arguments

filename	Path of the file to parse
...	ignored

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
 - datetime: Timestamp of the recording in format '%Y-%m-%d %H:%M:%S' and UTC timezone. If timezone is missing in source file, UTC time will be assumed (for reproducibility purposes across computers with different localtimes).
 - 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

```
res_jaz <- lr_parse_oceanoptics_jaz(system.file("testdata", "jazspec.jaz",
                                              package = "lightr"))
head(res_jaz$data)
res_jaz$metadata

res_jazirrad <- lr_parse_oceanoptics_jazirrad(system.file("testdata", "irrad.JazIrrad",
                                                         package = "lightr"))
head(res_jazirrad$data)
res_jazirrad$metadata

res_usb4000 <- lr_parse_oceanoptics_jaz(system.file("testdata", "00usb4000.txt",
                                                    package = "lightr"))
head(res_usb4000$data)
res_usb4000$metadata

res_transmission <- lr_parse_oceanoptics_jaz(
  system.file("testdata", "FMNH6834.00000001.Master.Transmission",
              package = "lightr")
)
head(res_transmission$data)
res_transmission$metadata
```

lr_parse_jdx

Parse OceanOptics JCAMP-DX (.jdx) file

Description

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

Usage

```
lr_parse_jdx(filename, ...)

lr_parse_oceanoptics_jdx(filename, ...)
```

Arguments

filename	Path of the file to parse
...	ignored

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
 - datetime: Timestamp of the recording in format '%Y-%m-%d %H:%M:%S' and UTC timezone. If timezone is missing in source file, UTC time will be assumed (for reproducibility purposes across computers with different localtimes).
 - 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

```
res_jdx <- lr_parse_oceanoptics_jdx(system.file("testdata", "OceanOptics_period.jdx",
package = "lightr"))
head(res_jdx$data)
res_jdx$metadata
```

lr_parse_procspec	<i>Parse OceanOptics ProcSpec file</i>
-------------------	--

Description

Parse OceanOptics/OceanInsight ProcSpec file. <https://www.oceanoptics.com/>

Usage

```
lr_parse_procspec(filename, verify_checksum = FALSE, ...)
```

```
lr_parse_oceanoptics_procspec(filename, verify_checksum = FALSE, ...)
```

Arguments

filename	Path of the file to parse
verify_checksum	Logical (defaults to FALSE). Should we check if the file has been modified since its creation by the spectrometer? An error will be returned if the check fails.
...	ignored

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
 - datetime: Timestamp of the recording in format '%Y-%m-%d %H:%M:%S' and UTC timezone. If timezone is missing in source file, UTC time will be assumed (for reproducibility purposes across computers with different localtimes).
 - 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.oceanoptics.com/software/>

Examples

```
res <- lr_parse_oceanoptics_procspec(system.file("testdata", "procspec_files",
                                                "OceanOptics_Linux.ProcSpec",
                                                package = "lightr"),
                                   verify_checksum = TRUE)

head(res$data)
res$metadata
```

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

Description

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

Usage

```
lr_parse_spc(filename, ...)

lr_parse_craic_spc(filename, ...)

lr_parse_oceanoptics_spc(filename, ...)
```

Arguments

filename	Path of the file to parse
...	ignored

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
 - datetime: Timestamp of the recording in format '%Y-%m-%d %H:%M:%S' and UTC timezone. If timezone is missing in source file, UTC time will be assumed (for reproducibility purposes across computers with different localtimes).

- 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

```
res <- lr_parse_craic_spc(
  system.file(
    "testdata", "compare", "CRAIC", "CRAIC.spc",
    package = "lightr"
  )
)
head(res$data)
res$metadata
```

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, ...)
```

```

lr_parse_irr8(filename, specnum = 1L, ...)

lr_parse_avantes_trm(filename, ...)

lr_parse_avantes_abs(filename, ...)

lr_parse_avantes_roh(filename, ...)

lr_parse_avantes_rfl8(filename, specnum = NULL, ...)

lr_parse_avantes_raw8(filename, specnum = NULL, ...)

lr_parse_avantes_irr8(filename, specnum = NULL, ...)

```

Arguments

filename	Path of the file to parse
...	ignored
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
 - datetime: Timestamp of the recording in format '%Y-%m-%d %H:%M:%S' and UTC timezone. If timezone is missing in source file, UTC time will be assumed (for reproducibility purposes across computers with different localtimes).
 - 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

```

res_trm <- lr_parse_avantes_trm(
  system.file("testdata", "avantes_trans.TRM", package = "lightr")
)
head(res_trm$data)
res_trm$metadata

res_roh <- lr_parse_avantes_roh(
  system.file("testdata", "avantes_reflect.ROH", package = "lightr")
)
head(res_roh$data)
res_roh$metadata

# This parser has a unique `specnum` argument
res_rfl8_1 <- lr_parse_avantes_rfl8(
  system.file("testdata", "compare", "Avantes", "feather.RFL8", package = "lightr"),
  specnum = 1
)
head(res_rfl8_1$data)
res_rfl8_1$metadata

res_rfl8_2 <- lr_parse_avantes_rfl8(
  system.file("testdata", "compare", "Avantes", "feather.RFL8", package = "lightr"),
  specnum = 2
)
head(res_rfl8_2$data)
res_rfl8_2$metadata

```

lr_parse_ttt

*Parse Avantes converted file***Description**

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

Usage

```

lr_parse_ttt(filename, ...)

lr_parse_trt(filename, ...)

lr_parse_avantes_ttt(filename, ...)

lr_parse_avantes_trt(filename, ...)

```

Arguments

filename	Path of the file to parse
...	ignored

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
 - datetime: Timestamp of the recording in format '%Y-%m-%d %H:%M:%S' and UTC timezone. If timezone is missing in source file, UTC time will be assumed (for reproducibility purposes across computers with different localtimes).
 - 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

```
res_ttt <- lr_parse_avantes_ttt(
  system.file("testdata", "avantes_export.ttt", package = "lightr")
)
head(res_ttt$data)
res_ttt$metadata

res_trt <- lr_parse_avantes_trt(
  system.file("testdata", "avantes_export2.trt", package = "lightr")
)
head(res_trt$data)
res_trt$metadata
```

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