Package: EML (via r-universe)

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Type Package

Title Read and Write Ecological Metadata Language Files

Description Work with Ecological Metadata Language ('EML') files. 'EML' is a widely used metadata standard in the ecological and environmental sciences, described in Jones et al. (2006), <doi:10.1146/annurev.ecolsys.37.091305.110031>.

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Contents

build_factors . . . 2
build_units_table . . . 3
detect_delim . . . 3
eml . . . 3
eml_get . . . 4
eml_validate . . . 4
get_attributes . . . 5
get_numberType . . . 6
get_unit_id . . . 7
get_unitList . . . 7
htmlwidgets_attributes . . . 8
is_standardUnit . . . 9
read_eml . . . 9
set_attributes . . . 10
set_coverage . . . 11
set_methods . . . 13
set_physical . . . 14
set_responsibleParty . . . 16
set_software . . . 18
set_taxonomicCoverage . . . 18
set_TextType . . . 20
set_unitList . . . 21
shiny_attributes . . . 22
table_to_r . . . 23
write_eml . . . 23

Index **25**

`build_factors` *build factor table*

Description

builds factor table for shiny app

Usage

```
build_factors(att_table, data)
```

Arguments

`att_table`	(data.frame) input attributes table
`data`	(data.frame) input data

build_units_table *build units table*

Description

builds unit table for shiny app

Usage

```
build_units_table(in_units, eml_units)
```

Arguments

in_units	input units
eml_units	eml units

detect_delim *Automatically detect line delimiters in a text file*

Description

This helper function was written expressly for set_physical to be able to automate its recordDelimiter argument.

Usage

```
detect_delim(path, nchar = 1000)
```

Arguments

path	(character) File to search for a delimiter
nchar	(numeric) Maximum number of characters to read from disk when searching

Value

(character) If found, the delimiter, it not, \r\n

eml *eml*

Description

eml

Format

A list with constructor functions

eml_get *eml_get*

Description

eml_get

Usage

```
eml_get(x, element, from = "list", ...)
```

Arguments

x	an EML object or child/descendant object
element	name of the element to be extracted. If multiple occurrences are found, will extract all
from	explicit type for the input format. Possible values: "xml", "json", "list", or "guess" with "list" as the default.
...	additional arguments

Examples

```
f <- system.file("tests", emld::eml_version(), "eml-datasetWithUnits.xml", package = "emld")
eml <- read_eml(f)
eml_get(eml, "physical")
eml_get(eml, "attributeList")

## The first argument need not be an "eml" class, it could be a child element; e.g.
eml_get(eml$dataset$dataTable, "physical")
```

eml_validate *eml_validate*

Description

eml_validate processes an EML document using the XSD schema for the appropriate version of EML and determines if the document is schema-valid as defined by the XSD specification

Usage

```
eml_validate(eml, encoding = "UTF-8", schema = NULL)
```

Arguments

eml	file path, xml_document,
encoding	optional encoding for files, default UTF-8.
schema	path to schema

Value

Whether the document is valid (logical)

Note

this function is simply an alias to ‘eml_validate‘ in ‘emld‘ package

Examples

```
f <- system.file("extdata", "example.xml", package = "emld")

## validate file directly from disk:
eml_validate(f)

## validate an eml object:
eml <- read_eml(f)
eml_validate(eml)
```

---

get_attributes *get_attributes*

Description

get_attributes

Usage

```
get_attributes(x, eml = NULL)
```

Arguments

x	an "attributeList" element from an emld object
eml	The full eml document, needed only if <references> outside of attributes must be resolved.

Details

EML metadata can use "references" elements which allow one attribute to use metadata declared elsewhere in the document. This function will automatically resolve these references and thus infer the correct metadata.

Value

a data frame whose rows are the attributes (names of each column in the data file) and whose columns describe metadata about those attributes. By default separate tables are given for each type

Examples

```
f <- system.file("tests", emld::eml_version(),
  "eml-datasetWithAttributelevelMethods.xml", package = "emld")
eml <- read_eml(f)
get_attributes(eml$dataset$dataTable$attributeList)
```

---

get_numberType *Get EML numberType*

Description

returns the EML numberType (either 'real', 'integer', 'whole', or 'natural') of input values

Usage

```
get_numberType(values)
```

Arguments

values	(numeric/character) a vector of values, if vector is non-numeric will return NA

Value

the numberType of values (either 'real', 'integer', 'whole', or 'natural').

Examples

```
## Not run:
# To get numberType for each column in a data.frame:

unlist(lapply(df, function(x) get_numberType(x)))

## End(Not run)
```

get_unit_id *get_unit_id*

Description

A function to assist with getting valid EML unit ids (see examples). Warning: ensure returned unit is equivalent to input unit (for example "pH" will return "picohenry" which may or may not be equivalent to the input unit "pH").

Usage

```
get_unit_id(input_units, eml_version = emld::eml_version())
```

Arguments

input_units	(character|vector) input units that needs valid EML unit ids
eml_version	(character) the eml schema version desired (there is a change in the way eml units are named from eml-2.1.1 to eml-2.2.0)

Value

(character) A valid EML unit id. If no valid EML unit id can be found, the function will output a warning, along with a preformatted custom unit id.

Examples

```
## Not run:
# The following all return the same id
get_unit_id("kilometersPerSquareSecond")
get_unit_id("kilometerPerSecondSquared")
get_unit_id("Kilometers per seconds squared")
get_unit_id("km/s^2")
get_unit_id("km s-2")
get_unit_id("s-2 /     kilometers-1") # this works but is not advised

## End(Not run)
```

get_unitList *get_unitList*

Description

get_unitList

Usage

```
get_unitList(x = NULL)
```

Arguments

x	an emld object

Details

If no unitList is provided, the function reads in the eml-unitDictionary defining all standard units and unitTypes. This provides a convenient way to look up standard units and their EML-recognized names when defining metadata, e.g. in the table passed to ‘set_attributes()‘.

Value

a list with two data.frames: "units", a table defining unit names, types, and conversions to SI, and "unitTypes", defining the type of unit. For instance, the unit table could define "Hertz" as a unit of unitType frequency, and the unitType define frequency as a type whose dimension is 1/time.

Examples

```
# Read in additional units defined in a EML file

f <- system.file("tests", emld::eml_version(),
  "eml-datasetWithUnits.xml",
  package = "emld"
)
eml <- read_eml(f)
unitList <- get_unitList(eml)

## Read in the definitions of standard units:
get_unitList()
```

---

htmlwidgets_attributes

Launch attributes htmlwidget

Description

Used to call handsontable html widget to build attributes

Usage

```
htmlwidgets_attributes(df, type = NULL)
```

Arguments

df	(data.frame) the data.frame of data that needs an attribute table
type	(character) either "attributes", "units", or "factors"

is_standardUnit *is_standardUnit*

Description

is_standardUnit

Usage

```
is_standardUnit(x)
```

Arguments

x	name of unit to check

Value

TRUE if unit is exact match to the id of a unit in the Standard Units Table, FALSE otherwise.

Examples

```
is_standardUnit("amperePerMeter") # TRUE
is_standardUnit("speciesPerSquareMeter") # FALSE
```

read_eml *read_eml*

Description

Read an EML file into R as an emld object.

Usage

```
read_eml(x, from = "xml")
```

Arguments

x	path to an EML file
from	explicit type for the input format. Possible values: "xml", "json", "list", or "guess" with "xml" as the default.

Value

an emld object (list / S3 object)

Examples

```
f <- system.file("extdata", "example.xml", package = "emld")
eml <- read_eml(f)
```

set_attributes *set_attributes*

Description

set_attributes

Usage

```
set_attributes(
  attributes,
  factors = NULL,
  col_classes = NULL,
  missingValues = NULL
)
```

Arguments

attributes	a joined table of all attribute metadata
factors	a table with factor code-definition pairs; see details
col_classes	optional, list of R column classes ('ordered', 'numeric', 'factor', 'Date', or 'character', case sensitive) will let the function infer missing 'domain' and 'measurementScale' values for attributes column. Should be in same order as attributeNames in the attributes table, or be a named list with names corresponding to attributeNames in the attributes table.
missingValues	optional, a table with missing value code-deinition pairs; see details

Details

The attributes data frame must use only the recognized column headers shown here. The attributes data frame must contain columns for required metadata. These are:

For all data:

- attributeName (required, free text field)
- attributeDefinition (required, free text field)
- measurementScale (required, "nominal", "ordinal", "ratio", "interval", or "dateTime", case sensitive) but it can be inferred from col_classes.
- domain (required, "numericDomain", "textDomain", "enumeratedDomain", or "dateTimeDomain", case sensitive) but it can be inferred from col_classes.

For numeric (ratio or interval) data:

- unit (required). Unitless values should use "dimensionless" as the unit.

For character (textDomain) data:

- definition (required)

For dateTime data:

- formatString (required)

Other optional allowed columns in the attributes table are: source, pattern, precision, numberType, missingValueCode, missingValueCodeExplanation, attributeLabel, storageType, minimum, maximum

The **factors** data frame, required for attributes in an enumerated domain, must use only the following recognized column headers:

- attributeName (required)
- code (required)
- definition (required)

The **missingValues** data frame, optional, can be used in the case that multiple missing value codes need to be set for the same attribute. This table must contain the following recognized column headers.

- attributeName (required)
- code (required)
- definition (required)

Value

an eml "attributeList" object

`set_coverage` *set_coverage*

Description

set_coverage

Usage

```
set_coverage(
  beginDate = character(),
  endDate = character(),
  date = character(),
  sci_names = character(),
  geographicDescription = character(),
  westBoundingCoordinate = numeric(),
  eastBoundingCoordinate = numeric(),
  northBoundingCoordinate = numeric(),
  southBoundingCoordinate = numeric(),
  altitudeMinimum = numeric(),
  altitudeMaximum = numeric(),
  altitudeUnits = character()
)
```

Arguments

beginDate	Starting date for temporal coverage range.
endDate	End date for temporal coverage range
date	give a single date, or vector of single dates covered (instead of beginDate and endDate)
sci_names	string (space separated) or list or data frame of scientific names for species covered. See details
geographicDescription	text string describing the geographic location
westBoundingCoordinate	Decimal longitude for west edge bounding box
eastBoundingCoordinate	Decimal longitude for east edge bounding box
northBoundingCoordinate	Decimal latitude value for north of bounding box
southBoundingCoordinate	Decimal latitude value for south edge of bounding box
altitudeMinimum	minimum altitude covered by the data (optional)
altitudeMaximum	maximum altitude covered by the data (optional)
altitudeUnits	name of the units used to measure altitude, if given

Details

set_coverage provides a simple and concise way to specify most common temporal, taxonomic, and geographic coverage metadata. For certain studies this will not be well suited, and users will need the more flexible but more verbose construction using "new()" methods; for instance, to specify temporal coverage in geological epoch instead of calendar dates, or to specify taxonomic coverage in terms of other ranks or identifiers.

Value

a coverage object for EML

Note

If "sci_names" is a data frame, column names of the data frame are rank names. For user-defined "sci_names", users must make sure that the order of rank names they specify is from high to low. Ex. "Kingdom","Phylum","Class","Order","Family","Genus","Species","Common"

Examples

```
coverage <-
  set_coverage(
    begin = "2012-06-01", end = "2013-12-31",
    sci_names = "Sarracenia purpurea",
```

```
    geographicDescription = "California coast, down through Baja, Mexico",
    west = -122.44, east = -117.15,
    north = 37.38, south = 30.00
  )
```

---

set_methods *set_methods*

Description

set_methods

Usage

```
set_methods(
  methods_file,
  instrumentation = character(),
  software = NULL,
  sampling_file = NULL,
  sampling_coverage = NULL,
  sampling_citation = NULL,
  qualityControl_file = NULL
)
```

Arguments

methods_file	Path to a file (markdown or .docx) containing a description of the methods used
instrumentation	optional, text describing instrumentation used in methods
software	optional, an EML software node describing software used in methods
sampling_file	optional, Path to a file (.md or .docx) describing sampling method
sampling_coverage	optional, coverage node for methods, e.g. set_coverage()
sampling_citation	optional, a citation element describing the sampling protocol
qualityControl_file	optional, path to a file (.md or .docx) describing quality control methods

Value

A methods object

Examples

```
f <- system.file("examples/hf205-methods.md", package = "EML")
set_methods(methods_file = f)

## Can also import from methods written in a .docx MS Word file.
f <- system.file("examples/hf205-methods.docx", package = "EML")
set_methods(methods_file = f)
```

`set_physical` *set_physical*

Description

Will calculate the file size, checksum, and checksum authentication method algorithm automatically if the argument `objectName` is a file that exists.

Usage

```
set_physical(
  objectName,
  id = character(),
  numHeaderLines = character(),
  numFooterLines = character(),
  recordDelimiter = detect_delim(objectName),
  fieldDelimiter = ",",
  collapseDelimiters = logical(),
  literalCharacter = character(),
  quoteCharacter = character(),
  attributeOrientation = "column",
  size = NULL,
  sizeUnit = "bytes",
  authentication = NULL,
  authMethod = NULL,
  characterEncoding = character(),
  encodingMethod = character(),
  compressionMethod = character(),
  url = character()
)
```

Arguments

`objectName`	name for the object, usually a filename like "hf205-1.csv"
`id`	optional, an id value for the <physical> element in EML, for use in referencing
`numHeaderLines`	Number of header lines preceding data. Lines are determined by the physicalLineDelimiter, or if it is absent, by the recordDelimiter. This value indicated the number of header lines that should be skipped before starting to parse the data.

`numFooterLines` Number of footer lines following data. Lines are determined by the physicalLineDelimiter, or if it is absent, by the recordDelimiter. This value indicated the number of footer lines that should be skipped after parsing the data. If this value is omitted, parsers should assume the data continues to the end of the data stream.

`recordDelimiter`

This element specifies the record delimiter character when the format is text. The record delimiter is usually a linefeed (\n) on UNIX, a carriage return (\r) on MacOS, or both (\r\n) on Windows/DOS. Multiline records are usually delimited with two line ending characters, for example on UNIX it would be two linefeed characters (\n\n). As record delimiters are often non-printing characters, one can use either the special value "\n" to represent a linefeed (ASCII 0x0a) and "\r" to represent a carriage return (ASCII 0x0d). Alternatively, one can use the hex value to represent character values (e.g., 0x0a).

`fieldDelimiter` "," character by default (for csv files). This element specifies a character to be used in the object for indicating the ending column for an attribute. The delimiter character itself is not part of the attribute value, but rather is present in the column following the last character of the value. Typical delimiter characters include commas, tabs, spaces, and semicolons. The only time the fieldDelimiter character is not interpreted as a delimiter is if it is contained in a quoted string (see quoteCharacter) or is immediately preceded by a literalCharacter. Non-printable quote characters can be provided as their hex values, and for tab characters by its ASCII string "\t". Processors should assume that the field starts in the column following the previous field if the previous field was fixed, or in the column following the delimiter from the previous field if the previous field was delimited.

`collapseDelimiters`

The collapseDelimiters element specifies whether sequential delimiters should be treated as a single delimiter or multiple delimiters. An example is when a space delimiter is used; often there may be several repeated spaces that should be treated as a single delimiter, but not always. The valid values are yes or no. If it is set to yes, then consecutive delimiters will be collapsed to one. If set to no or absent, then consecutive delimiters will be treated as separate delimiters. Default behavior is no; hence, consecutive delimiters will be treated as separate delimiters, by default.

`literalCharacter`

This element specifies a character to be used for escaping special character values so that they are treated as literal values. This allows "escaping" for special characters like quotes, commas, and spaces when they are intended to be used in an attribute value rather than being intended as a delimiter. The literalCharacter is typically a \.

`quoteCharacter` This element specifies a character to be used in the object for quoting values so that field delimiters can be used within the value. This basically allows delimiter "escaping". The quoteChacter is typically a " or '. When a processor encounters a quote character, it should not interpret any following characters as a delimiter until a matching quote character has been encountered (i.e., quotes come in pairs). It is an error to not provide a closing quote before the record ends. Non-printable quote characters can be provided as their hex values.

`attributeOrientation`
: Specifies whether the attributes described in the physical stream are found in columns or rows. The valid values are column or row. If set to 'column', then the attributes are in columns. If set to 'row', then the attributes are in rows. Row orientation is rare.

`size`
: This element contains information of the physical size of the entity, by default represented in bytes unless the sizeUnit attribute is provided to change the units.

`sizeUnit`
: the unit in which size is measured; default is 'bytes'

`authentication`
: This element describes authentication procedures or techniques, typically by giving a checksum value for the object. The method used to compute the authentication value (e.g., MD5) is listed in the method attribute.

`authMethod`
: the method for authentication checksum, e.g. MD5

`characterEncoding`
: This element contains the name of the character encoding. This is typically ASCII or UTF-8, or one of the other common encodings.

`encodingMethod`
: This element lists a encoding method used to encode the object, such as base64, BinHex.

`compressionMethod`
: This element lists a compression method used to compress the object, such as zip, compress, etc. Compression and encoding methods must be listed in the order in which they were applied, so that decompression and decoding should occur in the reverse order of the listing. For example, if a file is compressed using zip and then encoded using MIME base64, the compression method would be listed first and the encoding method second.

`url`
: optional. The complete url from which the data file can be downloaded, if possible.

Value

an EML physical object, such as used in a dataTable element to define the format of the data file.

Examples

```
set_physical("hf205-01-TPexp1.csv")
# FIXME set recordDelimiter based on user's system?
# FIXME richer distribution options? use set_distribution at top level?
```

`set_responsibleParty` *set_responsibleParty*

Description

set_responsibleParty

Usage

```
set_responsibleParty(
  givenName = NULL,
  surName = NULL,
  organizationName = NULL,
  positionName = NULL,
  address = NULL,
  phone = NULL,
  electronicMailAddress = NULL,
  onlineUrl = NULL,
  userId = NULL,
  id = NULL,
  email = NULL
)
```

Arguments

givenName	individual's given names (list or vector for multiple names). OR a person object.
surName	individual name
organizationName	if party is an organization instead of an individual, name for the org
positionName	individual's position, i.e. "Researcher", "Graduate Student", "Professor"
address	address object, see 'eml$address' to build an address object
phone	individual or organization phone number
electronicMailAddress	email address (alternatively, can use 'email' argument)
onlineUrl	a URL to the homepage of the individual or organization
userId	the user's ID, usually within a particular system (KNB, DataONE)
id	Identifier for this block, ideally an ORCID id (optional)
email	alias for electronicMailAddress

Value

A emld object for any responsibleParty (e.g. creator, contact, etc)

Examples

```
# Pass in either a person object or separate values to create an individual
carl <- set_responsibleParty(as.person("Carl Boettiger <cboettig@ropensci.org>"))
matt <- set_responsibleParty("Matthew", "Jones", email = "mbjones@nceas.ucsb.edu")

# To create an organization, use the named `organization` argument to
# specify the organization name
my_org <- set_responsibleParty(
  organization = "My Organization",
  email = "contact@example.org"
)
```

set_software *set_software*

Description

set_software

Usage

```
set_software(codemeta)
```

Arguments

codemeta	codemeta object, see examples

Value

an eml software element

Examples

```
cm <- jsonlite::read_json(system.file("extdata/codemeta.json", package = "EML"))
software <- set_software(cm)
my_eml <- eml$eml(packageId = "eml-1.2", system = "knb", software = software)

# write_eml(my_eml, "test.xml")
```

set_taxonomicCoverage *set_taxonomicCoverage*

Description

set_taxonomicCoverage

Usage

```
set_taxonomicCoverage(sci_names, expand = FALSE, db = "itis")
```

Arguments

sci_names	string (space separated) or list or data frame of scientific names for species covered.
expand	Set to TRUE to use '[taxadb]' to expand sci_names into full taxonomic classifications
db	The taxonomic database to query (when expand is set to TRUE). See '[taxadb::filter_name]' for valid options. Defaults to 'itis'.

Details

Turn a data.frame or a list of scientific names into a taxonomicCoverage block sci_names can be a space-separated character string or a data frame with column names as rank name or a list of user-defined taxonomicClassification

Value

a taxonomicCoverage object for EML

Note

If "sci_names" is a data frame, column names of the data frame are rank names. For user-defined "sci_names", users must make sure that the order of rank names they specify is from high to low. Ex. "Kingdom","Phylum","Class","Order","Family","Genus","Species","Common" EML permits any rank names provided they go in descending order.

Examples

```
taxon_coverage <- set_taxonomicCoverage("Macrocystis pyrifera")

sci_names <- data.frame(
  Kingdom = "Plantae",
  Phylum = "Phaeophyta",
  Class = "Phaeophyceae",
  Order = "Laminariales",
  Family = "Lessoniaceae",
  Genus = "Macrocystis",
  specificEpithet = "pyrifera"
)
taxon_coverage <- set_taxonomicCoverage(sci_names)

 # Examples that may take > 5s

## use a list of lists for multiple species
sci_names <- list(list(
  Kingdom = "Plantae",
  Phylum = "Phaeophyta",
  Class = "Phaeophyceae",
  Order = "Laminariales",
  Family = "Lessoniaceae",
  Genus = "Macrocystis",
  specificEpithet = "pyrifera"
))
set_taxonomicCoverage(sci_names)
```

set_TextType *set_TextType*

Description

For any EML element of class TextType, this function can be used to generate the appropriate EML from a markdown-formatted file.

Usage

```
set_TextType(file = NULL, text = NULL)
```

Arguments

file	path to a file providing formatted input text, see details.
text	a plain text character string which will be used directly as the content of the node if no file is given

Details

If the ‘rmarkdown‘ package is installed, then the input file can be a Microsoft Word (.docx) file, a markdown file, or other file recognized by Pandoc (see https://pandoc.org), which will automate the conversion to a docbook. Otherwise, the input file should already be in docbook format (with .xml or .dbk extension). Note that pandoc comes pre-installed in RStudio and is required for the rmarkdown package.

Value

a TextType object that can be coerced into any element inheriting from TextType, see examples

Examples

```
## using a simple character string
a <- set_TextType(text = "This is the abstract")

## Using an external markdown file
f <- system.file("examples/hf205-abstract.md", package = "EML")
a <- set_TextType(f)

## Can also import from methods written in a .docx MS Word file.
f <- system.file("examples/hf205-abstract.docx", package = "EML")
a <- set_TextType(f)

## Documents with title headings use `section` instead of `para` notation
f <- system.file("examples/hf205-methods.docx", package = "EML")
d <- set_TextType(f)
```

set_unitList *set_unitList*

Description

Define custom units, including new unitTypes. Note that it is not necessary to define most common units.

Usage

```
set_unitList(units, unitTypes = NULL, as_metadata = FALSE)
```

Arguments

units	a data.frame describing the custom units, see details.
unitTypes	optional, a data.frame defining any additional unitTypes not already defined
as_metadata	logical, default FALSE. If true, returns an 'additionalMetadata' element, see below.

Details

The units data.frame must have the following columns: - id: the referenced name of unit (singular). e.g. 'meter', 'second' - unitType: the base type of unit, e.g. 'length'. If not from a standard type, a new unitType must be provided - multiplierToSI: the multiplicative constant to convert to the SI unit. - parentSI: the name of the parent SI unit, e.g. second. - description: a text string describing the unit of measure. The following columns are optional: - name: usually the same as the id of the unit, e.g. second - abbreviation: common abbreviation, e.g. s - constantToSI: an additive constant to convert to the equivalent SI unit. If not given, default is "0"

In practice, researchers may save these tables of custom units they frequently use in an external .csv or other format and read them in to R for ready re-use.

The unitType table must have the following columns: - id: the name by which the unitType is referred to. - name: optional, default is same as the id - dimension: name of a base dimension of the unit - power: the power to which the dimension is raised (NA implies power of 1)

Value

unitList list object

Examples

```
## create the "unitType" table for custom unit
id <- c("speed", "speed", "acceleration", "acceleration", "frequency")
dimension <- c("length", "time", "length", "time", "time")
power <- c(NA, "-1", NA, "-2", "-1")
unitTypes <- data.frame(
  id = id, dimension = dimension,
  power = power, stringsAsFactors = FALSE
```

```
)

## Create the units table
id <- c("minute", "centimeter")
unitType <- c("time", "length")
parentSI <- c("second", "meter")
multiplierToSI <- c("0.0166", "1")
description <- c("one minute is 60 seconds", "centimeter is a 100th of a meter")
units <- data.frame(
  id = id, unitType = unitType, parentSI = parentSI,
  multiplierToSI = multiplierToSI, description = description,
  stringsAsFactors = FALSE
)

unitList <- set_unitList(units, unitTypes)
```

`shiny_attributes` *Create/Edit EML attributes*

Description

Create/edit EML attributes, custom units, and factors in a shiny environment.

Usage

```
shiny_attributes(data = NULL, attributes = NULL)
```

Arguments

`data`	(data.frame) the data.frame of data that needs an attribute table
`attributes`	(data.frame) an existing attributes table

Details

Attributes can be created from scratch using `shiny_attributes()`. Or an existing attribute table can be edited using `shiny_attributes(NULL, attributes)`. Or new attributes can be created from a data table using `shiny_attributes(data, NULL)`. If attributes are created from a data table, fields such as ‘attributeName‘ and ‘numberType‘ will be automatically completed based on the attributes within the data table. If both existing attributes and data table are entered (i.e. `shiny_attributes(data, attributes)`), any automatically generated fields based attributes within the data table **will not** override any non-empty fields in the entered attributes

Examples

```
## Not run:
# from scratch
out <- shiny_attributes(NULL, NULL)

# from data
```

```
data <- iris
out <- shiny_attributes(data, NULL)

# from exisiting attributes
file <- system.file("tests", emld::eml_version(),
  "eml-datasetWithAttributelevelMethods.xml",
  package = "emld"
)
eml <- read_eml(file)
x <- eml$dataset$dataTable$attributeList
df <- get_attributes(x, eml)
out <- shiny_attributes(NULL, df$attributes)

# from attributes and data
out <- shiny_attributes(data, df$attributes)

## End(Not run)
```

`table_to_r` *handsontable to r*

Description

Takes a handsontable and converts to r data.frame for shiny app

Usage

```
table_to_r(table)
```

Arguments

`table`	input table

`write_eml` *write_eml*

Description

write_eml

Usage

```
write_eml(eml, file, namespaces = NULL, ns = "eml", ...)
```

Arguments

eml	an emld class object
file	file name to write XML.
namespaces	named character vector of additional XML namespaces to use.
ns	root namespace abbreviation
...	additional arguments to write_xml

Value

If file is not specified, the result is a character string containing the resulting XML content. Otherwise return silently.

Examples

```
f <- system.file("extdata", "example.xml", package = "emld")
eml <- read_eml(f)
write_eml(eml, "test.xml")
eml_validate("test.xml")
unlink("test.xml") # clean up
```

Index

∗ **datasets**
eml, 3

build_factors, 2
build_units_table, 3

detect_delim, 3

eml, 3
eml_get, 4
eml_validate, 4

get_attributes, 5
get_numberType, 6
get_unit_id, 7
get_unitList, 7

htmlwidgets_attributes, 8

is_standardUnit, 9

read_eml, 9

set_attributes, 10
set_coverage, 11
set_methods, 13
set_physical, *3*, 14
set_responsibleParty, 16
set_software, 18
set_taxonomicCoverage, 18
set_TextType, 20
set_unitList, 21
shiny_attributes, 22

table_to_r, 23

write_eml, 23
write_xml, *24*