

Package ‘rezonateR’

July 23, 2021

Title A Support Package for Working with Rezonator in R

Version 0.0.0.9000

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Description This is a package, still in very early alpha, that a) takes data annotated in Rezonator and turns it into a format useable in R, and b) moulds data in formats currently unsupported by Rezonator into a shape importable into Rezonator.

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Encoding UTF-8

LazyData true

Roxygen list(markdown = TRUE)

RoxygenNote 7.1.1

Suggests testthat (>= 3.0.0)

Config/testthat/edition 3

Imports xml2,
rjson

Depends dplyr,
readr

R topics documented:

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importRez	<i>Import a Rez file</i>
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Description

Import a Rez file. This returns an object containing, among other things, a nodeMap object containing raw information, and data frames for tokens, units, chunks, track chain entries, track chains, containing only key information likely to be useful for the user.

Usage

```
importRez(paths, docnames = "", concatFields = c("word", "wordWylie", "lit"))
```

Arguments

docnames	A character vector of the document names. If left blank, a docname will be generated according to the filenames of files you import. For example, the document foo/bar.rez will be named 'bar'.
path	A character vector of paths to the files to be imported. For Windows users, please use / instead of \.

Value

rezRObj

rez_pangloss	<i>Read Pangloss file</i>
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Description

Read Pangloss file

Usage

```
rez_pangloss(url, filename)
```

Arguments

url	A URL for the Pangloss source.
filename	a filename for the export.

Value

rezImport

Examples

```
rez_pangloss("https://cocoon.huma-num.fr/data/mazaudon/masters/crdo-TAJ_ORIGTAM.xml", "tamang1.txt")
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