Package 'IalsaSynthesis'

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Title Synthesizing Information Across Collaborating Research
Description Synthesizing information across collaborating research. Created for Integrative Analysis of Longitudinal Studies of Aging (http://www.ialsa.org/).
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<pre>URL https://github.com/IALSA/IalsaSynthesis, http://www.ialsa.org/</pre>
BugReports https://github.com/IALSA/IalsaSynthesis/issues
Depends R(>= 3.0.0), stats
Imports testit
Suggests devtools, knitr, readr, testthat (>= 0.9)
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LazyData TRUE
VignetteBuilder knitr
Roxygen list(wrap = TRUE)
R topics documented:
extract
Index

2 IalsaSynthesis

extract

Extract the values within model output files.

Description

Functions that extract the values within model output files.

Usage

```
extract_output_filename(mplus_output)
```

Arguments

mplus_output Text containing model output.

Value

A numeric value corresponding to the desired quantity.

Author(s)

Will Beasley

Examples

library(IalsaSynthesis) #Load the package into the current R session.

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Description

IalsaSynthesis.

validate 3

validate

Functions that check the validty of values throughout the workflow.

Description

These functions help identify mistakes in formatting before the create difficult-to-diagnose problems later.

Usage

```
validate_filename_output(filename, path, file_extension_expected = "out",
  underscore_count_expected = 4L)
```

Arguments

filename The name of the file to be validated.

path The location of the file to be validated.

file_extension_expected

The extension of the file. This defaults to "out", which corresponds to Mplus

output.

underscore_count_expected

The number of underscores required in the name (not currently used).

Value

An invisible TRUE value if the filename is valid. Otherwise, an error is thrown.

Author(s)

Will Beasley

Examples

```
library(IalsaSynthesis) #Load the package into the current R session.
## Not run:
path <- "./studies/eas"
good_name <- "u1_male_aehplus_muscle_noCog_hand_noCogSpec.out"
validate_filename_output(good_name, path)
bad_name <- "missing_something.outttt"
validate_filename_output(bad_name, path)
## End(Not run)</pre>
```

Index