

Package ‘IalsaSynthesis’

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Title Synthesizing Information Across Collaborating Research

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Description Synthesizing information across collaborating research. Created for Integrative Analysis of Longitudinal Studies of Aging (<http://www.ialsa.org/>).

URL <https://github.com/IALSA/IalsaSynthesis>, <http://www.ialsa.org/>

BugReports <https://github.com/IALSA/IalsaSynthesis/issues>

Depends R(>= 3.0.0),
stats

Imports testit

Suggests devtools,
knitr,
testthat (>= 0.9)

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LazyData TRUE

VignetteBuilder knitr

Roxygen list(wrap = TRUE)

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IalsaSynthesis	<i>IalsaSynthesis.</i>
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Description

IalsaSynthesis.

validate*Functions that check the validity 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

<code>filename</code>	The name of the file to be validated.
<code>path</code>	The location of the file to be validated.
<code>file_extension_expected</code>	The extension of the file. This defaults to "out", which corresponds to Mplus output.
<code>underscore_count_expected</code>	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.outtttt"  
validate_filename_output(bad_name, path)  
  
## End(Not run)
```

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