# Package 'IalsaSynthesis'

April 30, 2015
Title Synthesizing Information Across Collaborating Research
Version 0.0-2
<b>Date</b> 2015-04-30
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<b>Description</b> Synthesizing information across collaborating research. Created for Integrative Analysis of Longitudinal Studies of Aging (http://www.ialsa.org/).
<pre>URL https://github.com/IALSA/IalsaSynthesis, http://www.ialsa.org/</pre>
BugReports https://github.com/IALSA/IalsaSynthesis/issues
<b>Depends</b> R(>= 3.0.0), stats
Imports testit
Suggests devtools, knitr, testthat (>= 0.9)
License MIT + file LICENSE
LazyData TRUE
VignetteBuilder knitr
<b>Roxygen</b> list(wrap = TRUE)
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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.

IalsaSynthesis

IalsaSynthesis.

# **Description**

IalsaSynthesis.

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)
```

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# **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>
```

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