# Package 'IalsaSynthesis'

September 1, 2015

Title Synthesizing Information Across Collaborating Research
<b>Description</b> Synthesizes information across collaborating research. Created specifically for Integrative Analysis of Longitudinal Studies of Aging (IALSA).
Version 0.1.5
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Maintainer Will Beasley <wibeasley@hotmail.com></wibeasley@hotmail.com>
<pre>URL https://github.com/IALSA/IalsaSynthesis, http://www.ialsa.org/</pre>
<pre>BugReports https://github.com/IALSA/IalsaSynthesis/issues</pre>
<b>Depends</b> R(>= 3.0.0), stats
Imports testit
Suggests devtools, knitr, readr, testthat (>= 0.9)
License GPL-2
LazyData TRUE
VignetteBuilder knitr
R topics documented:
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Synthesizing Information Across Collaborating Research

# **Description**

Synthesizing information across collaborating research. Created for Integrative Analysis of Longitudinal Studies of Aging (IALSA).

#### Note

The release version will be available through CRAN. The most recent development version is available through GitHub. Please see the installation examples below.

If you're having trouble with the package, please install the development version. If this doesn't solve your problem, please create a new issue, or email Will or Andrey.

# Author(s)

William Howard Beasley – Assistant Professor of Research, University of Oklahoma Health Sciences Center, Dept of Pediatrics, Biomedical and Behavioral Methodology Core (BBMC)

Andrey Koval -Post Doc, University of Victoria, Department of Psychology

# **Examples**

```
## Not run:
# Install/update IalsaSynthesis with the release version from CRAN.
install.packages('IalsaSynthesis')

# Install/update IalsaSynthesis with the development version from GitHub
#install.packages('devtools') #Uncomment if `devtools` isn't installed already.
devtools::install_github('IALSA/IalsaSynthesis')

## End(Not run)
```

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.
```

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#### Value

A numeric value corresponding to the desired quantity.

#### Author(s)

Will Beasley

# **Examples**

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

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>
```

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