# Package 'VADIS'

October 30, 2018
Title Variation-Based Distance & Similarity Modeling
<b>Version</b> 0.0.0.9000
<b>Description</b> Provides functions for computing the three lines of evidence used in the VADIS method. Includes vignette illustrating how to use the package.
<b>Depends</b> R (>= $3.5.1$ )
License GPL-3
Encoding UTF-8
<b>Imports</b> plyr (>= 1.8), magrittr (>= 1.5), dplyr (>= 0.7.6), readr (>= 1.1), reshape2 (>= 1.4), lme4 (>= 1.1), party (>= 1.2), ranger (>= 0.10), phangorn(>= 2.4), ade4 (>= 1.7), rstan (>= 2.17), brms (>= 2.4), e1071 (>= 1.7)
<b>Suggests</b> tidyverse (>= 1.2.1), broom (>= 0.5), edarf (>= 1.1.1), ggplot2 (>= 3.0), ggeffects (>= 0.5), scales (>= 1.0), knitr (>= 1.2), rmarkdown
LazyData true
RoxygenNote 6.1.0
VignetteBuilder knitr
Author Jason Grafmiller [aut, cre]
Maintainer Jason Grafmiller <j.grafmiller@bham.ac.uk></j.grafmiller@bham.ac.uk>
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create\_coef\_table

Create Table of Coefficients for VADIS analysis

# Description

Create Table of Coefficients for VADIS analysis

# Usage

```
create_coef_table(mod_list)
```

# **Arguments**

mod\_list A list of regression model objects.

path Path in which to save the output (as .csv file). If NULL, defaults to the current

working directory. Set path = FALSE if you do not wish to save to file.

#### **Details**

The function loops through a list of model objects, extracts the coefficient estimates, and compiles them in a single dataframe.

# Value

A dataframe

# Author(s)

Jason Grafmiller

```
## Not run:
lm_fnc <- function(x) lm(Sepal.Length ~ Petal.Length + Petal.Width, data = x)
rm_list <- fit.vadis.RM(iris, split.by = "Species", fit.func = lm_fnc,
    path = FALSE)
summary(rm_list[[1]])
create_coef_table(rm_list, path = FALSE)
## End(Not run)</pre>
```

create\_rank\_table 3

create\_rank\_table

Create Table of Variable Importance Rankings for VADIS analysis

#### **Description**

Create Table of Variable Importance Rankings for VADIS analysis

#### Usage

```
create_rank_table(mod_list, conditional = FALSE)
```

#### **Arguments**

mod\_list A list of random forest model objects, generated with fit\_vadis\_rf(). Cur-

rently supports objects of RandomForest-class, ranger, and randomForest

conditional logical. Should unconditional (default) or conditional permutation variable im-

portance be computed. Only applies to RandomForest-class models from the

party package.

path Path in which to save the output (as .csv file). If NULL, defaults to the current

working directory. Set path = FALSE if you do not wish to save to file.

#### **Details**

The function loops through a list of random forest objects, extracts the variable importance estimates, and compiles them in a single dataframe. For forests fit with ranger or randomForest, the importance argument must be specified. #'

#### Value

A dataframe

#### Author(s)

Jason Grafmiller

```
## Not run:
fmla <- Type ~ PossrAnimacyBin + PossrWordC + PossmWordC + FinalSibilant +
    TypeTokenRatio + ProtoSemanticRelation + PossrExpType

rf_fnc <- function(x) ranger::ranger(fmla, data = x, importance = "permutation")

rf_list <- fit_models(brown_genitives, split.by = "Genre", fit.func = rf_fnc, path = FALSE)

create_rank_table(rf_list, path = FALSE)

## End(Not run)</pre>
```

4 create\_signif\_table

# Description

Create Table of Significant Effects for VADIS analysis

# Usage

```
create_signif_table(mod_list)
```

# **Arguments**

mod\_list A list of regression model objects.

path Path in which to save the output (as .csv file). If NULL, defaults to the current

working directory. Set path = FALSE if you do not wish to save to file.

#### **Details**

The function loops through a list of model objects, extracts the coefficient estimates, and compiles them in a single dataframe.

# Value

A dataframe

# Author(s)

Jason Grafmiller

```
## Not run:
lm_fnc <- function(x) lm(Sepal.Length ~ Petal.Length + Petal.Width, data = x)
rm_list <- fit.vadis.RM(iris, split.by = "Species", fit.func = lm_fnc,
    path = FALSE)
summary(rm_list[[1]])
create_coef_table(rm_list, path = FALSE)
## End(Not run)</pre>
```

vadis\_line1 5

vadis\_line1

Calculate the first line of evidence for the VADIS method

# Description

Calculate the first line of evidence for the VADIS method

# Usage

```
vadis_line1(mod_list, path = NULL)
```

# **Arguments**

mod\_list A list of regression model objects.

Path in which to save the output as an R data file ( .rds). If NULL, defaults to the

current working directory. Set path = FALSE if you do not wish to save to file.

#### **Details**

The function loops through a list of model objects, extracts the coefficient estimates, and compiles them in a single dataframe.

# Value

A list of length 3.

# Author(s)

Jason Grafmiller

```
## Not run:
lm_fnc <- function(x) lm(Sepal.Length ~ Petal.Length + Petal.Width + Sepal.Width, data = x)
rm_list <- fit.vadis.RM(iris, split.by = "Species", fit.func = lm_fnc,
    path = FALSE)
summary(rm_list[[1]])
line1 <- calc_line1(rm_list, path = FALSE)
## End(Not run)</pre>
```

6 vadis\_line2

vadis\_line2

Calculate the second line of evidence for the VADIS method

# Description

Calculate the second line of evidence for the VADIS method

# Usage

```
vadis_line2(mod_list, path = NULL)
```

# **Arguments**

mod\_list A list of regression model objects.

Path in which to save the output as an R data file ( .rds). If NULL, defaults to the

current working directory. Set path = FALSE if you do not wish to save to file.

#### **Details**

The function loops through a list of model objects, extracts the coefficient estimates, and compiles them in a single dataframe.

# Value

A dataframe

# Author(s)

Jason Grafmiller

```
## Not run:
lm_fnc <- function(x) lm(Sepal.Length ~ Petal.Length + Petal.Width + Sepal.width, data = x)
rm_list <- fit.vadis.RM(iris, split.by = "Species", fit.func = lm_fnc,
    path = FALSE)
summary(rm_list[[1]])
line2 <- calc_line2(rm_list, path = FALSE)
## End(Not run)</pre>
```

vadis\_line3 7

vadis_line3 Calculate the third line of evidence for the VADIS method	
---	--

# **Description**

Calculate the third line of evidence for the VADIS method

#### Usage

```
vadis_line3(mod_list, path = NULL, conditional = FALSE)
```

#### **Arguments**

mod\_list A list of random forest model objects.

path Path in which to save the output as an R data file (.rds). If NULL, defaults to the

current working directory. Set path = FALSE if you do not wish to save to file.

conditional logical. Should unconditional (default) or conditional permutation variable im-

portance be computed. Only applies to RandomForest-class models from the

party package.

#### **Details**

The function loops through a list of model objects, extracts the coefficient estimates, and compiles them in a single dataframe.

# Value

A list of length 3

# Author(s)

Jason Grafmiller

```
## Not run:
lm_fnc <- function(x) lm(Sepal.Length ~ Petal.Length + Petal.Width, data = x)
rm_list <- fit.vadis.RM(iris, split.by = "Species", fit.func = lm_fnc,
    path = FALSE)
summary(rm_list[[1]])
line3 <- calc_line3(rm_list, path = FALSE)
## End(Not run)</pre>
```

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