Package 'spectratrait'

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Title A simple add-on package to aid in the fitting of leaf-level spectra-trait PLSR models	
Version 0.9.9	
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Description Provides functions to conduct standardized leaf-level spectra-trait PLSR model fitting is cluding uncertainty analysis that follow DOI: https://doi.org/10.1111/nph.16123	n-
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create_data_split

Create a calibration (training) / validation data split for PLSR model fitting and testing

Description

Create a calibration (training) / validation data split for PLSR model fitting and testing

Usage

```
create_data_split(
  dataset = NULL,
  approach = NULL,
  split_seed = 123456789,
  prop = 0.8,
  group_variables = NULL
)
```

Arguments

dataset input full PLSR dataset to split into cal/val datasets

approach approach to splitting the dataset. Options: base or dplyr

split_seed random seed to use for splitting data

prop the proportion of data to preserve for calibration (e.g. 0.8) and validation (0.2).

This sets the calibration proportion

group_variables

Use factor variables to conduct a stratfied sampling for cal/val

Value

output_list A list containing the calibration dataset (cal_data) and validation dataset (val_data)

Author(s)

Julien Lamour, Jeremiah Anderson, Shawn P. Serbin

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f.coef.valid f.coef.valid

Description

f.coef.valid

Usage

```
f.coef.valid(plsr.out, data_plsr, ncomp, inVar)
```

Arguments

plsr.out plsr model obtained with jaccknife = TRUE

data_plsr data used for the plsr model with Spectra the matrix of spectra

ncomp number of selection components

inVar Name of the PLSR model response variable

Value

B returns the intercept and the coefficients of the jackknife or bootstrap validation

Author(s)

Julien Lamour

f.plot.coef f.plot.coef

Description

f.plot.coef

Usage

```
f.plot.coef(
  Z,
  wv,
  xlim = NULL,
  position = "topright",
  type = "Coefficient",
  plot_label = NULL
)
```

f.plot.spec

Arguments

Z Coefficient matrix with each row corresponding to the coefficients and wave-

length in columns

wv vector of wavelengths

xlim vector to change the default xlim of the plots (ex xlim = c(500, 2400))

position Position of the legend (see base function legend for help)

type Name of the y axis and of the legend

plot_label optional plot label to include with the figure

Author(s)

Julien Lamour

f.plot.spec

f.plot.spec

Description

f.plot.spec

Usage

```
f.plot.spec(
   Z,
   wv,
   xlim = NULL,
   position = "topright",
   type = "Reflectance",
   plot_label = NULL
)
```

Arguments

Z Spectra matrix with each row corresponding to a spectra and wavelength in

columns

wv vector of wavelengths corresponding to the column of the spectra matrix Z

xlim vector to change the default xlim of the plots (ex xlim = c(500, 2400))

position Position of the legend (see base function legend for help)

type Name of the y axis and of the legend. E.g. Reflectance, Transmittance

plot_label optional plot label to include with the figure

Author(s)

Julien Lamour, Shawn P. Serbin

```
find_optimal_components
```

Apply different methods to determing the optimal number of PLSR model components

Description

Apply different methods to determing the optimal number of PLSR model components

Usage

```
find_optimal_components(
  dataset = NULL,
  method = "pls",
  maxComps = 20,
  iterations = 20,
  seg = 100,
  prop = 0.7,
  random_seed = 123456789
)
```

Arguments

dataset	input full PLSR dataset. Usually just the calibration dataset
method	Which approach to use to find optimal components. Options: pls, firstPlateau, firstMin
maxComps	maximum number of components to consider
iterations	how many different permutations to run
seg	For the built-in pls method, how many different data segments to select from the input dataset
prop	proportion of data to preserve for each permutation
random_seed	random seed to use for splitting data

Value

nComps the optimal number of PLSR components

Author(s)

Julien Lamour, Jeremiah Anderson, Shawn P. Serbin

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get_ecosis_data

Function to pull data from EcoSIS using the EcoSIS API

Description

Function to pull data from EcoSIS using the EcoSIS API

Usage

```
get_ecosis_data(ecosis_id = NULL)
```

Arguments

ecosis_id

the alphanumeric EcoSIS API dataset ID

Value

EcoSIS spectral dataset object

Author(s)

Shawn P. Serbin, Alexey Shiklomanov

Examples

```
## Not run:
ecosis_id <- "960dbb0c-144e-4563-8117-9e23d14f4aa9"
dat_raw <- get_ecosis_data(ecosis_id = ecosis_id)
head(dat_raw)
names(dat_raw)[1:40]
## End(Not run)</pre>
```

pls_permutation

Run a PLSR model permutation analysis. Can be used to determine the optimal number of components or conduct a boostrap uncertainty analysis

Description

See Serbin et al. (2019). DOI: https://doi.org/10.1111/nph.16123

Usage

```
pls_permutation(
  dataset = NULL,
  maxComps = 20,
  iterations = 20,
  seg = 100,
  prop = 0.7
)
```

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Arguments

dataset input full PLSR dataset. Usually just the calibration dataset maxComps maximum number of components to use for each PLSR fit

iterations how many different permutations to run

seg currently unused - should be removed from this function call

prop proportion of data to preserve for each permutation

Author(s)

Julien Lamour, Shawn P. Serbin

source_GitHubData

Function to source text data from GitHub

Description

Function to source text data from GitHub

Usage

```
source_GitHubData(url, sep = ",", header = TRUE)
```

Arguments

url http/https URL to the github dataset

sep dataset file delimiter

header TRUE/FALSE does the file have a column header?

Author(s)

gist.github.com/christophergandrud/4466237

testForPackage

Function to check for installed package not presently used

Description

Function to check for installed package not presently used

Usage

testForPackage(pkg)

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VIP

VIP returns all VIP values for all variables and all number of components, as a ncomp x nvars matrix.

Description

VIP returns all VIP values for all variables and all number of components, as a ncomp x nvars matrix.

Usage

```
VIP(object)
```

VIPjh

VIPjh returns the VIP of variable j with h components

Description

VIPjh returns the VIP of variable j with h components

Usage

```
VIPjh(object, j, h)
```

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