Package 'novelforestSG'

October 13, 2022

Title Dataset from the Novel Forests of Singapore

Version 2.0.0

Description The raw dataset and model used in Lai et al. (2021) Decoupled responses of native and exotic tree diversities to distance from old-growth forest and soil phosphorous in novel secondary forests. Applied Vegetation Science, 24, e12548.

License CC BY 4.0

URL https://hrlai.github.io/novelforestSG/,

https://github.com/hrlai/novelforestSG,

https://doi.org/10.1111/avsc.12548

BugReports https://github.com/hrlai/novelforestSG/issues

Encoding UTF-8

LazyData true

RoxygenNote 7.1.2

Depends R (>= 2.10)

Suggests brms (>= 2.10.0), testthat

NeedsCompilation no

Author Hao Ran Lai [aut, cre] (<https://orcid.org/0000-0001-6871-0146>), Kwek Yan Chong [aut] (<https://orcid.org/0000-0003-4754-8957>), Alex Thiam Koon Yee [aut] (<https://orcid.org/0000-0002-6465-0075>), Germaine Su Yin Tan [ctb], Louise Neo [ctb], Carmen Yingxin Kee [ctb], Hugh Tiang Wah Tan [ths]

Maintainer Hao Ran Lai <hrlai.ecology@gmail.com>

Repository CRAN

Date/Publication 2022-10-03 21:20:02 UTC

R topics documented:

backtransform																							•	•			2
download_model																				•							3
novelforestSG																											4
novelforest_data .					•	•	•	•	•			•	•	•		•	•	•		•	•	•			•	•	5
																											6

Index

backtransform

Backtransform Scaled Predictors

Description

Backtransform scaled predictors in the input data (obtained via download_model()) to their original scales. This is done by first back-scaling to the log-scale, and then backtranformed to the original scale (the predictors were log-transformed as described in Lai et al. 2021).

Usage

```
backtransform(data)
```

Arguments

data

Defaults to the data object in download_model() (see Examples), but could also be another data.frame with the same predictor names, should you wish to (back)scale and (back)center using the same means and standard deviations for any reason.

Value

A backtransformed data. frame with predictors at their original scales.

References

Lai, H.R., Tan, G.S.Y., Neo, L., Kee, C.Y., Yee, A.T.K., Tan, H.T.W. and Chong, K.Y. (2021) Decoupled responses of native and exotic tree diversities to distance from old-growth forest and soil phosphorous in novel secondary forests. *Applied Vegetation Science*, 24, e12548. doi:10.1111/avsc.12548

Examples

```
# download the model object containing input data
novelforest_model <- download_model()</pre>
```

```
dat <- backtransform(novelforest_model$data)
head(dat)</pre>
```

download_model

Description

Download the brms model fitted to novelforest_data (Lai et al. 2021). The model object is too large (16.5 MB) to be included with the package, so this function downloads the model from the developmental GitHub website. The generalised linear mixed-effect model was fitted via brms::brm so this package is recommended to make full use of the model object.

Usage

download_model(save_to = NULL)

Arguments

save_to	Path and name of the file where the R object is saved to. Defaults to NULL,
	which does not save the model object locally.

Value

A brms model output of class brmsfit, which is a list containing the input data and other slots that store the model components.

Notably, the data slot contains a data. frame with the following response variables:

SD_N_0 first-order native taxonomic diversity, i.e., species richness

SD_N_2 second-order native taxonomic diversity, i.e., inverse Simpson index

SD_E_0 first-order exotic taxonomic diversity

SD_E_2 second-order exotic taxonomic diversity

FD_N_0 first-order native functional diversity

- FD_N_2 second-order native functional diversity
- **FD_E_0** first-order exotic functional diversity

FD_E_2 second-order exotic functional diversity,

and the following explanatory variables (and measurement units if you backtransform them using backtransform):

dist Distance to old-growth forests (m)

size Patch area (km^2)

nitrogen Total soil nitrogen (mg/kg)

phosphorous Total extractable soil phosphorous (mg/kg)

potassium Total extractable soil potassium (mg/kg)

patch Forest patch ID

#' Note that all explanatory variables were log-transformed and standardised to zero mean and unit standard deviations. Use backtransform to obtain the variables in their original scales. See Lai et al. (2021) for more details on model building and data collection.

References

Lai, H.R., Tan, G.S.Y., Neo, L., Kee, C.Y., Yee, A.T.K., Tan, H.T.W. and Chong, K.Y. (2021) Decoupled responses of native and exotic tree diversities to distance from old-growth forest and soil phosphorous in novel secondary forests. *Applied Vegetation Science*, 24, e12548. doi:10.1111/avsc.12548

See Also

backtransform, brms::brmsfit, brms::brm

Examples

```
novelforest_model <- download_model()</pre>
```

library(brms) # recommended
summary(novelforest_model)

to obtain input data
novelforest_model\$data

novelforestSG novelforestSG

Description

Forest community data used in Lai et al. (2021); also comprised part of the raw data used in Neo et al. (2017).

Details

The main component of novelforestSG is the dataset, which can be accessed with novelforest_data. See ?novelforest_data for more details.

Author(s)

Hao Ran Lai, <hrlai.ecology@gmail.com> Kwek Yan Chong, <kwek@u.nus.edu> Alex Thiam Koon Yee Maintainer: Hao Ran Lai, <hrlai.ecology@gmail.com>

References

Lai, H.R., Tan, G.S.Y., Neo, L., Kee, C.Y., Yee, A.T.K., Tan, H.T.W. and Chong, K.Y. (2021) Decoupled responses of native and exotic tree diversities to distance from old-growth forest and soil phosphorous in novel secondary forests. *Applied Vegetation Science*, 24, e12548. doi:10.1111/avsc.12548

Neo, L., Yee, A. T. K., Chong, K. Y., Kee, C. Y. and Tan, H. T. W. (2017). Vascular plant species richness and composition in two types of post-cultivation tropical secondary forest. *Applied Vege-tation Science*, 20, 692-701. doi:10.1111/avsc.12322

novelforest_data Novel Forest Raw Dataset

Description

The raw data used in Lai et al. (2021), also comprised part of the data in Neo et al. (2017).

Usage

novelforest_data

Format

A data.frame containing the following variables:

patch Name of forest patch

plot Name of forest plot within patch

UID Unique stem identifier

species Species name following Chong et al. (2011)

stem Indicator column denoting whether a stem is the main trunk (=1) of an individual tree, or otherwise (=0)

dbh_2011 Diameter-at-breast-height (cm) measured in year 2011.

See Lai et al. (2021) for more details on data collection.

References

Chong, K. Y., Tan, H. T. W. and Corlett, R. T. (2011). A summary of the total vascular plant flora of Singapore. *Gardens' Bulletin Singapore*, 63, 197-204.

Lai, H.R., Tan, G.S.Y., Neo, L., Kee, C.Y., Yee, A.T.K., Tan, H.T.W. and Chong, K.Y. (2021) Decoupled responses of native and exotic tree diversities to distance from old-growth forest and soil phosphorous in novel secondary forests. *Applied Vegetation Science*, 24, e12548. doi:10.1111/avsc.12548

Neo, L., Yee, A. T. K., Chong, K. Y., Kee, C. Y. and Tan, H. T. W. (2017). Vascular plant species richness and composition in two types of post-cultivation tropical secondary forest. *Applied Vegetation Science*, 20, 692-701. doi:10.1111/avsc.12322

Examples

```
# To access the dataset
head(novelforest_data)
```

Index

* datasets
 novelforest_data, 5

backtransform, 2

 $\texttt{download_model}, 3$

novelforest_data, 5
novelforestSG, 4