

Package ‘Plasticity’

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Type Package

Title A package for computing plasticity indices

Version 0.2

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Description The Plasticity package allows to compute several plasticity indices as defined in Valladares et al. (2006) at Journal of Ecology

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Imports agricolae,
dplyr,
ggplot2,
psych

Encoding UTF-8

LazyData true

RoxygenNote 6.1.1

StagedInstall no

R topics documented:

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rdpi	<i>RDPI</i>
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Description

Function to compute the RDPI (Relative Distance Plasticity Index, Valladares et al, (2006) Quantitative estimation of phenotypic plasticity: bridging the gap between the evolutionary concept and its ecological applications, Journal of Ecology, 94(6):1103-1116.

Usage

```
rdpi(dataframe, sp, trait, factor, verbose = T)
```

Arguments

dataframe	The dataframe that contains the data
sp	The bare (unquoted) name of the column whose values will be used as independent variable. The function will compare RDPI values among values of this variable. It can be species, provenances, etc.
trait	The bare (unquoted) name of the column that holds the trait for which to calculate RDPI. Must be numeric
factor	the bare (unquoted) name of the column that holds the environmental factor for which we will calculate RDPI. By definition, RDPI computes distances between pairs of observations that are at different levels of this factor.
verbose	defines if we want to get a data frame with all the individual RDPI values calculated. By default is set to 'TRUE'; indicating that we will get the data frame. If set to 'FALSE', we only get a summary table and a bocplot

Value

This function computes RDPI to the environmental factor for each species of the dataset(or any other identifying variable defined in 'sp') Then it makes an ANOVA or t-test of the values of RDPI across species and plots the boxplot

Examples

```
data(ecophysio)
rdpi(ecophysio,sp,SB, Piso, verbose = F)

# if we want to store the values

foo <- rdpi(ecophysio,sp,SB, Piso, verbose = T)
```

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