

# Package ‘Plasticity’

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**Type** Package

**Title** A package for computing plasticity indices

**Version** 1.0

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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

**License** `use\_mit\_license()`

**Imports** agricolae,  
dplyr,  
ggplot2,  
psych

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.3.2

**StagedInstall** no

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ecophysio	<i>Ecophysiological data of seedlings in the Pyrenees</i>
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## Description

A dataset containing several ecophysiological and morphological variables for 191 seedlings of 4 tree species planted in the Pyrenees. The variables are as follows:

## Usage

ecophysio

**Format**

A data frame with 191 rows and 14 variables

**Stage** elevation belt at which the plant is located (Montane, Subalpine)

**Species** tree species (*Abies alba*, *Betula pendula*, *Pinus sylvestris*, *Pinus uncinata*)

**GLI** light availability measured through Gap Light Index in percentage

**PB** Plant biomass in g

**LB** Leaf biomass in g

**SB** Stem biomass in g

**RB** Root biomass in g

**LWR** leaf weight ratio (leaf weight/total weight)

**SWR** stem weight ratio (stem weight/total weight)

**RWR** root weight ratio (root weight/total weight)

**RMF** root mass fraction (leaf weight/total weight)

**RSR** root:shoot ratio

**SLA** specific leaf area in cm<sup>2</sup>/g(leaf weight/total weight)

**WatPot** water potential in bar

**Source**

<<https://annforsci.biomedcentral.com/articles/10.1007/s13595-019-0831-1>>

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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)
```

**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.

**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)

# if we want to store the values

foo <- rdpi(ecophysio,Species,SB, Stage)
```

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rdpi\_matrix

*rdpi\_matrix*


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**Description**

Function to compute a vector containing the relative distance plasticity index values (Valladares et al. 2006) of a given trait (trait) for a given categorical environmental variable (factor). It calculates rdpi for each pair of observation that does not belong to the same level of "factor", and returns a vector containing all the calculated rdpi values.

Function to compute a matrix with the relative distance plasticity index values (Valladares et al. 2006) of a given trait (trait) for a given categorical environmental variable (factor). It calculates rdpi for each pair of observation that does not belong to the same level of "factor", and returns a vector containing all the calculated rdpi values.

**Usage**

```
rdpi_matrix(data, trait, factor)

rdpi_matrix(data, trait, factor)
```

**Arguments**

data	The dataframe that contains the data
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.

**Value**

a vector containing all the calculated rdpi values for each pair of observations that do not belong to the same level of "factor"

a vector containing all the calculated rdpi values for each pair of observations that do not belong to the same level of "factor"

**Examples**

```
data(ecophysio)
compute_rdpi(ecophysio, SB, Stage)
data(ecophysio)
rdpi_matrix(ecophysio, SB, Stage)
```

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