

# R documentation

of 'alpha\_div.Rd'

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alpha.div

*Calculate alpha-diversity*

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## Description

Calculates two types of  $\alpha$ -diversity: the Simpson index and the Shannon Weiner index.

## Usage

```
alpha.div(x, method = "simpson")
```

## Arguments

x	Either a dataframe of taxa abundances in sites, with sites comprising rows and columns comprising taxa (rownames of the dataframe should give site names), or a numeric vector comprising taxa abundances for a single site.
method	One of "simpson" or "shannon". Partial matches allowed.

## Value

The output is a list with the following components:

1. p.i A dataframe containing proportional abundances of taxa in sites.
2. method The diversity method applied.
3. rn The rownames of the original community dataset (i.e., the site names).
4. div A dataframe of actual computed diversities.

## Author(s)

Intro to R class

## References

Magurran, A. E. (1988). Ecological diversity and its measurement. Princeton university press.

**Examples**

```
taxa <- c(sp1 = 3, sp2 = 4, sp3 = 9)
x <- alpha.div(taxa)
plot(x)
plot(x, TRUE)
```

# Index

`alpha.div`, [1](#)

`plot.a_div(alpha.div)`, [1](#)

`print.a_div(alpha.div)`, [1](#)