

Package ‘diversityArch’

November 25, 2025

Type Package

Title Computes Diversity Indices with Archaeological Data

Version 0.1.0

Description Companion package of Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025), ``Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave".

License GPL-2

Imports methods

Encoding UTF-8

LazyData true

RoxygenNote 7.3.2

Suggests testthat (>= 3.0.0)

Config/testthat.edition 3

NeedsCompilation no

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Depends R (>= 3.5.0)

Repository CRAN

Date/Publication 2025-11-25 20:42:09 UTC

Contents

all_indices	2
dec_equit	3
dec_shannon	4

dominance	5
equitability	6
evenness	7
margalev	8
menhinick	8
Qesem_f	9
Qesem_s	10
shannon	11
shannon_frag	12
simpson_D	13
simpson_E	14

Index**15****all_indices***Diversity indices***Description**

Computes and prints all the diversity indices

Usage

```
all_indices(x, groups = NULL)
```

Arguments

- x Vector of dimension S (spicies) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA.
- groups Vector of dimension S of factors indicating the groups.

Value

No return value. It prints the value of all indicators

References

"Quantifying Diversity through Entropy Decomposition: Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[shannon](#), [dec_shannon](#), [dominance](#), [equitability](#), [evenness](#), [margalev](#), [menhinick](#), [simpson_D](#), [simpson_E](#)

Examples

```
data(Qesem_s)
all_indices(Qesem_s$HU)
all_indices(Qesem_s$HU, Qesem_s$Group)
```

dec_equit*Decomposition of the equitability index***Description**

Computes equitability and its decomposition

Usage

```
dec_equit(x, groups)
```

Arguments

- | | |
|---------------------|--|
| <code>x</code> | Vector of dimension S (species) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA. |
| <code>groups</code> | Vector of dimension S of factors indicating the groups. |

Value

- `equitability`: Equitability index.
- `within`: Within groups equitability.
- `between`: Between groups equitability.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[dec_shannon](#)

Examples

```
data(Qesem_s)
dec_equit(Qesem_s$HU, Qesem_s$Group)
```

dec_shannon

*Shannon diversity decomposition***Description**

Computes Shannon diversity and its decomposition

Usage

```
dec_shannon(x, groups)
```

Arguments

- | | |
|--------|--|
| x | Vector of dimension S (species) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA. |
| groups | Vector of dimension S of factors indicating the groups. |

Value

- shannon: Shannon's total Entropy.
- within: Within groups entropy.
- between: Between groups entropy.
- groups: A data frame with information about each group: relative frequency, internal entropy and number of species.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[shannon](#)

Examples

```
data(Qesem_s)
dec_shannon(Qesem_s$HU, Qesem_s$Group)
```

dominance	<i>Dominance index</i>
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Description

Modification of the Simpson's dominance index to be restricted between 0 and 1.

Usage

```
dominance(x)
```

Arguments

- x Vector of dimension S (species) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA.

Value

Dominance index (Modified Simpson's dominance index).

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[simpson_D](#)

Examples

```
data(Qesem_s)
dominance(Qesem_s$HU)
```

equitability	<i>Equitability. J Pielou index</i>
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Description

Computes J Pielou index know as equitability

Usage

```
equitability(x)
```

Arguments

- x Vector of dimension S (spicies) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA.

Value

Equitability. J Pielou index.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[shannon](#)

Examples

```
data(Qesem_s)
equitability(Qesem_s$HU)
```

evenness	<i>Evenness index</i>
----------	-----------------------

Description

Modification of Simpson's evenness index to be restricted between 0 and 1

Usage

```
evenness(x)
```

Arguments

- x Vector of dimension S (species) with the numbers of observed individuals in each species. NA values are allowed. 0 values are converted to NA.

Value

Evenness index.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[simpson_E](#)

Examples

```
data(Qesem_s)
evenness(Qesem_s$HU)
```

margalev*Margalev index***Description**

Computes Margalev's index

Usage

```
margalev(x)
```

Arguments

- x Vector of dimension S (species) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA.

Value

Margalev index.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

Examples

```
data(Qesem_s)
margalev(Qesem_s$HU)
```

menhinick*Menhirnick index***Description**

Computes menhinick's index

Usage

```
menhinick(x)
```

Arguments

- x Vector of dimension S (species) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA.

Value

Mehhinick index.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

Examples

```
data(Qesem_s)
menhinick(Qesem_s$HU)
```

Qesem_f

Data used in the examples

Description

Data frame with number of fragments of bones of unidentified species but identified group for several levels in Qesem (Israel). Data source: Blasco, R., Rosell, J., Assaf, E., Barkai, R., Gopher, A., (2024)

Usage

```
data(Qesem_f)
```

Value

Data frame with 4 observations (groups) and 7 levels.

Author(s)

Blasco, R., Rosell, J., Assaf, E., Barkai, R., Gopher, A.

Source

Blasco, R., Rosell, J., Assaf, E., Barkai, R., Gopher, A., 2024. Exploring the lack of articular ends at the Middle Pleistocene site of Qesem Cave, Israel. Journal of Human Evolution 189, 103509. doi:10.1016/j.jhevol.2024.103509

References

Barat, A. Sansó, A. Arilla-Osuna, M. Blasco, R., Pérez-Fernández, I., Cifuentes-Alcobenda, G. Llorente, R., Vivar-Ríos, D., Assaf, E. Barkai, R., Gopher, A. & Rosell-Ardèvol, J., 2025. Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave. Blasco, R., Rosell, J., Assaf, E., Barkai, R., Gopher, A., 2024. Exploring the lack of articular ends at the Middle Pleistocene site of Qesem Cave, Israel. Journal of Human Evolution 189, 103509. doi:10.1016/j.jhevol.2024.103509

Examples

```
data(Qesem_f)
names(data)
# The following example replicates some of the results of Table 5 in
# Barat, A. Sansó, A. Arilla-Osuna, M. Blasco, R., Pérez-Fernández, I.,
# Cifuentes-Alcobenda, G. Llorente, R., Vivar-Ríos, D., Assaf, E. Barkai, R.,
# Gopher, A. & Rosell-Ardèvol, J., 2025. "Quantifying Diversity through Entropy
# Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave".
shannon_frag(Qesem_s$HU, Qesem_s$Group, Qesem_f$HU, Qesem_f$Group)
```

Qesem_s

Data used in the examples

Description

Data frame with number of bones of different species and different levels in Qesem (Israel), and a factor related to the size of the animal. Data source: Blasco, R., Rosell, J., Assaf, E., Barkai, R., Gopher, A., (2024)

Usage

```
data(Qesem_s)
```

Value

Data frame with 15 observations and 7 levels.

Author(s)

Blasco, R., Rosell, J., Assaf, E., Barkai, R., Gopher, A.

Source

Blasco, R., Rosell, J., Assaf, E., Barkai, R., Gopher, A., 2024. Exploring the lack of articular ends at the Middle Pleistocene site of Qesem Cave, Israel. Journal of Human Evolution 189, 103509. doi:10.1016/j.jhevol.2024.103509

References

Barat, A., Sansó, A., Arilla-Osuna, M., Blasco, R., Pérez-Fernández, I., Cifuentes-Alcobenda, G., Llorente, R., Vivar-Ríos, D., Assaf, E., Barkai, R., Gopher, A. & Rosell-Ardèvol, J., 2025. Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave. Blasco, R., Rosell, J., Assaf, E., Barkai, R., Gopher, A., 2024. Exploring the lack of articular ends at the Middle Pleistocene site of Qesem Cave, Israel. Journal of Human Evolution 189, 103509. doi:10.1016/j.jhevol.2024.103509

Examples

```
data(Qesem_s)
names(Qesem_s)
# The following example replicates some of the results of Table 3 in
# Barat, A. Sansó, A. Arilla-Osuna, M. Blasco, R., Pérez-Fernández, I.,
# Cifuentes-Alcobenda, G. Llorente, R., Vivar-Ríos, D., Assaf, E. Barkai, R.,
# Gopher, A. & Rosell-Ardèvol, J., 2025. "Quantifying Diversity through Entropy
# Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave".
all_indices(Qesem_s$HU)
```

shannon

Shannon diversity index

Description

Computes Shannon's diversity index

Usage

```
shannon(x)
```

Arguments

- x Vector of dimension S (species) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA.

Value

Shannon's diversity index (Entropy).

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[dec_shannon](#), [equitability](#)

Examples

```
data(Qesem_s)
shannon(Qesem_s$HU)
```

shannon_frag

Shannon diversity decomposition

Description

Computes Shannon diversity and its decomposition

Usage

```
shannon_frag(x, gx, f, gf)
```

Arguments

- | | |
|-----------------|--|
| <code>x</code> | Vector of dimension S (species) with the numbers of observed individuals in each species. NA values are allowed. 0 values are converted to NA. |
| <code>gx</code> | Vector of dimension S of factors indicating the groups. G groups. |
| <code>f</code> | Vector of dimension G with the number (>0) of fragments in each group |
| <code>gf</code> | Vector of dimension G of factors indicating the groups of f. |

Value

- `shannon`: Shannon's total Entropy.
- `within`: Within groups entropy.
- `between`: Between groups entropy.
- `groups`: A data frame with information about each group: relative frequency, internal entropy and number of species.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[shannon](#)

Examples

```
data(Qesem_s)
data(Qesem_f)
shannon_frag(Qesem_s$HU, Qesem_s$Group, Qesem_f$HU, Qesem_f$Group)
```

simpson_D

Simpson's dominance index

Description

Computes Simpson's dominance index.

Usage

```
simpson_D(x)
```

Arguments

- x Vector of dimension S (spicies) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA.

Value

Simpson's dominance index.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[dominance](#), [simpson_E](#)

Examples

```
data(Qesem_s)
simpson_D(Qesem_s$HU)
```

simpson_E*Simpson's evenness index***Description**

Computes Simpson's evenness index.

Usage

```
simpson_E(x)
```

Arguments

- x Vector of dimension S (species) with the numbers of observed individuals in each spicy. NA values are allowed. 0 values are converted to NA.

Value

Simpson's evenness index.

References

"Arnaud Barat, Andreu Sansó, Maite Arilla-Osuna, Ruth Blasco, Iñaki Pérez-Fernández, Gabriel Cifuentes-Alcobenda, Rubén Llorente, Daniel Vivar-Ríos, Ella Assaf, Ran Barkai, Avi Gopher, & Jordi Rosell-Ardèvol (2025): Quantifying Diversity through Entropy Decomposition. Insights into Hominin Occupation and Carcass Processing at Qesem cave"

See Also

[evenness](#), [simpson_D](#)

Examples

```
data(Qesem_s)
simpson_E(Qesem_s$HU)
```

Index

* Datasets

Qesem_f, [9](#)

Qesem_s, [10](#)

all_indices, [2](#)

dec_equit, [3](#)

dec_shannon, [2, 3, 4, 11](#)

dominance, [2, 5, 13](#)

equitability, [2, 6, 11](#)

evenness, [2, 7, 14](#)

margalev, [2, 8](#)

menhinick, [2, 8](#)

Qesem_f, [9](#)

Qesem_s, [10](#)

shannon, [2, 4, 6, 11, 12](#)

shannon_frag, [12](#)

simpson_D, [2, 5, 13, 14](#)

simpson_E, [2, 7, 13, 14](#)