

# Package ‘soundecology’

October 24, 2013

**Title** Soundscape ecology

**Version** 1.0

**Date** 2013-10-24

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**Description** Functions to calculate indices for soundscape ecology and other ecology research that uses audio recordings.

**Depends** R(>= 2.14.0), tuneR, ineq, vegan, parallel, seewave, pracma, oce

**Suggests** knitr

**License** GPL-3

**URL** <http://ljvillanueva.github.io/soundecology/>

**BugReports** <http://github.com/ljvillanueva/soundecology/issues>

**VignetteBuilder** knitr

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acoustic\_complexity     *Acoustic Complexity Index*


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### Description

Acoustic Complexity Index (ACI) from Pieretti, *et al.* 2011. The ACI is based on the "observation that many biotic sounds, such as bird songs, are characterized by an intrinsic variability of intensities, while some types of human generated noise (such as car passing or airplane transit) present very constant intensity values" (Pieretti, *et al.* 2011).

The index was tested to the ACItot calculated using SoundscapeMeter v 1.0.14.05.2012, courtesy of A. Farina.

### Usage

```
acoustic_complexity(soundfile, max_freq=NA, j=5, fft_w=512)
```

### Arguments

soundfile	an object of class Wave loaded with the function readWave of the tuneR package.
max_freq	maximum frequency to use when calculating the value, in Hertz. The default is the maximum for the file.
j	the cluster size, in seconds.
fft_w	FFT window to use.

### Value

Returns a list with three objects per channel

AciTotAll\_left the ACI total for the left channel

AciTotAll\_right  
the ACI total for the right channel

aci\_fl\_left\_vals  
values of ACI(fl) for the left channel

aci\_fl\_right\_vals  
values of ACI(fl) for the right channel

aci\_left\_matrix  
Matrix of all values before calculating ACI(fl) for the left channel

aci\_right\_matrix  
Matrix of all values before calculating ACI(fl) for the right channel

### References

Pieretti, N., A. Farina, and D. Morri. 2011. A new methodology to infer the singing activity of an avian community: The Acoustic Complexity Index (ACI). *Ecological Indicators* 11: 868-873. doi: 10.1016/j.ecolind.2010.11.005

**Examples**

```
data(tropicalsound)
ACI <- acoustic_complexity(tropicalsound)
print(ACI$AciTotAll_left)

summary(ACI)
```

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acoustic_diversity	<i>Acoustic Diversity Index</i>
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**Description**

Acoustic Diversity Index from Villanueva-Rivera *et al.* 2011. The ADI is calculated by dividing the spectrogram into bins (default 10) and taking the proportion of the signals in each bin above a threshold (default -50 dBFS). The ADI is the result of the Shannon index applied to these bins.

**Usage**

```
acoustic_diversity(soundfile, max_freq = 10000, db_threshold = -50,
  freq_step = 1000, shannon = TRUE)
```

**Arguments**

soundfile	an object of class "Wave" loaded with the function readWave of the tuneR package.
max_freq	maximum frequency to use when calculating the value, in Hertz.
db_threshold	threshold to use in the calculation.
freq_step	size of frequency bands.
shannon	TRUE to use the Shannon's diversity index to calculate the ADI (default).

**Value**

Returns a list with five objects per channel

adi_left	ADI value for the left channel
adi_right	ADI value for the right channel
left_band_values	vector of occupancy values for each band for the left channel
right_band_values	vector of occupancy values for each band for the right channel
left_bandrange_values	vector of frequency values for each band for the left channel
right_bandrange_values	vector of frequency values for each band for the right channel

## Note

The code to calculate the ADI has changed due to an error we detected in the original scripts in which the value was calculated using a different equation. In a test of ~38k files, both ways to calculate were highly correlated. This version of the function uses the Shannon's Diversity Index. To obtain a result using the old calculation, set the argument `shannon` to `FALSE`. Please check the vignette "Changes in the Acoustic Diversity Index", included in the package, for more details.

For audio files with one channel, the results are showed as the left channel, the right channel returns "NA".

## References

Villanueva-Rivera, L. J., B. C. Pijanowski, J. Doucette, and B. Pekin. 2011. A primer of acoustic analysis for landscape ecologists. *Landscape Ecology* 26: 1233-1246. doi: 10.1007/s10980-011-9636-9.

## Examples

```
data(tropicalsound)
result <- acoustic_diversity(tropicalsound)

print(result$adi_left)

summary(result)
```

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acoustic_eveness	<i>Acoustic Eveness Index</i>
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## Description

Acoustic Eveness Index from Villanueva-Rivera *et al.* 2011 (band eveness using the Gini index). The AEI is calculated by dividing the spectrogram into bins (default 10) and taking the proportion of the signals in each bin above a threshold (default -50 dBFS). The AEI is the result of the Gini index applied to these bins.

## Usage

```
acoustic_eveness(soundfile, max_freq=10000, db_threshold=-50, freq_step=1000)
```

## Arguments

<code>soundfile</code>	an object of class "Wave" loaded with the function <code>readWave</code> of the <code>tuneR</code> package.
<code>max_freq</code>	maximum frequency to use when calculating the value, in Hertz.
<code>db_threshold</code>	threshold to use in the calculation.
<code>freq_step</code>	size of frequency bands.

**Value**

Returns a list with five objects per channel

aei\_left            AEI for the left channel  
aei\_right          AEI for the right channel

**Note**

For audio files with one channel, the results are showed as the left channel, the right channel returns "NA".

**References**

Villanueva-Rivera, L. J., B. C. Pijanowski, J. Doucette, and B. Pekin. 2011. A primer of acoustic analysis for landscape ecologists. *Landscape Ecology* 26: 1233-1246. doi: 10.1007/s10980-011-9636-9.

**Examples**

```
data(tropicalsound)
result <- acoustic_eveness(tropicalsound)
print(result$aei_left)

summary(result)
```

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bioacoustic_index	<i>Bioacoustic Index</i>
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**Description**

Bioacoustic Index from Boelman, *et al.* 2007. Inspired on Matlab code courtesy of NT Boelman. Several parts where changed, in particular log math, so this won't be directly comparable to the original code in the paper.

The Bioacoustic Index is calculated as the "area under each curve included all frequency bands associated with the dB value that was greater than the minimum dB value for each curve. The area values are thus a function of both the sound level and the number of frequency bands used by the avifauna" (Boelman, *et al.* 2007).

**Usage**

```
bioacoustic_index(soundfile, min_freq=2000, max_freq=8000, fft_w=512)
```

**Arguments**

soundfile	an object of class "Wave" loaded with the function readWave of the tuneR package.
min_freq	minimum frequency to use when calculating the value, in Hertz.
max_freq	maximum frequency to use when calculating the value, in Hertz.
fft_w	FFT window size.

**Value**

Returns a list with one object per channel

left_area	area under the curve for the left channel
right_area	area under the curve for the right channel

**References**

Boelman NT, Asner GP, Hart PJ, Martin RE. 2007. Multi-trophic invasion resistance in Hawaii: bioacoustics, field surveys, and airborne remote sensing. *Ecological Applications* 17:2137-2144.

**Examples**

```
data(tropicalsound)
bioindex <- bioacoustic_index(tropicalsound)
print(bioindex$left_area)

summary(bioindex)
```

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multiple_sounds	<i>Multiple sound files</i>
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**Description**

Function to extract the specified index from all the wav files in a directory. The results, including the filename and wave technical details, are saved to a csv file. If the computer has multiple cores, it can run files in parallel.

**Usage**

```
multiple_sounds(directory, resultfile, soundindex, no_cores=1, ...)
```

**Arguments**

directory	a valid directory, readable by the user, that contains the wav files.
resultfile	name of the text file to which write the results in comma-separated values format.
soundindex	which index to calculate: <ul style="list-style-type: none"> <li>• ndsi</li> <li>• acoustic_complexity</li> <li>• acoustic_diversity</li> <li>• acoustic_evenness</li> <li>• bioacoustic_index</li> <li>• H from the seewave package</li> </ul>
no_cores	number of cores to use when calculating the indices. Can be "max" to use all cores, "-1" to use all but one core, or any positive integer. Default is 1. Uses the parallel package.
...	additional variables to pass to the selected function. See each function's help for details.

**Examples**

```
## Not run:
#Calculate the ACI of all the wav
# files in the directory "/home/user/wavs/"
# using the function acoustic_complexity
multiple_sounds(directory="/home/user/wavs/",
resultfile="/home/user/results.csv",
soundindex="acoustic_complexity")

#Calculate the same as above using 12000Hz as the
# maximum frequency instead of the default.
multiple_sounds(directory="/home/user/wavs/",
resultfile="/home/user/results.csv",
soundindex="acoustic_complexity", max_freq=12000)

#Calculate the same as above using two cores
multiple_sounds(directory="/home/user/wavs/",
resultfile="/home/user/results.csv",
soundindex="acoustic_complexity", no_cores=2)

#Calculate the same as above using all the cores
# the computer has
multiple_sounds(directory="/home/user/wavs/",
resultfile="/home/user/results.csv",
soundindex="acoustic_complexity", no_cores="max")

#Calculate the same as above using all but one cores
multiple_sounds(directory="/home/user/wavs/",
resultfile="/home/user/results.csv",
soundindex="acoustic_complexity", no_cores=-1)
```

```
## End(Not run)
```

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ndsi

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*Normalized Difference Soundscape Index*


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## Description

Normalized Difference Soundscape Index (NDSI) from REAL and Kasten, *et al.* 2012. The NDSI seeks to "estimate the level of anthropogenic disturbance on the soundscape by computing the ratio of human-generated (anthrophony) to biological (biophony) acoustic components found in field collected sound samples" (Kasten, *et al.* 2012).

Tested with Matlab code courtesy of S. Gage.

## Usage

```
ndsi(soundfile, fft_w=1024, anthro_min=1000, anthro_max=2000,
      bio_min=2000, bio_max=11000)
```

## Arguments

soundfile	an object of class "Wave" loaded with the function readWave of the tuneR package.
fft_w	FFT window size.
anthro_min	minimum value of the range of frequencies of the anthrophony.
anthro_max	maximum value of the range of frequencies of the anthrophony.
bio_min	minimum value of the range of frequencies of the biophony.
bio_max	maximum value of the range of frequencies of the biophony.

## Details

The bin size is determined as the difference between anthro\_max and anthro\_min, by default 1000Hz.

## Value

Returns a list with one object per channel

ndsi_left	NDSI value for the left channel
ndsi_right	NDSI value for the left channel

## References

Remote Environmental Assessment Laboratory. <http://www.real.msu.edu>

Kasten, Eric P., Stuart H. Gage, Jordan Fox, and Wooyeong Joo. 2012. The Remote Environmental Assessment Laboratory's Acoustic Library: An Archive for Studying Soundscape Ecology. *Ecological Informatics* 12: 50-67. doi: 10.1016/j.ecoinf.2012.08.001



**Examples**

```
data(tropicalsound)
NDSI <- ndsi(tropicalsound)
print(NDSI$ndsi_left)

summary(NDSI)
```

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soundecology

*Soundscape ecology*

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**Description**

Functions to calculate indices for soundscape ecology and other ecology research that uses audio recordings.

**Details**

Package: soundecology  
Type: Package  
Version: 1.0  
Date: 2013-10-24  
License: GPLv3

**Author(s)**

Luis J. Villanueva-Rivera and Bryan C. Pijanowski

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tropicalsound

*tropicalsound sound example*

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**Description**

Sample sound of a digital recording of a chorus of tropical frogs.

**Usage**

```
data(tropicalsound)
```

**Format**

A "Wave" object.

**Details**

Duration = 20 sec. Sampling rate = 22050 Hz.

**Source**

Recording made at a tropical rainforest in Puerto Rico by Luis J. Villanueva-Rivera.

**Examples**

```
data(tropicalsound)
```

```
tropicalsound
```

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