

Package ‘soundgen’

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Maintainer Andrey Anikin <rty_anik@yahoo.com>

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Description Performs parametric synthesis of sounds with harmonic and noise components such as animal vocalizations or human voice. Also offers tools for audio manipulation and acoustic analysis, including pitch tracking, spectral analysis, audio segmentation, pitch and formant shifting, etc. Includes four interactive web apps for synthesizing and annotating audio, manually correcting pitch contours, and measuring formant frequencies. Reference: Anikin (2019) <[doi:10.3758/s13428-018-1095-7](https://doi.org/10.3758/s13428-018-1095-7)>.

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Author Andrey Anikin [aut, cre]

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Description

Adds sinusoidal or logistic amplitude modulation to a sound. This produces additional harmonics in the spectrum at $\pm am_freq$ around each original harmonic and makes the sound rough. The optimal frequency for creating a perception of roughness is ~ 70 Hz (Fastl & Zwicker "Psychoacoustics"). Sinusoidal AM creates a single pair of new harmonics, while non-sinusoidal AM creates more extra harmonics (see examples).

Usage

```
addAM(
  x,
  samplingRate = NULL,
  amDep = 25,
  amFreq = 30,
  amType = c("logistic", "sine")[1],
  amShape = 0,
  invalidArgAction = c("adjust", "abort", "ignore")[1],
  plot = FALSE,
  play = FALSE,
  saveAudio = NULL,
  reportEvery = NULL,
  cores = 1
)
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>amDep</code>	amplitude modulation (AM) depth, %. 0: no change; 100: AM with amplitude range equal to the dynamic range of the sound (anchor format)
<code>amFreq</code>	AM frequency, Hz (anchor format)
<code>amType</code>	"sine" = sinusoidal, "logistic" = logistic (default)
<code>amShape</code>	ignore if <code>amType</code> = "sine", otherwise determines the shape of non-sinusoidal AM: 0 = ~sine, -1 = notches, +1 = clicks (anchor format)
<code>invalidArgAction</code>	what to do if an argument is invalid or outside the range in <code>permittedValues</code> : 'adjust' = reset to default value, 'abort' = stop execution, 'ignore' = throw a warning and continue (may crash)
<code>plot</code>	if TRUE, plots the amplitude modulation
<code>play</code>	if TRUE, plays the processed audio
<code>saveAudio</code>	full (!) path to folder for saving the processed audio; NULL = don't save, "" = same as input folder (NB: overwrites the originals!)
<code>reportEvery</code>	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
<code>cores</code>	number of cores for parallel processing

Examples

```

sound1 = soundgen(pitch = c(200, 300), addSilence = 0)
s1 = addAM(sound1, 16000, amDep = c(0, 50, 0), amFreq = 75, plot = TRUE)
# playme(s1)
## Not run:
# Parameters can be specified as in the soundgen() function, eg:
s2 = addAM(sound1, 16000,
           amDep = list(time = c(0, 50, 52, 200, 201, 300),
                       value = c(0, 0, 35, 25, 0, 0)),
           plot = TRUE, play = TRUE)

# Sinusoidal AM produces exactly 2 extra harmonics at  $\pm$ am_freq
# around each f0 harmonic:
s3 = addAM(sound1, 16000, amDep = 30, amFreq = c(50, 80),
           amType = 'sine', plot = TRUE, play = TRUE)
spectrogram(s3, 16000, windowLength = 150, ylim = c(0, 2))

# Non-sinusoidal AM produces multiple new harmonics,
# which can resemble subharmonics...
s4 = addAM(sound1, 16000, amDep = 70, amFreq = 50, amShape = -1,
           plot = TRUE, play = TRUE)
spectrogram(s4, 16000, windowLength = 150, ylim = c(0, 2))

# ...but more often look like sidebands
sound3 = soundgen(syllLen = 600, pitch = c(800, 1300, 1100), addSilence = 0)
s5 = addAM(sound3, 16000, amDep = c(0, 30, 100, 40, 0),
           amFreq = 105, amShape = -.3,
           plot = TRUE, play = TRUE)
spectrogram(s5, 16000, ylim = c(0, 5))

# Feel free to add AM stochastically:
s6 = addAM(sound1, 16000,
           amDep = rnorm(10, 40, 20), amFreq = rnorm(20, 70, 20),
           plot = TRUE, play = TRUE)
spectrogram(s6, 16000, windowLength = 150, ylim = c(0, 2))

# If am_freq is locked to an integer ratio of f0, we can get subharmonics
# For ex., here is with pitch 400-600-400 Hz (soundgen interpolates pitch
# on a log scale and am_freq on a linear scale, so we align them by extracting
# a long contour on a log scale for both)
con = getSmoothContour(anchors = c(400, 600, 400),
                      len = 20, thisIsPitch = TRUE)
s = soundgen(syllLen = 1500, pitch = con, amFreq = con/3, amDep = 30,
           plot = TRUE, play = TRUE, ylim = c(0, 3))

# Process all files in a folder and save the modified audio
addAM('~Downloads/temp', saveAudio = '~/Downloads/temp/AM',
      amFreq = 70, amDep = c(0, 50))

## End(Not run)

```

 addFormants

Add formants

Description

A spectral filter that either adds or removes formants from a sound - that is, amplifies or dampens certain frequency bands, as in human vowels. See [soundgen](#) and [getSpectralEnvelope](#) for more information. With `action = 'remove'` this function can perform inverse filtering to remove formants and obtain raw glottal output, provided that you can specify the correct formant structure. Instead of formants, any arbitrary spectral filtering function can be applied using the `spectralEnvelope` argument (eg for a low/high/bandpass filter).

Usage

```
addFormants(
  x,
  samplingRate = NULL,
  formants = NULL,
  spectralEnvelope = NULL,
  zFun = NULL,
  action = c("add", "remove")[1],
  vocalTract = NA,
  formantDep = 1,
  formantDepStoch = 1,
  formantWidth = 1,
  formantCeiling = 2,
  lipRad = 6,
  noseRad = 4,
  mouthOpenThres = 0,
  mouth = NA,
  temperature = 0.025,
  formDrift = 0.3,
  formDisp = 0.2,
  smoothing = list(),
  windowLength_points = 800,
  overlap = 75,
  normalize = c("max", "orig", "none")[1],
  play = FALSE,
  saveAudio = NULL,
  reportEvery = NULL,
  cores = 1,
  ...
)
```

Arguments

`x` path to a folder, one or more wav or mp3 files `c('file1.wav', 'file2.mp3')`, Wave object, numeric vector, or a list of Wave objects or numeric vectors

samplingRate	sampling frequency, Hz
formants	either a character string like "aau" referring to default presets for speaker "M1" or a list of formant times, frequencies, amplitudes, and bandwidths (see ex. below). formants = NA defaults to schwa. Time stamps for formants and mouthOpening can be specified in ms or an any other arbitrary scale. See getSpectralEnvelope for more details
spectralEnvelope	(optional): as an alternative to specifying formant frequencies, we can provide the exact filter - a vector of non-negative numbers specifying the power in each frequency bin on a linear scale (interpolated to length equal to windowLength_points/2). A matrix specifying the filter for each STFT step is also accepted. The easiest way to create this matrix is to call <code>soundgen::getSpectralEnvelope</code> or to use the spectrum of a recorded sound
zFun	(optional) an arbitrary function to apply to the spectrogram prior to iSTFT, where "z" is the spectrogram - a matrix of complex values (see examples)
action	'add' = add formants to the sound, 'remove' = remove formants (inverse filtering)
vocalTract	the length of vocal tract, cm. Used for calculating formant dispersion (for adding extra formants) and formant transitions as the mouth opens and closes. If NULL or NA, the length is estimated based on specified formant frequencies, if any (anchor format)
formantDep	scale factor of formant amplitude (1 = no change relative to amplitudes in formants)
formantDepStoch	the amplitude of additional stochastic formants added above the highest specified formant, dB (only if temperature > 0)
formantWidth	scale factor of formant bandwidth (1 = no change)
formantCeiling	frequency to which stochastic formants are calculated, in multiples of the Nyquist frequency; increase up to ~10 for long vocal tracts to avoid losing energy in the upper part of the spectrum
lipRad	the effect of lip radiation on source spectrum, dB/oct (the default of +6 dB/oct produces a high-frequency boost when the mouth is open)
noseRad	the effect of radiation through the nose on source spectrum, dB/oct (the alternative to lipRad when the mouth is closed)
mouthOpenThres	open the lips (switch from nose radiation to lip radiation) when the mouth is open >mouthOpenThres, 0 to 1
mouth	mouth opening (0 to 1, 0.5 = neutral, i.e. no modification) (anchor format)
temperature	hyperparameter for regulating the amount of stochasticity in sound generation
formDrift, formDisp	scaling factors for the effect of temperature on formant drift and dispersal, respectively
smoothing	a list of parameters passed to getSmoothContour to control the interpolation and smoothing of contours: <code>interpol</code> (approx / spline / loess), <code>loessSpan</code> , <code>discontThres</code> , <code>jumpThres</code>

windowLength_points	length of FFT window, points
overlap	FFT window overlap, %. For allowed values, see istft
normalize	"orig" = same as input (default), "max" = maximum possible peak amplitude, "none" = no normalization
play	if TRUE, plays the synthesized sound using the default player on your system. If character, passed to play as the name of player to use, eg "aplay", "play", "vlc", etc. In case of errors, try setting another default player for play
saveAudio	path + filename for saving the output, e.g. '~/Downloads/temp.wav'. If NULL = doesn't save
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
...	other plotting parameters passed to spectrogram

Details

Algorithm: converts input from a time series (time domain) to a spectrogram (frequency domain) through short-time Fourier transform (STFT), multiplies by the spectral filter containing the specified formants, and transforms back to a time series via inverse STFT. This is a subroutine for voice synthesis in [soundgen](#), but it can also be applied to a recording.

See Also

[getSpectralEnvelope](#) [transplantFormants](#) [soundgen](#)

Examples

```

sound = c(rep(0, 1000), runif(8000) * 2 - 1, rep(0, 1000)) # white noise
# NB: pad with silence to avoid artefacts if removing formants
# playme(sound)
# spectrogram(sound, samplingRate = 16000)

# add F1 = 900, F2 = 1300 Hz
sound_filtered = addFormants(sound, samplingRate = 16000,
                             formants = c(900, 1300))
# playme(sound_filtered)
# spectrogram(sound_filtered, samplingRate = 16000)

# ...and remove them again (assuming we know what the formants are)
sound_inverse_filt = addFormants(sound_filtered,
                                 samplingRate = 16000,
                                 formants = c(900, 1300),
                                 action = 'remove')
# playme(sound_inverse_filt)
# spectrogram(sound_inverse_filt, samplingRate = 16000)

## Not run:
## Perform some user-defined manipulation of the spectrogram with zFun

```



```

# Ex.: noise removal - silence all bins under threshold,
# say -0 dB below the max value
s_noisy = soundgen(syllLen = 200, addSilence = 0,
                  noise = list(time = c(-100, 300), value = -20))
spectrogram(s_noisy, 16000)
# playme(s_noisy)
zFun = function(z, cutoff = -50) {
  az = abs(z)
  thres = max(az) * 10 ^ (cutoff / 20)
  z[which(az < thres)] = 0
  return(z)
}
s_denoised = addFormants(s_noisy, samplingRate = 16000,
                        formants = NA, zFun = zFun, cutoff = -40)
spectrogram(s_denoised, 16000)
# playme(s_denoised)

# If neither formants nor spectralEnvelope are defined, only lipRad has an effect
# For ex., we can boost low frequencies by 6 dB/oct
noise = runif(8000)
noise1 = addFormants(noise, 16000, lipRad = -6)
seewave::meanspec(noise1, f = 16000, dB = 'max0')

# Arbitrary spectra can be defined with spectralEnvelope. For ex., we can
# have a flat spectrum up to 2 kHz (Nyquist / 4) and -3 dB/kHz above:
freqs = seq(0, 16000 / 2, length.out = 100)
n = length(freqs)
idx = (n / 4):n
sp_dB = c(rep(0, n / 4 - 1), (freqs[idx] - freqs[idx[1]]) / 1000 * (-3))
plot(freqs, sp_dB, type = 'b')
noise2 = addFormants(noise, 16000, lipRad = 0, spectralEnvelope = 10 ^ (sp_dB / 20))
seewave::meanspec(noise2, f = 16000, dB = 'max0')

## Use the spectral envelope of an existing recording (bleating of a sheep)
# (see also the same example with noise as source in ?generateNoise)
# (NB: this can also be achieved with a single call to transplantFormants)
data(sheep, package = 'seewave') # import a recording from seewave
sound_orig = as.numeric(scale(sheep@left))
samplingRate = sheep@samp.rate
sound_orig = sound_orig / max(abs(sound_orig)) # range -1 to +1
# playme(sound_orig, samplingRate)

# get a few pitch anchors to reproduce the original intonation
pitch = analyze(sound_orig, samplingRate = samplingRate,
               pitchMethod = c('autocor', 'dom'))$detailed$pitch
pitch = pitch[!is.na(pitch)]

# extract a frequency-smoothed version of the original spectrogram
# to use as filter
specEnv_bleating = spectrogram(sound_orig, windowLength = 5,
                              samplingRate = samplingRate, output = 'original', plot = FALSE)
# image(t(log(specEnv_bleating)))

```

```

# Synthesize source only, with flat spectrum
sound_unfilt = soundgen(syllLen = 2500, pitch = pitch,
  rolloff = 0, rolloffOct = 0, rolloffKHz = 0,
  temperature = 0, jitterDep = 0, subDep = 0,
  formants = NULL, lipRad = 0, samplingRate = samplingRate,
  invalidArgAction = 'ignore') # prevent soundgen from increasing samplingRate
# playme(sound_unfilt, samplingRate)
# seewave::meanspec(sound_unfilt, f = samplingRate, dB = 'max0') # ~flat

# Force spectral envelope to the shape of target
sound_filt = addFormants(sound_unfilt, formants = NULL,
  spectralEnvelope = specEnv_bleating, samplingRate = samplingRate)
# playme(sound_filt, samplingRate) # playme(sound_orig, samplingRate)
# spectrogram(sound_filt, samplingRate) # spectrogram(sound_orig, samplingRate)

# The spectral envelope is now similar to the original recording. Compare:
par(mfrow = c(1, 2))
seewave::meanspec(sound_orig, f = samplingRate, dB = 'max0', alim = c(-50, 20))
seewave::meanspec(sound_filt, f = samplingRate, dB = 'max0', alim = c(-50, 20))
par(mfrow = c(1, 1))
# NB: but the source of excitation in the original is actually a mix of
# harmonics and noise, while the new sound is purely tonal

## End(Not run)

```

addVectors

Add overlapping vectors

Description

Adds two partly overlapping vectors, such as two waveforms, to produce a longer vector. The location at which vector 2 is pasted is defined by `insertionPoint`. Algorithm: both vectors are padded with zeros to match in length and then added. All NA's are converted to 0.

Usage

```
addVectors(v1, v2, insertionPoint = 1, normalize = TRUE)
```

Arguments

<code>v1, v2</code>	numeric vectors
<code>insertionPoint</code>	the index of element in vector 1 at which vector 2 will be inserted (any integer, can also be negative)
<code>normalize</code>	if TRUE, the output is normalized to range from -1 to +1

See Also

[soundgen](#)

Examples

```

v1 = 1:6
v2 = rep(100, 3)
addVectors(v1, v2, insertionPoint = 5, normalize = FALSE)
addVectors(v1, v2, insertionPoint = -4, normalize = FALSE)
# note the asymmetry: insertionPoint refers to the first arg
addVectors(v2, v1, insertionPoint = -4, normalize = FALSE)

v3 = rep(100, 15)
addVectors(v1, v3, insertionPoint = -4, normalize = FALSE)
addVectors(v2, v3, insertionPoint = 7, normalize = FALSE)

```

analyze

Acoustic analysis

Description

Acoustic analysis of one or more sounds: pitch tracking, basic spectral characteristics, formants, estimated loudness (see [getLoudness](#)), roughness (see [modulationSpectrum](#)), novelty (see [ssm](#)), etc. The default values of arguments are optimized for human non-linguistic vocalizations. See vignette('acoustic_analysis', package = 'soundgen') for details. The defaults and reasonable ranges of all arguments can be found in [defaults_analyze](#). For high-precision work, first extract and manually correct pitch contours with [pitch_app](#), PRAAT, or whatever, and then run `analyze(pitchManual = ...)` with these manual contours.

Usage

```

analyze(
  x,
  samplingRate = NULL,
  scale = NULL,
  from = NULL,
  to = NULL,
  dynamicRange = 80,
  silence = 0.04,
  windowLength = 50,
  step = 25,
  overlap = 50,
  specType = c("spectrum", "reassign", "spectralDerivative")[1],
  wn = "gaussian",
  zp = 0,
  cutFreq = NULL,
  nFormants = 3,
  formants = list(),
  loudness = list(SPL_measured = 70),
  roughness = list(windowLength = 15, step = 3, amRes = 10),
  novelty = list(input = "melspec", kernellLen = 100),

```

```

pitchMethods = c("dom", "autocor"),
pitchManual = NULL,
entropyThres = 0.6,
pitchFloor = 75,
pitchCeiling = 3500,
priorMean = 300,
priorSD = 6,
priorAdapt = TRUE,
nCands = 1,
minVoicedCands = NULL,
pitchDom = list(domThres = 0.1, domSmooth = 220),
pitchAutocor = list(autocorThres = 0.7, autocorSmooth = 7, autocorUpsample = 25,
  autocorBestPeak = 0.975, interpol = "sinc", wn = "hanning"),
pitchCep = list(cepThres = 0.75, cepZp = 0),
pitchSpec = list(specThres = 0.05, specPeak = 0.25, specHNRSlope = 0.8, specSmooth =
  150, specMerge = 0.1, specSinglePeakCert = 0.4, specRatios = 3),
pitchHps = list(hpsNum = 5, hpsThres = 0.1, hpsNorm = 2, hpsPenalty = 2),
pitchZc = list(zcThres = 0.1, zcWin = 5),
harmHeight = list(harmThres = 3, harmTol = 0.25, harmPerSel = 5),
subh = list(method = c("cep", "pitchCands", "harm")[1], nSubh = 5, tol = 0.05, nHarm =
  5, harmThres = 12, harmTol = 0.25, amRange = c(10, 200)),
flux = list(thres = 0.15, smoothWin = 100),
amRange = c(10, 200),
fmRange = c(5, 1000/step/2),
shortestSyl = 20,
shortestPause = 60,
interpol = list(win = 75, tol = 0.3, cert = 0.3),
pathfinding = c("none", "fast", "slow")[2],
annealPars = list(maxit = 5000, temp = 1000),
certWeight = 0.5,
snakeStep = 0,
snakePlot = FALSE,
smooth = 1,
smoothVars = c("pitch", "dom"),
summaryFun = c("mean", "median", "sd"),
invalidArgAction = c("adjust", "abort", "ignore")[1],
reportEvery = NULL,
cores = 1,
plot = FALSE,
osc = "linear",
showLegend = TRUE,
savePlots = NULL,
pitchPlot = list(col = rgb(0, 0, 1, 0.75), lwd = 3, showPrior = TRUE),
extraContour = NULL,
ylim = NULL,
xlab = "Time",
ylab = NULL,
main = NULL,

```

```

width = 900,
height = 500,
units = "px",
res = NA,
...
)

```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
scale	maximum possible amplitude of input used for normalization of input vector (only needed if x is a numeric vector)
from, to	if NULL (default), analyzes the whole sound, otherwise from...to (s)
dynamicRange	dynamic range, dB. All values more than one dynamicRange under maximum are treated as zero
silence	(0 to 1 as proportion of max amplitude) frames with RMS amplitude below $\text{silence} * \text{max_ampl}$ adjusted by scale are not analyzed at all.
windowLength	length of FFT window, ms
step	you can override overlap by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
overlap	overlap between successive FFT frames, %
specType	plot the original FFT ('spectrum'), reassigned spectrogram ('reassigned'), or spectral derivative ('spectralDerivative')
wn	window type accepted by ftwindow , currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop
zp	window length after zero padding, points
cutFreq	if specified, spectral descriptives (peakFreq, specCentroid, specSlope, and quartiles) are calculated only between cutFreq[1] and cutFreq[2], Hz. If a single number is given, analyzes frequencies from 0 to cutFreq. For ex., when analyzing recordings with varying sampling rates, set to half the lowest sampling rate to make the spectra more comparable. Note that "entropyThres" applies only to this frequency range, which also affects which frames will not be analyzed with pitchAutocor.
nFormants	the number of formants to extract per STFT frame (0 = no formant analysis, NULL = as many as possible)
formants	a list of arguments passed to findformants - an external function called to perform LPC analysis
loudness	a list of parameters passed to getLoudness for measuring subjective loudness, namely SPL_measured, Pref, spreadSpectrum. NULL = skip loudness analysis

roughness	a list of parameters passed to <code>modulationSpectrum</code> for measuring roughness. NULL = skip roughness analysis
novelty	a list of parameters passed to <code>ssm</code> for measuring spectral novelty. NULL = skip novelty analysis
pitchMethods	methods of pitch estimation to consider for determining pitch contour: 'autocor' = autocorrelation (~PRAAT), 'cep' = cepstral, 'spec' = spectral (~BaNa), 'dom' = lowest dominant frequency band, 'hps' = harmonic product spectrum, NULL = no pitch analysis
pitchManual	manually corrected pitch contour. For a single sound, provide a numeric vector of any length. For multiple sounds, provide a dataframe with columns "file" and "pitch" (or path to a csv file) as returned by <code>pitch_app</code> , ideally with the same <code>windowLength</code> and <code>step</code> as in current call to <code>analyze</code> . A named list with pitch vectors per file is also OK (eg as returned by <code>pitch_app</code>)
entropyThres	pitch tracking is only performed for frames with Weiner entropy below <code>entropyThres</code> , but other spectral descriptives are still calculated (NULL = analyze everything)
pitchFloor, pitchCeiling	absolute bounds for pitch candidates (Hz)
priorMean, priorSD	specifies the mean (Hz) and standard deviation (semitones) of gamma distribution describing our prior knowledge about the most likely pitch values for this file. For ex., <code>priorMean = 300, priorSD = 6</code> gives a prior with mean = 300 Hz and SD = 6 semitones (half an octave). To avoid using any priors, set <code>priorMean = NA, priorAdapt = FALSE</code>
priorAdapt	adaptive second-pass prior: if TRUE, optimal pitch contours are estimated first with a prior determined by <code>priorMean, priorSD</code> , and then with a new prior adjusted according to this first-pass pitch contour
nCands	maximum number of pitch candidates per method, normally 1..4 (except for <code>dom</code> , which returns at most one candidate per frame)
minVoicedCands	minimum number of pitch candidates that have to be defined to consider a frame voiced (if NULL, defaults to 2 if <code>dom</code> is among other candidates and 1 otherwise)
pitchDom	a list of control parameters for pitch tracking using the lowest dominant frequency band or "dom" method; see details and <code>?soundgen:::getDom</code>
pitchAutocor	a list of control parameters for pitch tracking using the autocorrelation or "autocor" method; see details and <code>?soundgen:::getPitchAutocor</code>
pitchCep	a list of control parameters for pitch tracking using the cepstrum or "cep" method; see details and <code>?soundgen:::getPitchCep</code>
pitchSpec	a list of control parameters for pitch tracking using the BaNa or "spec" method; see details and <code>?soundgen:::getPitchSpec</code>
pitchHps	a list of control parameters for pitch tracking using the harmonic product spectrum or "hps" method; see details and <code>?soundgen:::getPitchHps</code>
pitchZc	a list of control parameters for pitch tracking based on zero crossings in bandpass-filtered audio or "zc" method; see <code>getPitchZc</code>
harmHeight	a list of control parameters for estimating how high harmonics reach in the spectrum; see details and <code>?soundgen:::harmHeight</code>

subh	a list of control parameters for estimating the strength of subharmonics per frame - that is, spectral energy at integer ratios of f0: see <code>?soundgen:::subhToHarm</code>
flux	a list of control parameters for calculating feature-based flux (not spectral flux) passed to <code>getFeatureFlux</code>
amRange	target range of frequencies for amplitude modulation, Hz: a vector of length 2 (affects both <code>amMsFreq</code> and <code>amEnvFreq</code>)
fmRange	target range of frequencies for analyzing frequency modulation, Hz (<code>fmFreq</code>): a vector of length 2
shortestSyl	the smallest length of a voiced segment (ms) that constitutes a voiced syllable (shorter segments will be replaced by NA, as if unvoiced)
shortestPause	the smallest gap between voiced syllables (ms): large value = interpolate and merge, small value = treat as separate syllables separated by an unvoiced gap
interpol	a list of parameters (currently <code>win</code> , <code>tol</code> , <code>cert</code>) passed to <code>soundgen:::pathfinder</code> for interpolating missing pitch candidates (NULL = no interpolation)
pathfinding	method of finding the optimal path through pitch candidates: 'none' = best candidate per frame, 'fast' = simple heuristic, 'slow' = annealing. See <code>soundgen:::pathfinder</code>
annealPars	a list of control parameters for postprocessing of pitch contour with SANN algorithm of <code>optim</code> . This is only relevant if <code>pathfinding = 'slow'</code>
certWeight	(0 to 1) in pitch postprocessing, specifies how much we prioritize the certainty of pitch candidates vs. pitch jumps / the internal tension of the resulting pitch curve
snakeStep	optimized path through pitch candidates is further processed to minimize the elastic force acting on pitch contour. To disable, set <code>snakeStep = 0</code>
snakePlot	if TRUE, plots the snake
smooth, smoothVars	if <code>smooth</code> is a positive number, outliers of the variables in <code>smoothVars</code> are adjusted with median smoothing. <code>smooth</code> of 1 corresponds to a window of ~100 ms and tolerated deviation of ~4 semitones. To disable, set <code>smooth = 0</code>
summaryFun	functions used to summarize each acoustic characteristic, eg <code>"c('mean', 'sd')"</code> ; user-defined functions are fine (see examples); NAs are omitted automatically for <code>mean/median/sd/min/max/range/sum</code> , otherwise take care of NAs yourself
invalidArgAction	what to do if an argument is invalid or outside the range in <code>defaults_analyze</code> : 'adjust' = reset to default value, 'abort' = stop execution, 'ignore' = throw a warning and continue (may crash)
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
plot	if TRUE, produces a spectrogram with pitch contour overlaid
osc	"none" = no oscillogram; "linear" = on the original scale; "dB" = in decibels
showLegend	if TRUE, adds a legend with pitch tracking methods
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)

pitchPlot	a list of graphical parameters for displaying the final pitch contour. Set to <code>list(type = 'n')</code> to suppress
extraContour	name of an output variable to overlap on the pitch contour plot, eg 'peakFreq' or 'loudness'; can also be a list with extra graphical parameters, eg <code>extraContour = list(x = 'harmHeight', col = 'red')</code>
ylim	frequency range to plot, kHz (defaults to 0 to Nyquist frequency). NB: still in kHz, even if <code>yScale = bark, mel, or ERB</code>
xlab, ylab, main	plotting parameters
width, height, units, res	parameters passed to png if the plot is saved
...	other graphical parameters passed to spectrogram

Details

Each pitch tracker is controlled by its own list of settings, as follows:

- pitchDom (lowest dominant frequency band)** • `domThres` (0 to 1) to find the lowest dominant frequency band, we do short-term FFT and take the lowest frequency with amplitude at least `domThres`
- `domSmooth` the width of smoothing interval (Hz) for finding `dom`
- pitchAutocor (autocorrelation)** • `autocorThres` voicing threshold (unitless, ~0 to 1)
- `autocorSmooth` the width of smoothing interval (in bins) for finding peaks in the autocorrelation function. Defaults to 7 for sampling rate 44100 and smaller odd numbers for lower values of sampling rate
 - `autocorUpsample` upsamples `acf` to this resolution (Hz) to improve accuracy in high frequencies
 - `autocorBestPeak` amplitude of the lowest best candidate relative to the absolute max of the `acf`
- pitchCep (cepstrum)** • `cepThres` voicing threshold (unitless, ~0 to 1)
- `cepZp` zero-padding of the spectrum used for cepstral pitch detection (final length of spectrum after zero-padding in points, e.g. 2^{13})
- pitchSpec (ratio of harmonics - BaNa algorithm)** • `specThres` voicing threshold (unitless, ~0 to 1)
- `specPeak`, `specHNRSlope` when looking for putative harmonics in the spectrum, the threshold for peak detection is calculated as `specPeak * (1 - HNR * specHNRSlope)`
 - `specSmooth` the width of window for detecting peaks in the spectrum, Hz
 - `specMerge` pitch candidates within `specMerge` semitones are merged with boosted certainty
 - `specSinglePeakCert` (0 to 1) if `F0` is calculated based on a single harmonic ratio (as opposed to several ratios converging on the same candidate), its certainty is taken to be `specSinglePeakCert`
- pitchHps (harmonic product spectrum)** • `hpsNum` the number of times to downsample the spectrum
- `hpsThres` voicing threshold (unitless, ~0 to 1)

- `hpsNorm` the amount of inflation of hps pitch certainty (0 = none)
- `hpsPenalty` the amount of penalizing hps candidates in low frequencies (0 = none)

Each of these lists also accepts graphical parameters that affect how pitch candidates are plotted, eg `pitchDom = list(domThres = .5, col = 'yellow')`. Other arguments that are lists of subroutine-specific settings include:

`harmonicHeight` (**finding how high harmonics reach in the spectrum**) • `harmThres` minimum height of spectral peak, dB

- `harmPerSel` the number of harmonics per sliding selection
- `harmTol` maximum tolerated deviation of peak frequency from multiples of `f0`, proportion of `f0`

Value

Returns a list with `$detailed` frame-by-frame descriptives and a `$summary` with one row per file, as determined by `summaryFun` (e.g., mean / median / SD of each acoustic variable across all STFT frames). Output measures include:

duration total duration, s

duration_noSilence duration from the beginning of the first non-silent STFT frame to the end of the last non-silent STFT frame, s (NB: depends strongly on `windowLength` and silence settings)

time time of the middle of each frame (ms)

amEnvFreq,amEnvDep frequency (Hz) and depth (0 to 1) of amplitude modulation estimated from a smoothed amplitude envelope

amMsFreq,amMsPurity frequency and purity of amplitude modulation estimated via `modulationSpectrum`

ampl root mean square of amplitude per frame, calculated as `sqrt(mean(frame ^ 2))`

ampl_noSilence same as `ampl`, but ignoring silent frames

CPP Cepstral Peak Prominence, dB (a measure of pitch quality, the ratio of the highest peak in the cepstrum to the regression line drawn through it)

dom lowest dominant frequency band (Hz) (see "Pitch tracking methods / Dominant frequency" in the vignette)

entropy Weiner entropy of the spectrum of the current frame. Close to 0: pure tone or tonal sound with nearly all energy in harmonics; close to 1: white noise

entropySh Normalized Shannon entropy of the spectrum of the current frame: 0 = pure tone, 1 = white noise

f1_freq, f1_width, ... the frequency and bandwidth of the first `nFormants` formants per STFT frame, as calculated by `phonTools::findformants`

flux feature-based flux, the rate of change in acoustic features such as pitch, HNR, etc. (0 = none, 1 = max); "epoch" is an audio segment between two peaks of flux that exceed a threshold of `flux = list(thres = ...)` (listed in `output$detailed` only)

fmFreq frequency of frequency modulation (FM) such as vibrato or jitter, Hz

fmDep depth of FM, semitones

fmPurity purity or dominance of the main FM frequency (`fmFreq`), 0 to 1

harmEnergy the amount of energy in upper harmonics, namely the ratio of total spectral mass above $1.25 \times F_0$ to the total spectral mass below $1.25 \times F_0$ (dB)

harmHeight how high harmonics reach in the spectrum, based on the best guess at pitch (or the manually provided pitch values)

HNR harmonics-to-noise ratio (dB), a measure of harmonicity returned by `soundgen:::getPitchAutocor` (see "Pitch tracking methods / Autocorrelation"). If HNR = 0 dB, there is as much energy in harmonics as in noise

loudness subjective loudness, in sone, corresponding to the chosen `SPL_measured` - see [getLoudness](#)

novelty spectral novelty - a measure of how variable the spectrum is on a particular time scale, as estimated by [ssm](#)

peakFreq the frequency with maximum spectral power (Hz)

pitch post-processed pitch contour based on all F_0 estimates

quartile25, quartile50, quartile75 the 25th, 50th, and 75th quantiles of the spectrum of voiced frames (Hz)

roughness the amount of amplitude modulation, see `modulationSpectrum`

specCentroid the center of gravity of the frame's spectrum, first spectral moment (Hz)

specSlope the slope of linear regression fit to the spectrum below `cutFreq` (dB/kHz)

subDep estimated depth of subharmonics per frame: 0 = none, 1 = as strong as f_0 . NB: this depends critically on accurate pitch tracking

subRatio the ratio of f_0 to subharmonics frequency with strength `subDep`: 2 = period doubling, 3 = $f_0 / 3$, etc.

voiced is the current STFT frame voiced? TRUE / FALSE

See Also

[pitch_app](#) [getLoudness](#) [segment](#) [getRMS](#)

Examples

```

sound = soundgen(syllLen = 300, pitch = c(500, 400, 600),
  noise = list(time = c(0, 300), value = c(-40, 0)),
  temperature = 0.001,
  addSilence = 50) # NB: always have some silence before and after!!!
# playme(sound, 16000)
a = analyze(sound, samplingRate = 16000, plot = TRUE)
str(a$detailed) # frame-by-frame
a$summary      # summary per sound

## Not run:
# For maximum processing speed (just basic spectral descriptives):
a = analyze(sound, samplingRate = 16000,
  plot = FALSE,          # no plotting
  pitchMethods = NULL,  # no pitch tracking
  loudness = NULL,      # no loudness analysis
  novelty = NULL,       # no novelty analysis
  roughness = NULL,     # no roughness analysis

```

```

nFormants = 0          # no formant analysis
)

# Take a sound hard to analyze b/c of subharmonics and jitter
sound2 = soundgen(syllLen = 900, pitch = list(
  time = c(0, .3, .8, 1), value = c(300, 900, 400, 2300)),
  noise = list(time = c(0, 900), value = c(-40, -20)),
  subDep = 10, jitterDep = 0.5,
  temperature = 0.001, samplingRate = 44100, pitchSamplingRate = 44100)
# playme(sound2, 44100)
a2 = analyze(sound2, samplingRate = 44100, priorSD = 24,
  plot = TRUE, ylim = c(0, 5))

# Compare the available pitch trackers
analyze(sound2, 44100,
  pitchMethods = c('dom', 'autocor', 'spec', 'cep', 'hps', 'zc'),
  # don't use priors to see weird pitch candidates better
  priorMean = NA, priorAdapt = FALSE,
  plot = TRUE, yScale = 'bark')

# Fancy plotting options:
a = analyze(sound2, samplingRate = 44100, plot = TRUE,
  xlab = 'Time, ms', colorTheme = 'seewave',
  contrast = .5, ylim = c(0, 4), main = 'My plot',
  pitchMethods = c('dom', 'autocor', 'spec', 'hps', 'cep'),
  priorMean = NA, # no prior info at all
  pitchDom = list(col = 'red', domThres = .25),
  pitchPlot = list(col = 'black', pch = 9, lty = 3, lwd = 3),
  extraContour = list(x = 'peakFreq', type = 'b', pch = 4, col = 'brown'),
  osc = 'dB', heights = c(2, 1))

# Analyze an entire folder in one go, saving spectrograms with pitch contours
# plus an html file for easy access
s2 = analyze('~Downloads/temp',
  savePlots = '', # save in the same folder as audio
  showLegend = TRUE, yScale = 'bark',
  width = 20, height = 12,
  units = 'cm', res = 300, ylim = c(0, 5))
s2$summary[, 1:5]

# Different options for summarizing the output
a = analyze(sound1, 44100,
  summaryFun = c('mean', 'range'))
a$summary # one row per sound
# ...with custom summaryFun, eg time of peak relative to duration (0 to 1)
timePeak = function(x) which.max(x) / length(x) # without omitting NAs
timeTrough = function(x) which.min(x) / length(x)
a = analyze(sound2, samplingRate = 16000,
  summaryFun = c('mean', 'timePeak', 'timeTrough'))
colnames(a$summary)

# Analyze a selection rather than the whole sound
a = analyze(sound, samplingRate = 16000, from = .1, to = .3, plot = TRUE)

```

```

# Use only a range of frequencies when calculating spectral descriptives
# (ignore everything below 100 Hz and above 8000 Hz as irrelevant noise)
a = analyze(sound, samplingRate = 16000, cutFreq = c(100, 8000))

## Amplitude and loudness: analyze() should give the same results as
# dedicated functions getRMS() / getLoudness()
# Create 1 kHz tone
samplingRate = 16000; dur_ms = 50
sound3 = sin(2*pi*1000/samplingRate*(1:(dur_ms/1000*samplingRate)))
a1 = analyze(sound3, samplingRate = samplingRate, scale = 1,
             windowLength = 25, overlap = 50,
             loudness = list(SPL_measured = 40),
             pitchMethods = NULL, plot = FALSE)
a1$detailed$loudness # loudness per STFT frame (1 sone by definition)
getLoudness(sound3, samplingRate = samplingRate, windowLength = 25,
            overlap = 50, SPL_measured = 40, scale = 1)$loudness
a1$detailed$ampl # RMS amplitude per STFT frame
getRMS(sound3, samplingRate = samplingRate, windowLength = 25,
        overlap = 50, scale = 1)$detailed
# or even simply: sqrt(mean(sound3 ^ 2))

# The same sound as above, but with half the amplitude
a_half = analyze(sound3 / 2, samplingRate = samplingRate, scale = 1,
                windowLength = 25, overlap = 50,
                loudness = list(SPL_measured = 40),
                pitchMethods = NULL, plot = FALSE)
a1$detailed$ampl / a_half$detailed$ampl # rms amplitude halved
a1$detailed$loudness / a_half$detailed$loudness
# loudness is not a linear function of amplitude

# Analyzing ultrasounds (slow but possible, just adjust pitchCeiling)
s = soundgen(syllen = 100, addSilence = 10,
            pitch = c(25000, 35000, 30000),
            formants = NA, rolloff = -12, rolloffKHz = 0,
            pitchSamplingRate = 350000, samplingRate = 350000, windowLength = 5,
            pitchCeiling = 45000, invalidArgAction = 'ignore')
# s is a bat-like ultrasound inaudible to humans
a = analyze(s, 350000, plot = TRUE,
            pitchCeiling = 45000, priorMean = NA,
            windowLength = 2, overlap = 0,
            nFormants = 0, loudness = NULL)
# NB: ignore formants and loudness estimates for such non-human sounds

# download 260 sounds from Anikin & Persson (2017)
# http://cogsci.se/publications/anikin-persson_2017_nonlinguistic-vocs/260sounds_wav.zip
# unzip them into a folder, say '~/Downloads/temp'
myfolder = '~/Downloads/temp' # 260 .wav files live here
s = analyze(myfolder) # ~ 10-20 minutes!
# s = write.csv(s, paste0(myfolder, '/temp.csv')) # save a backup

# Check accuracy: import manually verified pitch values (our "key")
# pitchManual # "ground truth" of mean pitch per sound

```

```

# pitchContour # "ground truth" of complete pitch contours per sound
files_manual = paste0(names(pitchManual), '.wav')
idx = match(s$file, files_manual) # in case the order is wrong
s$key = pitchManual[idx]

# Compare manually verified mean pitch with the output of analyze:
cor(s$key, s$summary$pitch_median, use = 'pairwise.complete.obs')
plot(s$key, s$summary$pitch_median, log = 'xy')
abline(a=0, b=1, col='red')

# Re-running analyze with manually corrected contours gives correct
pitch-related descriptives like amplVoiced and harmonics (NB: you get it "for
free" when running pitch_app)
s1 = analyze(myfolder, pitchManual = pitchContour)
plot(s$summary$harmonics_median, s1$summary$harmonics_median)
abline(a=0, b=1, col='red')

## End(Not run)

```

annotation_app

Annotation app

Description

Starts a shiny app for annotating audio. This is a simplified and faster version of [formant_app](#) intended only for making annotations.

Usage

```
annotation_app()
```

Value

Every time a new annotation is added, the app creates a backup csv file and creates or updates a global object called "my_annot", which contains all the annotations. When the app is terminated, it also returns the results as a dataframe.

Examples

```

## Not run:
ann = annotation_app() # runs in default browser such as Firefox or Chrome

# To change system default browser, run something like:
options('browser' = '/usr/bin/firefox') # path to the executable on Linux

## End(Not run)

```

audSpectrogram *Auditory spectrogram*

Description

Produces an auditory spectrogram by extracting a bank of bandpass filters (work in progress). While `tuneR::audspec` is based on FFT, here we convolve the sound with a bank of filters. The main difference is that we don't window the signal and de facto get variable temporal resolution in different frequency channels, as with a wavelet transform. The filters are currently third-order Butterworth bandpass filters implemented in [butter](#).

Usage

```
audSpectrogram(
  x,
  samplingRate = NULL,
  scale = NULL,
  from = NULL,
  to = NULL,
  step = 1,
  dynamicRange = 80,
  nFilters = 128,
  minFreq = 20,
  maxFreq = samplingRate/2,
  minBandwidth = 10,
  reportEvery = NULL,
  cores = 1,
  plot = TRUE,
  savePlots = NULL,
  osc = c("none", "linear", "dB")[2],
  heights = c(3, 1),
  ylim = NULL,
  yScale = c("bark", "mel", "ERB", "log")[1],
  contrast = 0.2,
  brightness = 0,
  maxPoints = c(1e+05, 5e+05),
  padWithSilence = TRUE,
  colorTheme = c("bw", "seewave", "heat.colors", "...")[1],
  col = NULL,
  extraContour = NULL,
  xlab = NULL,
  ylab = NULL,
  xaxp = NULL,
  mar = c(5.1, 4.1, 4.1, 2),
  main = NULL,
  grid = NULL,
  width = 900,
```

```

    height = 500,
    units = "px",
    res = NA,
    ...
)

```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>scale</code>	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
<code>from, to</code>	if NULL (default), analyzes the whole sound, otherwise from...to (s)
<code>step</code>	step, ms (determines time resolution). <code>step = NULL</code> means no downsampling at all (ncol of output = length of input audio)
<code>dynamicRange</code>	dynamic range, dB. All values more than one <code>dynamicRange</code> under maximum are treated as zero
<code>nFilters</code>	the number of filters (determines frequency resolution)
<code>minFreq, maxFreq</code>	the range of frequencies to analyze
<code>minBandwidth</code>	minimum filter bandwidth, Hz (otherwise filters may become too narrow when <code>nFilters</code> is high)
<code>reportEvery</code>	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
<code>cores</code>	number of cores for parallel processing
<code>plot</code>	should a spectrogram be plotted? TRUE / FALSE
<code>savePlots</code>	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
<code>osc</code>	"none" = no oscillogram; "linear" = on the original scale; "dB" = in decibels
<code>heights</code>	a vector of length two specifying the relative height of the spectrogram and the oscillogram (including time axes labels)
<code>ylim</code>	frequency range to plot, kHz (defaults to 0 to Nyquist frequency). NB: still in kHz, even if <code>yScale = bark, mel, or ERB</code>
<code>yScale</code>	scale of the frequency axis: 'linear' = linear, 'log' = logarithmic (musical), 'bark' = bark with hz2bark , 'mel' = mel with hz2mel , 'ERB' = Equivalent Rectangular Bandwidths with HzToERB
<code>contrast</code>	spectrum is exponentiated by contrast (any real number, recommended -1 to +1). Contrast >0 increases sharpness, <0 decreases sharpness
<code>brightness</code>	how much to "lighten" the image (>0 = lighter, <0 = darker)
<code>maxPoints</code>	the maximum number of "pixels" in the oscillogram (if any) and spectrogram; good for quickly plotting long audio files; defaults to c(1e5, 5e5)

padWithSilence if TRUE, pads the sound with just enough silence to resolve the edges properly (only the original region is plotted, so the apparent duration doesn't change)
colorTheme black and white ('bw'), as in seewave package ('seewave'), or any palette from [palette](#) such as 'heat.colors', 'cm.colors', etc
col actual colors, eg rev(rainbow(100)) - see ?hcl.colors for colors in base R (overrides colorTheme)
extraContour a vector of arbitrary length scaled in Hz (regardless of yScale!) that will be plotted over the spectrogram (eg pitch contour); can also be a list with extra graphical parameters such as lwd, col, etc. (see examples)
xlab, ylab, main, mar, xaxp graphical parameters for plotting
grid if numeric, adds n = grid dotted lines per kHz
width, height, units, res graphical parameters for saving plots passed to [png](#)
... other graphical parameters

Examples

```

# synthesize a sound with gradually increasing hissing noise
sound = soundgen(syllLen = 200, temperature = 0.001,
  noise = list(time = c(0, 350), value = c(-40, 0)),
  formantsNoise = list(f1 = list(freq = 5000, width = 10000)),
  addSilence = 25)
# playme(sound, samplingRate = 16000)

# auditory spectrogram
as = audSpectrogram(sound, samplingRate = 16000, nFilters = 48)
dim(as$audSpec)

# compare to FFT-based spectrogram with similar time and frequency resolution
fs = spectrogram(sound, samplingRate = 16000, yScale = 'bark',
  windowLength = 5, step = 1)
dim(fs)

## Not run:
# add bells and whistles
audSpectrogram(sound, samplingRate = 16000,
  yScale = 'ERB',
  osc = 'dB', # plot oscillogram in dB
  heights = c(2, 1), # spectro/osc height ratio
  brightness = -.1, # reduce brightness
  # colorTheme = 'heat.colors', # pick color theme...
  col = hcl.colors(30, palette = 'Plasma'), # ...or specify the colors
  cex.lab = .75, cex.axis = .75, # text size and other base graphics pars
  grid = 5, # to customize, add manually with graphics::grid()
  ylim = c(0.1, 5), # always in kHz
  main = 'My auditory spectrogram' # title
  # + axis labels, etc
)

```



```

# change dynamic range
audSpectrogram(sound, samplingRate = 16000, dynamicRange = 40)
audSpectrogram(sound, samplingRate = 16000, dynamicRange = 120)

# remove the oscillogram
audSpectrogram(sound, samplingRate = 16000, osc = 'none')

# save auditory spectrograms of all audio files in a folder
audSpectrogram('~Downloads/temp',
  savePlots = '~/Downloads/temp/audSpec', cores = 4)

## End(Not run)

```

bandpass

Bandpass/stop filters

Description

Filtering in the frequency domain with FFT-iFFT: low-pass, high-pass, bandpass, and bandstop filters. Similar to [ffilter](#), but here we use FFT instead of STFT - that is, the entire sound is processed at once. This works best for relatively short sounds (seconds), but gives us maximum precision (e.g., for precise notch filtering) and doesn't affect the attack and decay. NAs are accepted and can be interpolated or preserved in the output. Because we don't do STFT, arbitrarily short vectors are also fine as input - for example, we can apply a low-pass filter prior to decimation when changing the sampling rate without aliasing. Note that, unlike [pitchSmoothPraat](#), bandpass by default applies an abrupt cutoff instead of a smooth gaussian filter, but this behavior can be adjusted with the `bw` argument.

Usage

```

bandpass(
  x,
  samplingRate = NULL,
  lwr = NULL,
  upr = NULL,
  action = c("pass", "stop")[1],
  dB = Inf,
  bw = 0,
  na.rm = TRUE,
  from = NULL,
  to = NULL,
  normalize = FALSE,
  reportEvery = NULL,
  cores = 1,
  saveAudio = NULL,
  plot = FALSE,
  savePlots = NULL,
  width = 900,

```

```

    height = 500,
    units = "px",
    res = NA,
    ...
)

```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>lwr</code> , <code>upr</code>	cutoff frequencies, Hz. Specifying just <code>lwr</code> gives a high-pass filter, just <code>upr</code> low-pass filter with <code>action = 'pass'</code> (or vice versa with <code>action = 'stop'</code>). Specifying both <code>lwr</code> and <code>upr</code> a bandpass/bandstop filter, depending on <code>'action'</code>
<code>action</code>	"pass" = preserve the selected frequency range (bandpass), "stop" = remove the selected frequency range (bandstop)
<code>dB</code>	a positive number giving the strength of effect in dB (defaults to Inf - complete removal of selected frequencies)
<code>bw</code>	bandwidth of the filter cutoffs, Hz. Defaults to 0 (abrupt, step function), a positive number corresponds to the standard deviation of a Gaussian curve, and two numbers set different bandwidths for the lower and upper cutoff points
<code>na.rm</code>	if TRUE, NAs are interpolated, otherwise they are preserved in the output
<code>from</code> , <code>to</code>	if NULL (default), analyzes the whole sound, otherwise from...to (s)
<code>normalize</code>	if TRUE, resets the output to the original scale (otherwise filtering often reduces the amplitude)
<code>reportEvery</code>	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
<code>cores</code>	number of cores for parallel processing
<code>saveAudio</code>	full path to the folder in which to save the processed audio
<code>plot</code>	should a spectrogram be plotted? TRUE / FALSE
<code>savePlots</code>	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
<code>width</code> , <code>height</code> , <code>units</code> , <code>res</code>	graphical parameters for saving plots passed to png
<code>...</code>	other graphical parameters passed to <code>plot()</code> as well as to meanspec

Details

Algorithm: fill in NAs with constant interpolation at the edges and linear interpolation in the middle; perform FFT; set the frequency ranges to be filtered out to 0; perform inverse FFT; set to the original scale; put the NAs back in.

Examples

```

# Filter white noise
s1 = fade(c(runif(2000, -1, 1)), samplingRate = 16000)

# low-pass
bandpass(s1, 16000, upr = 2000, plot = TRUE)

# high-pass by 40 dB
bandpass(s1, 16000, lwr = 2000, dB = 40, plot = TRUE, wl = 1024)
# wl is passed to seewave::meanspec for plotting

# bandstop
bandpass(s1, 16000, lwr = 1000, upr = 1800, action = 'stop', plot = TRUE)

# bandpass
s2 = bandpass(s1, 16000, lwr = 2000, upr = 2100, plot = TRUE)
# playme(rep(s2, 5))
# spectrogram(s2, 16000)

# low-pass and interpolate a short vector with some NAs
x = rnorm(150, 10) + 3 * sin((1:50) / 5)
x[sample(1:length(x), 50)] = NA
plot(x, type = 'l')
x_bandp = bandpass(x, samplingRate = 100, upr = 10)
points(x_bandp, type = 'l', col = 'blue')

## Not run:
# add 200 dB with a Gaussian-shaped filter instead of step function
s3 = bandpass(s1, 16000, lwr = 1700, upr = 2100, bw = 200,
  dB = 20, plot = TRUE)
spectrogram(s3, 16000)
s4 = bandpass(s1, 16000, lwr = 2000, upr = 4300, bw = c(100, 500),
  dB = 60, action = 'stop', plot = TRUE)
spectrogram(s4, 16000)

# precise notch filtering is possible, even in low frequencies
whiteNoise = runif(16000, -1, 1)
s3 = bandpass(whiteNoise, 16000, lwr = 30, upr = 40, normalize = TRUE,
  plot = TRUE, xlim = c(0, 500))
playme(rep(s3, 5))
spectrogram(s3, 16000, windowLength = 150, yScale = 'log')

# compare the same with STFT
s4 = seewave::ffilter(whiteNoise, f = 16000, from = 30, to = 40)
spectrogram(s4, 16000, windowLength = 150, yScale = 'log')
# (note: works better as wl approaches length(s4))

# high-pass all audio files in a folder
bandpass('~Downloads/temp', saveAudio = '~Downloads/temp/hp2000/',
  lwr = 2000, savePlots = '~Downloads/temp/hp2000/')

## End(Not run)

```

 beat

Generate beat

Description

Generates percussive sounds from clicks through drum-like beats to sliding tones. The principle is to create a sine wave with rapid frequency modulation and to add a fade-out. No extra harmonics or formants are added. For this specific purpose, this is vastly faster and easier than to tinker with [soundgen](#) settings, especially since percussive syllables tend to be very short.

Usage

```
beat(
  nSyl = 10,
  sylLen = 200,
  pauseLen = 50,
  pitch = c(200, 10),
  samplingRate = 16000,
  fadeOut = TRUE,
  play = FALSE
)
```

Arguments

<code>nSyl</code>	the number of syllables to generate
<code>sylLen</code>	average duration of each syllable, ms
<code>pauseLen</code>	average duration of pauses between syllables, ms
<code>pitch</code>	fundamental frequency, Hz - a vector or <code>data.frame(time = ..., value = ...)</code>
<code>samplingRate</code>	sampling frequency, Hz
<code>fadeOut</code>	if TRUE, a linear fade-out is applied to the entire syllable
<code>play</code>	if TRUE, plays the synthesized sound using the default player on your system. If character, passed to <code>play</code> as the name of player to use, eg "aplay", "play", "vlc", etc. In case of errors, try setting another default player for <code>play</code>

Value

Returns a non-normalized waveform centered at zero.

See Also

[soundgen generateNoise fart](#)

Examples

```

playback = c(TRUE, FALSE)[2]
# a drum-like sound
s = beat(nSyl = 1, syllLen = 200,
        pitch = c(200, 100), play = playback)
# plot(s, type = 'l')

# a dry, muted drum
s = beat(nSyl = 1, syllLen = 200,
        pitch = c(200, 10), play = playback)

# sci-fi laser guns
s = beat(nSyl = 3, syllLen = 300,
        pitch = c(1000, 50), play = playback)

# machine guns
s = beat(nSyl = 10, syllLen = 10, pauseLen = 50,
        pitch = c(2300, 300), play = playback)

```

compareSounds

Compare two sounds

Description

Computes similarity between two sounds based on comparing their spectrogram-like representations. If the input is audio, two methods of producing spectrograms are available: `specType = 'linear'` calls `powspec` for an power spectrogram with frequencies in Hz, and `specType = 'mel'` calls `melfcc` for an auditory spectrogram with frequencies in Mel. For more customized options, just produce your spectrograms or feature matrices (time in column, features like pitch, peak frequency etc in rows) with your favorite function before calling `compareSounds` because it also accepts matrices as input. To be directly comparable, the two matrices are made into matrices of the same size. In case of differences in sampling rates, only frequencies below the lower Nyquist frequency or below `maxFreq` are kept. In case of differences in duration, the shorter sound is padded with 0 (silence) or NA, as controlled by arguments `padWith`, `padDir`. Then the matrices are compared using methods like cross-correlation or Dynamic Time Warp.

Usage

```

compareSounds(
  x,
  y,
  samplingRate = NULL,
  windowLength = 40,
  overlap = 50,
  step = NULL,
  dynamicRange = 80,
  method = c("cor", "cosine", "diff", "dtw"),
  specType = c("linear", "mel")[2],

```

```

specPars = list(),
dtwPars = list(),
padWith = NA,
padDir = c("central", "left", "right")[1],
maxFreq = NULL
)

```

Arguments

<code>x, y</code>	either two matrices (spectrograms or feature matrices) or two sounds to be compared (numeric vectors, Wave objects, or paths to wav/mp3 files)
<code>samplingRate</code>	if one or both inputs are numeric vectors, specify sampling rate, Hz. A vector of length 2 means the two inputs have different sampling rates, in which case spectrograms are compared only up to the lower Nyquist frequency
<code>windowLength</code>	length of FFT window, ms
<code>overlap</code>	overlap between successive FFT frames, %
<code>step</code>	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
<code>dynamicRange</code>	parts of the spectra quieter than $-\text{dynamicRange}$ dB are not compared
<code>method</code>	method of comparing mel-transformed spectra of two sounds: "cor" = Pearson's correlation; "cosine" = cosine similarity; "diff" = absolute difference between each bin in the two spectrograms; "dtw" = multivariate Dynamic Time Warp with dtw
<code>specType</code>	"linear" = power spectrogram with powspec , "mel" = mel-frequency spectrogram with melfcc
<code>specPars</code>	a list of parameters passed to melfcc
<code>dtwPars</code>	a list of parameters passed to dtw
<code>padWith</code>	if the duration of <code>x</code> and <code>y</code> is not identical, the compared spectrograms are padded with either silence (<code>padWith = 0</code>) or with NA's (<code>padWith = NA</code>) to have the same number of columns. Padding with NA implies that only the overlapping part is of relevance, whereas padding with 0 means that the added silent part is also compared with the longer sound, usually resulting in lower similarity (see examples)
<code>padDir</code>	if padding, specify where to add zeros or NAs: before the sound ('left'), after the sound ('right'), or on both sides ('central')
<code>maxFreq</code>	parts of the spectra above <code>maxFreq</code> Hz are not compared

Value

Returns a dataframe with two columns: "method" for the method(s) used, and "sim" for the similarity between the two sounds calculated with that method. The range of similarity measures is [-1, 1] for "cor", [0, 1] for "cosine" and "diff", and (-Inf, Inf) for "dtw".

Examples

```

data(orni, peewit, package = 'seewave')
compareSounds(orni, peewit)
# spectrogram(orni); playme(orni)
# spectrogram(peewit); playme(peewit)

## Not run:
s1 = soundgen(formants = 'a', play = TRUE)
s2 = soundgen(formants = 'ae', play = TRUE)
s3 = soundgen(formants = 'eae', sylLen = 700, play = TRUE)
s4 = runif(8000, -1, 1) # white noise
compareSounds(s1, s2, samplingRate = 16000)
compareSounds(s1, s4, samplingRate = 16000)

# the central section of s3 is more similar to s1 than is the beg/eng of s3
compareSounds(s1, s3, samplingRate = 16000, padDir = 'left')
compareSounds(s1, s3, samplingRate = 16000, padDir = 'central')

# padding with 0 penalizes differences in duration, whereas padding with NA
# is like saying we only care about the overlapping part
compareSounds(s1, s3, samplingRate = 16000, padWith = 0)
compareSounds(s1, s3, samplingRate = 16000, padWith = NA)

# comparing linear (Hz) vs mel-spectrograms produces quite different results
compareSounds(s1, s3, samplingRate = 16000, specType = 'linear')
compareSounds(s1, s3, samplingRate = 16000, specType = 'mel')

# pass additional control parameters to dtw and melfcc
compareSounds(s1, s3, samplingRate = 16000,
              specPars = list(nbands = 128),
              dtwPars = list(dist.method = "Manhattan"))

# use feature matrices instead of spectrograms (time in columns, features in rows)
a1 = t(as.matrix(analyze(s1, samplingRate = 16000)$detailed))
a1 = a1[4:nrow(a1), ]; a1[is.na(a1)] = 0
a2 = t(as.matrix(analyze(s2, samplingRate = 16000)$detailed))
a2 = a2[4:nrow(a2), ]; a2[is.na(a2)] = 0
a4 = t(as.matrix(analyze(s4, samplingRate = 16000)$detailed))
a4 = a4[4:nrow(a4), ]; a4[is.na(a4)] = 0
compareSounds(a1, a2, method = c('cosine', 'dtw'))
compareSounds(a1, a4, method = c('cosine', 'dtw'))

# a demo for comparing different similarity metrics
target = soundgen(sylLen = 500, formants = 'a',
                  pitch = data.frame(time = c(0, 0.1, 0.9, 1),
                                      value = c(100, 150, 135, 100)),
                  temperature = 0.001)
spec1 = soundgen:::getMelSpec(target, samplingRate = 16000)

parsToTry = list(
  list(formants = 'i', # wrong
        pitch = data.frame(time = c(0, 1), # wrong

```

```

                                value = c(200, 300)),
list(formants = 'i',                                # wrong
     pitch = data.frame(time = c(0, 0.1, 0.9, 1),    # right
                          value = c(100, 150, 135, 100))),
list(formants = 'a',                                # right
     pitch = data.frame(time = c(0,1),              # wrong
                          value = c(200, 300))),
list(formants = 'a',                                # right
     pitch = data.frame(time = c(0, 0.1, 0.9, 1),    # right
                          value = c(100, 150, 135, 100))) # right
)

sounds = list()
for (s in 1:length(parsToTry)) {
  sounds[[length(sounds) + 1]] = do.call(soundgen,
    c(parsToTry[[s]], list(temperature = 0.001, syllLen = 500)))
}
lapply(sounds, playme)

method = c('cor', 'cosine', 'diff', 'dtw')
df = matrix(NA, nrow = length(parsToTry), ncol = length(method))
colnames(df) = method
df = as.data.frame(df)
for (i in 1:nrow(df)) {
  df[i, ] = compareSounds(
    x = spec1, # faster to calculate spec1 once
    y = sounds[[i]],
    samplingRate = 16000,
    method = method
  )[, 2]
}
df$av = rowMeans(df, na.rm = TRUE)
# row 1 = wrong pitch & formants, ..., row 4 = right pitch & formants
df$formants = c('wrong', 'wrong', 'right', 'right')
df$pitch = c('wrong', 'right', 'wrong', 'right')
df

## End(Not run)

```

crossFade

Join two waveforms by cross-fading

Description

crossFade joins two input vectors (waveforms) by cross-fading. First it truncates both input vectors, so that amp11 ends with a zero crossing and amp12 starts with a zero crossing, both on an upward portion of the soundwave. Then it cross-fades both vectors linearly with an overlap of crossLen or crossLenPoints. If the input vectors are too short for the specified length of cross-faded region, the two vectors are concatenated at zero crossings instead of cross-fading. Soundgen uses crossFade for gluing together epochs with different regimes of pitch effects (see the vignette on

sound generation), but it can also be useful for joining two separately generated sounds without audible artifacts.

Usage

```
crossFade(
  ampl1,
  ampl2,
  crossLenPoints = 240,
  crossLen = NULL,
  samplingRate = NULL,
  shape = c("lin", "exp", "log", "cos", "logistic", "gaussian")[1],
  steepness = 1
)
```

Arguments

<code>ampl1, ampl2</code>	two numeric vectors (waveforms) to be joined
<code>crossLenPoints</code>	(optional) the length of overlap in points
<code>crossLen</code>	the length of overlap in ms (overrides <code>crossLenPoints</code>)
<code>samplingRate</code>	the sampling rate of input vectors, Hz (needed only if <code>crossLen</code> is given in ms rather than points)
<code>shape</code>	controls the type of fade function: 'lin' = linear, 'exp' = exponential, 'log' = logarithmic, 'cos' = cosine, 'logistic' = logistic S-curve
<code>steepness</code>	scaling factor regulating the steepness of fading curves (except for shapes 'lin' and 'cos'): 0 = linear, >1 = steeper than default

Value

Returns a numeric vector.

See Also

[fade](#)

Examples

```
sound1 = sin(1:100 / 9)
sound2 = sin(7:107 / 3)
plot(c(sound1, sound2), type = 'b')
# an ugly discontinuity at 100 that will make an audible click

sound = crossFade(sound1, sound2, crossLenPoints = 5)
plot(sound, type = 'b') # a nice, smooth transition
length(sound) # but note that cross-fading costs us ~60 points
# because of trimming to zero crossings and then overlapping

## Not run:
# Actual sounds, alternative shapes of fade-in/out
```

```

sound3 = soundgen(formants = 'a', pitch = 200,
                  addSilence = 0, attackLen = c(50, 0))
sound4 = soundgen(formants = 'u', pitch = 200,
                  addSilence = 0, attackLen = c(0, 50))

# simple concatenation (with a click)
playme(c(sound3, sound4), 16000)

# concatenation from zc to zc (no click, but a rough transition)
playme(crossFade(sound3, sound4, crossLen = 0), 16000)

# linear crossFade over 35 ms - brief, but smooth
playme(crossFade(sound3, sound4, crossLen = 35, samplingRate = 16000), 16000)

# s-shaped cross-fade over 300 ms (shortens the sound by ~300 ms)
playme(crossFade(sound3, sound4, samplingRate = 16000,
                 crossLen = 300, shape = 'cos'), 16000)

## End(Not run)

```

defaults

Shiny app defaults

Description

A list of default values for Shiny app `soundgen_app()` - mostly the same as the defaults for `soundgen()`. NB: if defaults change, this has to be updated!!!

Usage

```
defaults
```

Format

An object of class `list` of length 69.

defaults_analyze

Defaults and ranges for analyze()

Description

A dataset containing defaults and ranges of key variables for `analyze()` and `pitch_app()`. Adjust as needed.

Usage

```
defaults_analyze
```

Format

A matrix with 58 rows and 4 columns:

default default value
low lowest permitted value
high highest permitted value
step increment for adjustment ...

defaults_analyze_pitchCand

Defaults for plotting with analyze()

Description

Default plotting settings for each pitch tracker in `analyze()` and `pitch_app()`. Adjust as needed.

Usage

`defaults_analyze_pitchCand`

Format

A dataframe with 8 rows and 5 columns:

method pitch tracking method
col color
pch point character
lwd line width
lty line type ...

detectNLP

Detect NLP

Description

(Experimental) A function for automatically detecting and annotating nonlinear vocal phenomena (NLP). Algorithm: analyze the audio using [analyze](#) and [phasegram](#), then use the extracted frame-by-frame descriptives to classify each frame as having no NLP ("none"), subharmonics ("sh"), sibebands / amplitude modulation ("sb"), or deterministic chaos ("chaos"). The classification is performed by a [naiveBayes](#) algorithm adapted to autocorrelated time series and pretrained on a manually annotated corpus of vocalizations. Whenever possible, check and correct pitch tracks prior to running the algorithm. See [naiveBayes](#) for tips on using adaptive priors and "clumping" to account for the fact that NLP typically occur in continuous segments spanning multiple frames.

Usage

```

detectNLP(
  x,
  samplingRate = NULL,
  predictors = c("nPeaks", "d2", "subDep", "amEnvDep", "entropy", "HNR", "CPP",
    "roughness"),
  thresProb = 0.4,
  unvoicedToNone = FALSE,
  train = soundgen::detectNLP_training_nonv,
  scale = NULL,
  from = NULL,
  to = NULL,
  pitchManual = NULL,
  pars_analyze = list(windowLength = 50, roughness = list(windowLength = 15, step = 3)),
  pars_phasegram = list(nonlinStats = "d2"),
  pars_naiveBayes = list(prior = "static", wlClumper = 3),
  jumpThres = 14,
  jumpWindow = 100,
  reportEvery = NULL,
  cores = 1,
  plot = FALSE,
  savePlots = NULL,
  main = NULL,
  xlab = NULL,
  ylab = NULL,
  ylim = NULL,
  width = 900,
  height = 500,
  units = "px",
  res = NA,
  ...
)

```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>predictors</code>	variables to include in NLP classification. The default is to include all 7 variables in the training corpus. NA values are fine (they do not cause the entire frame to be dropped as long as at least one variable is measured).
<code>thresProb</code>	minimum probability of NLP for the frame to be classified as non-"none", which is good for reducing false alarms ($< 1/nClasses$ means just go for the highest probability)
<code>unvoicedToNone</code>	if TRUE, frames treated as unvoiced are set to "none" (mostly makes sense with manual pitch tracking)

train	training corpus, namely the result of running naiveBayes_train on audio with known NLP episodes. Currently implemented: <code>soundgen::detectNLP_training_nonv</code> = manually annotated human nonverbal vocalizations, <code>soundgen::detectNLP_training_synth</code> = synthetic, <code>soundgen()</code> -generated sounds with various NLP. To train your own, run <code>detectNLP</code> on a collection of recordings, provide ground truth classification of NLP per frame (normally this would be converted from NLP annotations), and run naiveBayes_train .
scale	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
from, to	if NULL (default), analyzes the whole sound, otherwise from...to (s)
pitchManual	manually corrected pitch contour. For a single sound, provide a numeric vector of any length. For multiple sounds, provide a dataframe with columns "file" and "pitch" (or path to a csv file) as returned by pitch_app , ideally with the same <code>windowLength</code> and <code>step</code> as in current call to analyze. A named list with pitch vectors per file is also OK (eg as returned by <code>pitch_app</code>)
pars_analyze	arguments passed to analyze . NB: drop everything unnecessary to speed up the process, e.g. <code>nFormants = 0</code> , <code>loudness = NULL</code> , etc. If you have manual pitch contours, pass them as <code>pitchManual = . . .</code> . Make sure the "silence" threshold is appropriate, and ideally normalize the audio (silent frames are automatically assigned to "none")
pars_phasegram	arguments passed to phasegram . NB: only <code>d2</code> and <code>nPeaks</code> are used for NLP detection because they proved effective in the training corpus; other nonlinear statistics are not calculated to save time.
pars_naiveBayes	arguments passed to naiveBayes . It is strongly recommended to use some clumpering, with <code>wlClumper</code> given as frames (multiple by <code>step</code> to get the corresponding minimum duration of an NLP segment in ms), and/or dynamic priors.
jumpThres	frames in which pitch changes by <code>jumpThres</code> octaves/s more than in the surrounding frames are classified as containing "pitch jumps". Note that this is the rate of frequency change PER SECOND, not from one frame to the next
jumpWindow	the window for calculating the median pitch slope around the analyzed frame, ms
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
plot	if TRUE, produces a spectrogram with annotated NLP regimes
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
main, xlab, ylab, ...	graphical parameters passed to spectrogram
ylim	frequency range to plot, kHz (defaults to 0 to Nyquist frequency). NB: still in kHz, even if <code>yScale = bark</code> , <code>mel</code> , or <code>ERB</code>
width, height, units, res	parameters passed to png if the plot is saved

Value

Returns a dataframe with frame-by-frame descriptives, posterior probabilities of each NLP type per frame, and the tentative classification (the NLP type with the highest posterior probability, possibly corrected by clumping). The time step is equal to the larger of the steps passed to `analyze()` and `phasegram()`.

Returns a list of datasets, one per input file, with acoustic descriptives per frame (returned by `analyze` and `phasegram`), probabilities of each NLP type per frame, and the putative classification of NLP per frame.

Examples

```
## Not run:
target = soundgen(syllLen = 1600, addSilence = 0, temperature = 1e-6,
  pitch = c(380, 550, 500, 220), subDep = c(0, 0, 40, 0, 0, 0, 0, 0),
  amDep = c(0, 0, 0, 0, 80, 0, 0, 0), amFreq = 80,
  noise = c(-10, rep(-40, 5)),
  jitterDep = c(0, 0, 0, 0, 0, 3))

# classifier trained on manually annotated recordings of human nonverbal
# vocalizations
nlp = detectNLP(target, 16000, plot = TRUE, ylim = c(0, 4))

# classifier trained on synthetic, soundgen()-generated sounds
nlp = detectNLP(target, 16000, train = soundgen::detectNLP_training_synth,
  plot = TRUE, ylim = c(0, 4))
head(nlp[, c('time', 'pr')])
table(nlp$pr)
plot(nlp$amEnvDep, type = 'l')
plot(nlp$subDep, type = 'l')
plot(nlp$entropy, type = 'l')
plot(nlp$none, type = 'l')
points(nlp$sb, type = 'l', col = 'blue')
points(nlp$sh, type = 'l', col = 'green')
points(nlp$chaos, type = 'l', col = 'red')

# detection of pitch jumps
s1 = soundgen(syllLen = 1200, temperature = .001, pitch = list(
  time = c(0, 350, 351, 890, 891, 1200),
  value = c(140, 230, 460, 330, 220, 200))
playme(s1, 16000)
detectNLP(s1, 16000, plot = TRUE, ylim = c(0, 3))

# process all files in a folder
nlp = detectNLP('/home/allgoodguys/Downloads/temp260/',
  pitchManual = soundgen::pitchContour, cores = 4, plot = TRUE,
  savePlots = '', ylim = c(0, 3))

## End(Not run)
```

`detectNLP_training_nonv`

Nonlinear phenomena: Naive Bayes classifier trained on human non-verbal vocalizations

Description

The results of running `naiveBayes_train` on acoustically analyzed 969 human nonverbal vocalizations (>83K frames). It is used by `detectNLP`.

Usage`detectNLP_training_nonv`**Format**

An object of class `list` of length 9.

`detectNLP_training_synth`

Nonlinear phenomena: Naive Bayes classifier trained on synthetic sounds

Description

The results of running `naiveBayes_train` on 5000 synthetic sounds with or without NLP created with `soundgen()`. It is used by `detectNLP`.

Usage`detectNLP_training_synth`**Format**

An object of class `list` of length 9.

ERBToHz *Convert Hz to ERB rate*

Description

Converts from Hz to the number of Equivalent Rectangular Bandwidths (ERBs) below input frequency. See <https://www2.ling.su.se/staff/hartmut/bark.htm> and https://en.wikipedia.org/wiki/Equivalent_rectangular_bandwidth

Usage

```
ERBToHz(e, method = c("linear", "quadratic")[1])
```

Arguments

e vector or matrix of frequencies in ERB rate
method approximation to use

See Also

[HzToERB](#) [HzToSemitones](#) [HzToNotes](#)

Examples

```
freqs_Hz = c(-20, 20, 100, 440, 1000, 20000, NA)
e_lin = HzToERB(freqs_Hz, 'linear')
ERBToHz(e_lin, 'linear')

e_quad = HzToERB(freqs_Hz, 'quadratic')
ERBToHz(e_quad, 'quadratic')
```

estimateVTL *Estimate vocal tract length*

Description

Estimates the length of vocal tract based on formant frequencies. If method = 'meanFormant', vocal tract length (VTL) is calculated separately for each formant, and then the resulting VTLs are averaged. The equation used is $(2 * \text{formant_number} - 1) * \text{speedSound} / (4 * \text{formant_frequency})$ for a closed-open tube (mouth open) and $\text{formant_number} * \text{speedSound} / (2 * \text{formant_frequency})$ for an open-open or closed-closed tube (eg closed mouth in mmm or open mouth and open glottis in whispering). If method = 'meanDispersion', formant dispersion is calculated as the mean distance between formants, and then VTL is calculated as $\text{speedofsound} / 2 / \text{formantdispersion}$. If method = 'regression', formant dispersion is estimated using the regression method described in Reby et al. (2005) "Red deer stags use formants as assessment cues during intrasexual agonistic interactions". For a review of these and other VTL-related summary measures of formant frequencies, refer to Pisanski et al. (2014) "Vocal indicators of body size in men and women: a meta-analysis". See also [schwa](#) for VTL estimation with additional information on formant frequencies.

Usage

```
estimateVTL(
  formants,
  method = c("regression", "meanDispersion", "meanFormant")[1],
  interceptZero = TRUE,
  tube = c("closed-open", "open-open")[1],
  speedSound = 35400,
  checkFormat = TRUE,
  output = c("simple", "detailed")[1],
  plot = FALSE
)
```

Arguments

formants	formant frequencies in any format recognized by soundgen : a vector of formant frequencies like <code>c(550, 1600, 3200)</code> ; a list with multiple values per formant like <code>list(f1 = c(500, 550), f2 = 1200)</code> ; or a character string like <code>aaui</code> referring to default presets for speaker "M1" in <code>soundgen</code> presets
method	the method of estimating vocal tract length (see details)
interceptZero	if TRUE, forces the regression curve to pass through the origin. This reduces the influence of highly variable lower formants, but we have to commit to a particular model of the vocal tract: closed-open or open-open/closed-closed (method = "regression" only)
tube	the vocal tract is assumed to be a cylindrical tube that is either "closed-open" or "open-open" (same as closed-closed)
speedSound	speed of sound in warm air, by default 35400 cm/s. Stevens (2000) "Acoustic phonetics", p. 138
checkFormat	if FALSE, only a list of properly formatted formant frequencies is accepted
output	"simple" (default) = just the VTL; "detailed" = a list of additional stats (see Value below)
plot	if TRUE, plots the regression line whose slope gives formant dispersion (method = "regression" only). Label sizes show the influence of each formant, and the blue line corresponds to each formant being an integer multiple of F1 (as when harmonics are misidentified as formants); the second plot shows how VTL varies depending on the number of formants used

Value

If `output = 'simple'` (default), returns the estimated vocal tract length in cm. If `output = 'detailed'` and `method = 'regression'`, returns a list with extra stats used for plotting. Namely, `$regressionInfo$infl` gives the influence of each observation calculated as the absolute change in VTL with vs without the observation * 10 + 1 (the size of labels on the first plot). `$vtlPerFormant$vtl` gives the VTL as it would be estimated if only the first `nFormants` were used.

See Also

[schwa](#)

Examples

```

estimateVTL(NA)
estimateVTL(500)
estimateVTL(c(600, 1850, 2800, 3600, 5000), plot = TRUE)
estimateVTL(c(600, 1850, 2800, 3600, 5000), plot = TRUE, output = 'detailed')
estimateVTL(c(1200, 2000, 2800, 3800, 5400, 6400),
  tube = 'open-open', interceptZero = FALSE, plot = TRUE)
estimateVTL(c(1200, 2000, 2800, 3800, 5400, 6400),
  tube = 'open-open', interceptZero = TRUE, plot = TRUE)

# Multiple measurements are OK
estimateVTL(
  formants = list(f1 = c(540, 600, 550),
    f2 = 1650, f3 = c(2400, 2550)),
  plot = TRUE, output = 'detailed')
# NB: this is better than averaging formant values. Cf.:
estimateVTL(
  formants = list(f1 = mean(c(540, 600, 550)),
    f2 = 1650, f3 = mean(c(2400, 2550))),
  plot = TRUE)

# Missing values are OK
estimateVTL(c(600, 1850, 3100, NA, 5000), plot = TRUE)
estimateVTL(list(f1 = 500, f2 = c(1650, NA, 1400), f3 = 2700), plot = TRUE)

# Note that VTL estimates based on the commonly reported 'meanDispersion'
# depend only on the first and last formants
estimateVTL(c(500, 1400, 2800, 4100), method = 'meanDispersion')
estimateVTL(c(500, 1100, 2300, 4100), method = 'meanDispersion') # identical
# ...but this is not the case for 'meanFormant' and 'regression' methods
estimateVTL(c(500, 1400, 2800, 4100), method = 'meanFormant')
estimateVTL(c(500, 1100, 2300, 4100), method = 'meanFormant') # much longer

## Not run:
# Compare the results produced by the three methods
nIter = 1000
out = data.frame(meanFormant = rep(NA, nIter), meanDispersion = NA, regression = NA)
for (i in 1:nIter) {
  # generate a random formant configuration
  f = runif(1, 300, 900) + (1:6) * rnorm(6, 1000, 200)
  out$meanFormant[i] = estimateVTL(f, method = 'meanFormant')
  out$meanDispersion[i] = estimateVTL(f, method = 'meanDispersion')
  out$regression[i] = estimateVTL(f, method = 'regression')
}
pairs(out)
cor(out)
# 'meanDispersion' is pretty different, while 'meanFormant' and 'regression'
# give broadly comparable results

## End(Not run)

```

fade

*Fade***Description**

Applies fade-in and/or fade-out of variable length, shape, and steepness. The resulting effect softens the attack and release of a waveform.

Usage

```
fade(
  x,
  fadeIn = 50,
  fadeOut = 50,
  fadeIn_points = NULL,
  fadeOut_points = NULL,
  samplingRate = NULL,
  scale = NULL,
  shape = c("lin", "exp", "log", "cos", "logistic", "gaussian")[1],
  steepness = 1,
  reportEvery = NULL,
  cores = 1,
  saveAudio = NULL,
  plot = FALSE,
  savePlots = NULL,
  width = 900,
  height = 500,
  units = "px",
  res = NA,
  ...
)
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>fadeIn</code> , <code>fadeOut</code>	length of segments for fading in and out, ms (0 = no fade)
<code>fadeIn_points</code> , <code>fadeOut_points</code>	length of segments for fading in and out, points (if specified, override <code>fadeIn/fadeOut</code>)
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>scale</code>	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
<code>shape</code>	controls the type of fade function: 'lin' = linear, 'exp' = exponential, 'log' = logarithmic, 'cos' = cosine, 'logistic' = logistic S-curve

steepness	scaling factor regulating the steepness of fading curves (except for shapes 'lin' and 'cos'): 0 = linear, >1 = steeper than default
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
saveAudio	full path to the folder in which to save audio files (one per detected syllable)
plot	if TRUE, produces an oscillogram of the waveform after fading
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
width, height, units, res	graphical parameters for saving plots passed to png
...	other graphical parameters

Value

Returns a numeric vector of the same length as input

See Also

[crossFade](#)

Examples

```
#' # Fading a real sound: say we want fast attack and slow release
s = soundgen(attack = 0, windowLength = 10,
             syllLen = 500, addSilence = 0)
# playme(s)
s1 = fade(s, fadeIn = 40, fadeOut = 350,
          samplingRate = 16000, shape = 'cos', plot = TRUE)
# playme(s1)

# Illustration of fade shapes
x = runif(5000, min = -1, max = 1) # make sure to zero-center input!!!
# plot(x, type = 'l')
y = fade(x, fadeIn_points = 1000, fadeOut_points = 0, plot = TRUE)
y = fade(x, fadeIn_points = 1000, fadeOut_points = 1500,
         shape = 'exp', steepness = 1, plot = TRUE)
y = fade(x, fadeIn_points = 1500, fadeOut_points = 500,
         shape = 'log', steepness = 1, plot = TRUE)
y = fade(x, fadeIn_points = 1500, fadeOut_points = 500,
         shape = 'log', steepness = 3, plot = TRUE)
y = fade(x, fadeIn_points = 1500, fadeOut_points = 1500,
         shape = 'cos', plot = TRUE)
y = fade(x, fadeIn_points = 1500, fadeOut_points = 1500,
         shape = 'logistic', steepness = 1, plot = TRUE)
y = fade(x, fadeIn_points = 1500, fadeOut_points = 1500,
         shape = 'logistic', steepness = 3, plot = TRUE)
y = fade(x, fadeIn_points = 1500, fadeOut_points = 1500,
         shape = 'gaussian', steepness = 1.5, plot = TRUE)
```

```
## Not run:
fade('~Downloads/temp', fadeIn = 500, fadeOut = 500, savePlots = '')

## End(Not run)
```

fart

Fart

Description

While the same sounds can be created with `soundgen()`, this facetious function produces the same effect more efficiently and with very few control parameters. With default settings, execution time is ~ 10 ms per second of audio sampled at 16000 Hz. Principle: creates separate glottal cycles with harmonics, but no formants. See [soundgen](#) for more details.

Usage

```
fart(
  glottis = c(50, 200),
  pitch = 65,
  temperature = 0.25,
  syllLen = 600,
  rolloff = -10,
  samplingRate = 16000,
  play = FALSE,
  plot = FALSE
)
```

Arguments

<code>glottis</code>	anchors for specifying the proportion of a glottal cycle with closed glottis, % (0 = no modification, 100 = closed phase as long as open phase); numeric vector or dataframe specifying time and value (anchor format)
<code>pitch</code>	a numeric vector of f_0 values in Hz or a dataframe specifying the time (ms or 0 to 1) and value (Hz) of each anchor, hereafter "anchor format". These anchors are used to create a smooth contour of fundamental frequency f_0 (pitch) within one syllable
<code>temperature</code>	hyperparameter for regulating the amount of stochasticity in sound generation
<code>syllLen</code>	syllable length, ms (not vectorized)
<code>rolloff</code>	rolloff of harmonics in source spectrum, dB/octave (not vectorized)
<code>samplingRate</code>	sampling frequency, Hz
<code>play</code>	if TRUE, plays the synthesized sound using the default player on your system. If character, passed to play as the name of player to use, eg "aplay", "play", "vlc", etc. In case of errors, try setting another default player for play
<code>plot</code>	if TRUE, plots the waveform

Value

Returns a normalized waveform.

See Also

[soundgen](#) [generateNoise](#) [beat](#)

Examples

```
f = fart()
# playme(f)

## Not run:
while (TRUE) {
  fart(syllLen = 300, temperature = .5, play = TRUE)
  Sys.sleep(rexp(1, rate = 1))
}

## End(Not run)
```

filterMS

Filter modulation spectrum

Description

Filters a modulation spectrum by removing a certain range of amplitude modulation (AM) and frequency modulation (FM) frequencies. Conditions can be specified either separately for AM and FM with `amCond = ...`, `fmCond = ...`, implying an OR combination of conditions, or jointly on AM and FM with `jointCond`. `jointCond` is more general, but using `amCond/fmCond` is ~100 times faster.

Usage

```
filterMS(
  ms,
  amCond = NULL,
  fmCond = NULL,
  jointCond = NULL,
  action = c("remove", "preserve")[1],
  plot = TRUE
)
```

Arguments

`ms` a modulation spectrum as returned by [modulationSpectrum](#) - a matrix of real or complex values, AM in columns, FM in rows

`amCond`, `fmCond` character strings with valid conditions on amplitude and frequency modulation (see examples)

jointCond	character string with a valid joint condition amplitude and frequency modulation
action	should the defined AM-FM region be removed ('remove') or preserved, while everything else is removed ('preserve')?
plot	if TRUE, plots the filtered modulation spectrum

Value

Returns the filtered modulation spectrum - a matrix of the original dimensions, real or complex.

Examples

```
ms = modulationSpectrum(soundgen(), samplingRate = 16000,
                        returnComplex = TRUE)$complex
# Remove all AM over 25 Hz
ms_filt = filterMS(ms, amCond = 'abs(am) > 25')

# amCond and fmCond are OR-conditions
filterMS(ms, amCond = 'abs(am) > 15', fmCond = 'abs(fm) > 5', action = 'remove')
filterMS(ms, amCond = 'abs(am) > 15', fmCond = 'abs(fm) > 5', action = 'preserve')
filterMS(ms, amCond = 'abs(am) > 10 & abs(am) < 25', action = 'remove')

# jointCond is an AND-condition
filterMS(ms, jointCond = 'am * fm < 5', action = 'remove')
filterMS(ms, jointCond = 'am^2 + (fm*3)^2 < 200', action = 'preserve')

# So:
filterMS(ms, jointCond = 'abs(am) > 5 | abs(fm) < 5') # slow but general
# ...is the same as:
filterMS(ms, amCond = 'abs(am) > 5', fmCond = 'abs(fm) < 5') # fast
```

filterSoundByMS

Filter sound by modulation spectrum

Description

Manipulates the modulation spectrum (MS) of a sound so as to remove certain frequencies of amplitude modulation (AM) and frequency modulation (FM). Algorithm: produces a modulation spectrum with `modulationSpectrum`, modifies it with `filterMS`, converts the modified MS to a spectrogram with `msToSpec`, and finally inverts the spectrogram with `invertSpectrogram`, thus producing a sound with (approximately) the desired characteristics of the MS. Note that the last step of inverting the spectrogram introduces some noise, so the resulting MS is not precisely the same as the intermediate filtered version. In practice this means that some residual energy will still be present in the filtered-out frequency range (see examples).

Usage

```

filterSoundByMS(
  x,
  samplingRate = NULL,
  from = NULL,
  to = NULL,
  logSpec = FALSE,
  windowLength = 25,
  step = NULL,
  overlap = 80,
  wn = "hamming",
  zp = 0,
  amCond = NULL,
  fmCond = NULL,
  jointCond = NULL,
  action = c("remove", "preserve")[1],
  initialPhase = c("zero", "random", "spssi")[3],
  nIter = 50,
  reportEvery = NULL,
  cores = 1,
  play = FALSE,
  saveAudio = NULL,
  plot = TRUE,
  savePlots = NULL,
  width = 900,
  height = 500,
  units = "px",
  res = NA
)

```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>from</code> , <code>to</code>	if <code>NULL</code> (default), analyzes the whole sound, otherwise from...to (s)
<code>logSpec</code>	if <code>TRUE</code> , the spectrogram is log-transformed prior to taking 2D FFT
<code>windowLength</code>	length of FFT window, ms
<code>step</code>	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
<code>overlap</code>	overlap between successive FFT frames, %
<code>wn</code>	window type accepted by <code>ftwindow</code> , currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop
<code>zp</code>	window length after zero padding, points

amCond, fmCond	character strings with valid conditions on amplitude and frequency modulation (see examples)
jointCond	character string with a valid joint condition amplitude and frequency modulation
action	should the defined AM-FM region be removed ('remove') or preserved, while everything else is removed ('preserve')?
initialPhase	initial phase estimate: "zero" = set all phases to zero; "random" = Gaussian noise; "spsi" (default) = single-pass spectrogram inversion (Beauregard et al., 2015)
nIter	the number of iterations of the GL algorithm (Griffin & Lim, 1984), 0 = don't run
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
play	if TRUE, plays back the reconstructed audio
saveAudio	full (!) path to folder for saving the processed audio; NULL = don't save, "" = same as input folder (NB: overwrites the originals!)
plot	if TRUE, produces a triple plot: original MS, filtered MS, and the MS of the output sound
savePlots	if a valid path is specified, a plot is saved in this folder (defaults to NA)
width, height, units, res	parameters passed to png if the plot is saved

Value

Returns the filtered audio as a numeric vector normalized to [-1, 1] with the same sampling rate as input.

See Also

[invertSpectrogram](#) [filterMS](#)

Examples

```
# Create a sound to be filtered
s = soundgen(pitch = rnorm(n = 20, mean = 200, sd = 25),
  amFreq = 25, amDep = 50, samplingRate = 16000,
  addSilence = 50, plot = TRUE, osc = TRUE)
# playme(s, 16000)

# Filter
s_filt = filterSoundByMS(s, samplingRate = 16000,
  amCond = 'abs(am) > 15', fmCond = 'abs(fm) > 5',
  action = 'remove', nIter = 10, plot = TRUE)
# playme(s_filt, samplingRate = 16000)

## Not run:
# Process all files in a folder, save filtered audio and plots
```

```

s_filt = filterSoundByMS('~Downloads/temp2',
  saveAudio = '~Downloads/temp2/ms', savePlots = '',
  amCond = 'abs(am) > 15', fmCond = 'abs(fm) > 5',
  action = 'remove', nIter = 10)

# Download an example - a bit of speech (sampled at 16000 Hz)
download.file('http://cogsci.se/soundgen/audio/speechEx.wav',
  destfile = '~Downloads/speechEx.wav') # modify as needed
target = '~Downloads/speechEx.wav'
samplingRate = tuneR::readWave(target)@samp.rate
playme(target)
spectrogram(target, osc = TRUE)

# Remove AM above 3 Hz from a bit of speech (remove most temporal details)
s_filt1 = filterSoundByMS(target, amCond = 'abs(am) > 3',
  action = 'remove', nIter = 15)
playme(s_filt1, samplingRate)
spectrogram(s_filt1, samplingRate = samplingRate, osc = TRUE)

# Intelligible when AM in 5-25 Hz is preserved:
s_filt2 = filterSoundByMS(target, amCond = 'abs(am) > 5 & abs(am) < 25',
  action = 'preserve', nIter = 15)
playme(s_filt2, samplingRate)
spectrogram(s_filt2, samplingRate = samplingRate, osc = TRUE)

# Remove slow AM/FM (prosody) to achieve a "robotic" voice
s_filt3 = filterSoundByMS(target, jointCond = 'am^2 + (fm*3)^2 < 300',
  nIter = 15)
playme(s_filt3, samplingRate)
spectrogram(s_filt3, samplingRate = samplingRate, osc = TRUE)

## An alternative manual workflow w/o calling filterSoundByMS()
# This way you can modify the MS directly and more flexibly
# than with the filterMS() function called by filterSoundByMS()

# (optional) Check that the target spectrogram can be successfully inverted
spec = spectrogram(s, 16000, windowLength = 25, overlap = 80,
  wn = 'hanning', osc = TRUE, padWithSilence = FALSE)
s_rev = invertSpectrogram(spec, samplingRate = 16000,
  windowLength = 25, overlap = 80, wn = 'hamming', play = FALSE)
# playme(s_rev, 16000) # should be close to the original
spectrogram(s_rev, 16000, osc = TRUE)

# Get modulation spectrum starting from the sound...
ms = modulationSpectrum(s, samplingRate = 16000, windowLength = 25,
  overlap = 80, wn = 'hanning', amRes = NULL, maxDur = Inf, logSpec = FALSE,
  power = NA, returnComplex = TRUE, plot = FALSE)$complex
# ... or starting from the spectrogram:
# ms = specToMS(spec)
image(x = as.numeric(colnames(ms)), y = as.numeric(rownames(ms)),
  z = t(log(abs(ms)))) # this is the original MS

```

```

# Filter as needed - for ex., remove AM > 10 Hz and FM > 3 cycles/kHz
# (removes f0, preserves formants)
am = as.numeric(colnames(ms))
fm = as.numeric(rownames(ms))
idx_row = which(abs(fm) > 3)
idx_col = which(abs(am) > 10)
ms_filt = ms
ms_filt[idx_row, ] = 0
ms_filt[, idx_col] = 0
image(x = as.numeric(colnames(ms_filt)), y = as.numeric(rownames(ms_filt)),
      t(log(abs(ms_filt)))) # this is the filtered MS

# Convert back to a spectrogram
spec_filt = msToSpec(ms_filt)
image(t(log(abs(spec_filt))))

# Invert the spectrogram
s_filt = invertSpectrogram(abs(spec_filt), samplingRate = 16000,
  windowLength = 25, overlap = 80, wn = 'hanning')
# NB: use the same settings as in "spec = spectrogram(s, ...)" above

# Compare with the original
playme(s, 16000)
spectrogram(s, 16000, osc = TRUE)
playme(s_filt, 16000)
spectrogram(s_filt, 16000, osc = TRUE)

ms_new = modulationSpectrum(s_filt, samplingRate = 16000,
  windowLength = 25, overlap = 80, wn = 'hanning', maxDur = Inf,
  plot = TRUE, returnComplex = TRUE)$complex
image(x = as.numeric(colnames(ms_new)), y = as.numeric(rownames(ms_new)),
      z = t(log(abs(ms_new))))
plot(as.numeric(colnames(ms)), log(abs(ms[nrow(ms) / 2, ])), type = 'l')
points(as.numeric(colnames(ms_new)), log(ms_new[nrow(ms_new) / 2, ]), type = 'l',
  col = 'red', lty = 3)
# AM peaks at 25 Hz are removed, but inverting the spectrogram adds a lot of noise

## End(Not run)

```

findInflections

Find inflections

Description

Finds inflections in discrete time series such as pitch contours. When there are no missing values and no thresholds, this can be accomplished with a fast one-liner like `which(diff(diff(x) > 0) != 0) + 1`. Missing values are interpolated by repeating the first and last non-missing values at the head and tail, respectively, and by linear interpolation in the middle. Setting a threshold means that small "wiggling" no longer counts. To use an analogy with ocean waves, smoothing (low-pass filtering) removes the ripples and only leaves the slow roll, while thresholding preserves only waves that are sufficiently high, whatever their period.

Usage

```
findInflections(x, thres = NULL, step = NULL, plot = FALSE, main = "")
```

Arguments

x	numeric vector with or without NAs
thres	minimum vertical distance between two extrema for them to count as two independent inflections
step	distance between values in s (only needed for plotting)
plot	if TRUE, produces a simple plot
main	plot title

Value

Returns a vector of indices giving the location of inflections.

Examples

```
x = sin(2 * pi * (1:100) / 15) * seq(1, 5, length.out = 100)
idx_na = c(1:4, 6, 7, 14, 25, 30:36, 39, 40, 42, 45:50,
          57, 59, 62, 66, 71:79, 98)
x[idx_na] = NA
soundgen:::findInflections(x, plot = TRUE)
soundgen:::findInflections(x, thres = 5, plot = TRUE)

for (i in 1:10) {
  temp = soundgen:::getRandomWalk(len = runif(1, 10, 100), rw_range = 10,
                                rw_smoothing = runif(1, 0, 1))
  soundgen:::findInflections(temp, thres = 1, plot = TRUE)
  invisible(readline(prompt="Press [enter] to continue"))
}
```

 findJumps

Find frequency jumps

Description

This function flags frames with apparent pitch jumps (frequency jumps, voice breaks), defined as relatively large and sudden changes in voice pitch or some other frequency measure (peak frequency, a formant frequency, etc). It is called by `detectNLP`. Algorithm: a frame is considered to contain a frequency jump if the absolute slope at this frame exceeds the average slope over \pm jumpWindow around it by more than jumpThres. Note that the slope is considered per second rather than per time step - that is, taking into account the sampling rate of the frequency track. Thus, it's not just the change from frame to frame that defines what is considered a jump, but a change that differs from the trend in the surrounding frames (see examples). If several consecutive frames contain apparent jumps, only the greatest of them is preserved.

Usage

```
findJumps(
  pitch,
  step,
  jumpThres = 8,
  jumpWindow = 80,
  plot = FALSE,
  xlab = "Time, ms",
  ylab = "f0, Hz",
  ...
)
```

Arguments

pitch	vector of frequencies per frame, Hz
step	time step between frames, ms
jumpThres	frames in which pitch changes by jumpThres octaves/s more than in the surrounding frames are classified as containing "pitch jumps". Note that this is the rate of frequency change PER SECOND, not from one frame to the next
jumpWindow	the window for calculating the median pitch slope around the analyzed frame, ms
plot	if TRUE, plots the pitch contour with putative frequency jumps marked by arrows
xlab, ylab, ...	graphical parameters passed to plot

Value

Returns a boolean vector of the same length as pitch, where TRUE values correspond to frames with detected pitch jumps.

Examples

```
pitch = getSmoothContour(anchors = list(
  time = c(0, 350, 351, 890, 891, 1200),
  value = c(140, 230, 460, 330, 220, 200)), len = 40)
step = 25
pj = findJumps(pitch, step, plot = TRUE)

# convert frame indices to time in ms
step = 25
which(pj) * step
# or consider pj's to occur midway between the two frames
which(pj) * step - step / 2

# even very rapid changes are not considered jumps if they match
# the surrounding trend
pitch = getSmoothContour(anchors = list(
  time = c(0, 350, 351, 700),
```

```

value = c(340, 710, 850, 1200)), len = 20)
findJumps(pitch, step, plot = TRUE)
diff(HzToSemitones(pitch)) * (1000 / step) / 12
# the slope at frame 10 (10.4 oct/s) exceeds the jumpThres (8 oct/s), but not
# 10.4 minus the average slope around frame 10 (~3 oct/s, so 10 - 3 < 8)

```

flatEnv

Flat envelope / compressor

Description

Applies a compressor - that is, flattens the amplitude envelope of a waveform, reducing the difference in amplitude between loud and quiet sections. This is achieved by dividing the waveform by some function of its smoothed amplitude envelope (Hilbert, peak or root mean square).

Usage

```

flatEnv(
  x,
  samplingRate = NULL,
  scale = NULL,
  compression = 1,
  method = c("hil", "rms", "peak")[1],
  windowLength = 50,
  windowLength_points = NULL,
  killDC = FALSE,
  dynamicRange = 40,
  reportEvery = NULL,
  cores = 1,
  saveAudio = NULL,
  plot = FALSE,
  savePlots = NULL,
  col = "blue",
  width = 900,
  height = 500,
  units = "px",
  res = NA,
  ...
)

compressor(
  x,
  samplingRate = NULL,
  scale = NULL,
  compression = 1,
  method = c("hil", "rms", "peak")[1],
  windowLength = 50,

```

```

    windowLength_points = NULL,
    killDC = FALSE,
    dynamicRange = 40,
    reportEvery = NULL,
    cores = 1,
    saveAudio = NULL,
    plot = FALSE,
    savePlots = NULL,
    col = "blue",
    width = 900,
    height = 500,
    units = "px",
    res = NA,
    ...
)

```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
scale	maximum possible amplitude of input used for normalization of input vector (only needed if x is a numeric vector)
compression	the amount of compression to apply: 0 = none, 1 = maximum
method	hil = Hilbert envelope, rms = root mean square amplitude, peak = peak amplitude per window
windowLength	the length of smoothing window, ms
windowLength_points	the length of smoothing window, points. If specified, overrides windowLength
killDC	if TRUE, dynamically removes DC offset or similar deviations of average waveform from zero (see examples)
dynamicRange	parts of sound quieter than -dynamicRange dB will not be amplified
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
saveAudio	full path to the folder in which to save the compressed sound(s)
plot	if TRUE, plots the original sound, the smoothed envelope, and the compressed sound
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
col	the color of amplitude contours
width, height, units, res	graphical parameters for saving plots passed to png
...	other graphical parameters passed to <code>points()</code> that control the appearance of amplitude contours, eg <code>lwd</code> , <code>lty</code> , etc.

Value

If the input is a single audio (file, Wave, or numeric vector), returns the compressed waveform as a numeric vector with the original sampling rate and scale. If the input is a folder with several audio files, returns a list of compressed waveforms, one for each file.

Examples

```
a = rnorm(500) * seq(1, 0, length.out = 500)
b = flatEnv(a, 1000, plot = TRUE, windowLength_points = 5) # too short
c = flatEnv(a, 1000, plot = TRUE, windowLength_points = 450) # too long
d = flatEnv(a, 1000, plot = TRUE, windowLength_points = 100) # about right

## Not run:
s = soundgen(syllLen = 1000, ampl = c(0, -40, 0), plot = TRUE)
# playme(s)
s_flat1 = flatEnv(s, 16000, dynamicRange = 60, plot = TRUE,
                 windowLength = 50, method = 'hil')
s_flat2 = flatEnv(s, 16000, dynamicRange = 60, plot = TRUE,
                 windowLength = 10, method = 'rms')
s_flat3 = flatEnv(s, 16000, dynamicRange = 60, plot = TRUE,
                 windowLength = 10, method = 'peak')
# playme(s_flat2)

# Remove DC offset
s1 = c(rep(0, 50), runif(1000, -1, 1), rep(0, 50)) +
     seq(.3, 1, length.out = 1100)
s2 = flatEnv(s1, 16000, plot = TRUE, windowLength_points = 50, killDC = FALSE)
s3 = flatEnv(s1, 16000, plot = TRUE, windowLength_points = 50, killDC = TRUE)

# Compress and save all audio files in a folder
s4 = flatEnv('~Downloads/temp2',
            method = 'peak', compression = .5,
            saveAudio = '~Downloads/temp2/compressed',
            savePlots = '~Downloads/temp2/compressed',
            col = 'green', lwd = 5)
osc(s4[[1]])

## End(Not run)
```

flatSpectrum

Flat spectrum

Description

Flattens the spectrum of a sound by smoothing in the frequency domain. Can be used for removing formants without modifying pitch contour or voice quality (the balance of harmonic and noise components), followed by the addition of a new spectral envelope (cf. [transplantFormants](#)). Algorithm: makes a spectrogram, flattens the real part of the smoothed spectrum of each STFT frame, and transforms back into time domain with inverse STFT (see also [addFormants](#)).

Usage

```
flatSpectrum(
  x,
  samplingRate = NULL,
  freqWindow = NULL,
  dynamicRange = 80,
  windowLength = 50,
  step = NULL,
  overlap = 90,
  wn = "gaussian",
  zp = 0,
  play = FALSE,
  saveAudio = NULL,
  reportEvery = NULL,
  cores = 1
)
```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
freqWindow	the width of smoothing window, Hz. Defaults to median pitch estimated by analyze
dynamicRange	dynamic range, dB. All values more than one dynamicRange under maximum are treated as zero
windowLength	length of FFT window, ms
step	you can override overlap by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of 1/samplingRate, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
overlap	overlap between successive FFT frames, %
wn	window type accepted by ftwindow , currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop
zp	window length after zero padding, points
play	if TRUE, plays the processed audio
saveAudio	full (!) path to folder for saving the processed audio; NULL = don't save, "" = same as input folder (NB: overwrites the originals!)
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing

Value

Returns a numeric vector with the same sampling rate as the input.

See Also

[addFormants](#) [transplantFormants](#)

Examples

```

sound_aii = soundgen(formants = 'aai')
# playme(sound_aai, 16000)
seewave::meanspec(sound_aai, f = 16000, dB = 'max0')

sound_flat = flatSpectrum(sound_aai, freqWindow = 150, samplingRate = 16000)
# playme(sound_flat, 16000)
seewave::meanspec(sound_flat, f = 16000, dB = 'max0')
# harmonics are still there, but formants are gone and can be replaced

## Not run:
# Now let's make a sheep say "aai"
data(sheep, package = 'seewave') # import a recording from seewave
playme(sheep)
sheep_flat = flatSpectrum(sheep)
playme(sheep_flat, sheep@samp.rate)
seewave::spec(sheep_flat, f = sheep@samp.rate, dB = 'max0')

# So far we have a sheep bleating with a flat spectrum;
# now let's add new formants
sheep_aai = addFormants(sheep_flat,
  samplingRate = sheep@samp.rate,
  formants = 'aai',
  lipRad = -3) # negative lipRad to counter unnatural flat source
playme(sheep_aai, sheep@samp.rate)
spectrogram(sheep_aai, sheep@samp.rate)
seewave::spec(sheep_aai, f = sheep@samp.rate, dB = 'max0')

## End(Not run)

```

formant_app

Interactive formant tracker

Description

Starts a shiny app for manually correcting formant measurements. For more tips, see [pitch_app](#) and <http://cogsci.se/soundgen.html>.

Usage

```
formant_app()
```

Details

Suggested workflow: load one or several audio files (wav/mp3), preferably not longer than a minute or so. Select a region of interest in the spectrogram - for example, a sustained vowel with clear and relatively steady formants. Double-click within the selection to create a new annotation (you may add a text label if needed). If you are satisfied with the automatically calculated formant frequencies, proceed to the next region of interest. If not, there are 4 ways to adjust them: (1) type in the correct number in one of the formant boxes in the top right corner; (2) click a spectrogram within selection (pick the formant number to adjust by clicking the formant boxes); (3) single-click the spectrum to use the cursor's position, or (4) double-click the spectrum to use the nearest spectral peak. When done with a file, move on to the next one in the queue. Use the orange button to download the results. To continue work, upload the output file from the previous session together with the audio files (you can rename it, but keep the .csv extension). Use hotkeys (eg spacebar to play/stop) and avoid working with very large files.

Value

Every time a new annotation is added, the app creates a backup csv file and creates or updates a global object called "my_formants", which contains all the annotations. When the app is terminated, it also returns the results as a dataframe.

See Also

[pitch_app](#)

Examples

```
## Not run:
f = formant_app() # runs in default browser such as Firefox or Chrome

# To change system default browser, run something like:
options('browser' = '/usr/bin/firefox') # path to the executable on Linux

## End(Not run)
```

gaussianSmooth2D

Gaussian smoothing in 2D

Description

Takes a matrix of numeric values and smoothes it by convolution with a symmetric Gaussian window function.

Usage

```
gaussianSmooth2D(
  m,
  kernelSize = 5,
  kernelSD = 0.5,
```

```

    action = c("blur", "unblur")[1],
    plotKernel = FALSE
  )

```

Arguments

<code>m</code>	input matrix (numeric, on any scale, doesn't have to be square)
<code>kernelSize</code>	the size of the Gaussian kernel, in points
<code>kernelSD</code>	the SD of the Gaussian kernel relative to its size (.5 = the edge is two SD's away)
<code>action</code>	'blur' = kernel-weighted average, 'unblur' = subtract kernel-weighted average
<code>plotKernel</code>	if TRUE, plots the kernel

Value

Returns a numeric matrix of the same dimensions as input.

See Also

[modulationSpectrum](#)

Examples

```

s = spectrogram(soundgen(), samplingRate = 16000, windowLength = 10,
  output = 'original', plot = FALSE)
s = log(s + .001)
# image(s)
s1 = gaussianSmooth2D(s, kernelSize = 5, plotKernel = TRUE)
# image(s1)

## Not run:
# more smoothing in time than in frequency
s2 = gaussianSmooth2D(s, kernelSize = c(5, 15))
image(s2)

# vice versa - more smoothing in frequency
s3 = gaussianSmooth2D(s, kernelSize = c(25, 3))
image(s3)

# sharpen the image by deconvolution with the kernel
s4 = gaussianSmooth2D(s1, kernelSize = 5, action = 'unblur')
image(s4)

s5 = gaussianSmooth2D(s, kernelSize = c(15, 1), action = 'unblur')
image(s5)

## End(Not run)

```

generateNoise	<i>Generate noise</i>
---------------	-----------------------

Description

Generates noise of length `len` and with spectrum defined by rolloff parameters OR by a specified filter `spectralEnvelope`. This function is called internally by `soundgen`, but it may be more convenient to call it directly when synthesizing non-biological noises defined by specific spectral and amplitude envelopes rather than formants: the wind, whistles, impact noises, etc. See `fart` and `beat` for similarly simplified functions for tonal non-biological sounds.

Usage

```
generateNoise(
  len,
  rolloffNoise = 0,
  noiseFlatSpec = 1200,
  rolloffNoiseExp = 0,
  spectralEnvelope = NULL,
  noise = NULL,
  temperature = 0.1,
  attackLen = 10,
  windowLength_points = 1024,
  samplingRate = 16000,
  overlap = 75,
  dynamicRange = 80,
  smoothing = list(),
  invalidArgAction = c("adjust", "abort", "ignore")[1],
  play = FALSE
)
```

Arguments

<code>len</code>	length of output
<code>rolloffNoise</code> , <code>noiseFlatSpec</code>	linear rolloff of the excitation source for the unvoiced component, <code>rolloffNoise</code> dB/kHz (anchor format) applied above <code>noiseFlatSpec</code> Hz
<code>rolloffNoiseExp</code>	exponential rolloff of the excitation source for the unvoiced component, dB/oct (anchor format) applied above 0 Hz
<code>spectralEnvelope</code>	(optional): as an alternative to using <code>rolloffNoise</code> , we can provide the exact filter - a vector of non-negative numbers specifying the desired spectrum on a linear scale up to Nyquist frequency (<code>samplingRate / 2</code>). The length doesn't matter as it can be interpolated internally to <code>windowLength_points/2</code> . A matrix specifying the filter for each STFT step is also accepted. The easiest way to

	obtain spectralEnvelope is to call <code>soundgen::getSpectralEnvelope</code> or to use the spectrum / spectrogram of a recorded sound
noise	loudness of turbulent noise (0 dB = as loud as voiced component, negative values = quieter) such as aspiration, hissing, etc (anchor format)
temperature	hyperparameter for regulating the amount of stochasticity in sound generation
attackLen	duration of fade-in / fade-out at each end of syllables and noise (ms): a vector of length 1 (symmetric) or 2 (separately for fade-in and fade-out)
windowLength_points	the length of fft window, points
samplingRate	sampling frequency, Hz
overlap	FFT window overlap, %. For allowed values, see istft
dynamicRange	dynamic range, dB. Harmonics and noise more than dynamicRange under maximum amplitude are discarded to save computational resources
smoothing	a list of parameters passed to getSmoothContour to control the interpolation and smoothing of contours: <code>interpol</code> (approx / spline / loess), <code>loessSpan</code> , <code>discontThres</code> , <code>jumpThres</code>
invalidArgAction	what to do if an argument is invalid or outside the range in <code>permittedValues</code> : 'adjust' = reset to default value, 'abort' = stop execution, 'ignore' = throw a warning and continue (may crash)
play	if TRUE, plays the synthesized sound using the default player on your system. If character, passed to play as the name of player to use, eg "aplay", "play", "vlc", etc. In case of errors, try setting another default player for play

Details

Algorithm: paints a spectrogram with desired characteristics, sets phase to zero, and generates a time sequence via inverse FFT.

See Also

[soundgen fart beat](#)

Examples

```
# .5 s of white noise
samplingRate = 16000
noise1 = generateNoise(len = samplingRate * .5,
  samplingRate = samplingRate)
# playme(noise1, samplingRate)
# seewave::meanspec(noise1, f = samplingRate)

# Percussion (run a few times to notice stochasticity due to temperature = .25)
noise2 = generateNoise(len = samplingRate * .15, noise = c(0, -80),
  rolloffNoise = c(4, -6), attackLen = 5, temperature = .25)
noise3 = generateNoise(len = samplingRate * .25, noise = c(0, -40),
  rolloffNoise = c(4, -20), attackLen = 5, temperature = .25)
```

```

# playme(c(noise2, noise3), samplingRate)

## Not run:
playback = list(TRUE, FALSE, 'aplay', 'vlc')[[1]]
# 1.2 s of noise with rolloff changing from 0 to -12 dB above 2 kHz
noise = generateNoise(len = samplingRate * 1.2,
  rolloffNoise = c(0, -12), noiseFlatSpec = 2000,
  samplingRate = samplingRate, play = playback)
# spectrogram(noise, samplingRate, osc = TRUE)

# Similar, but using the dataframe format to specify a more complicated
# contour for rolloffNoise:
noise = generateNoise(len = samplingRate * 1.2,
  rolloffNoise = data.frame(time = c(0, .3, 1), value = c(-12, 0, -12)),
  noiseFlatSpec = 2000, samplingRate = samplingRate, play = playback)
# spectrogram(noise, samplingRate, osc = TRUE)

# To create a sibilant [s], specify a single strong, broad formant at ~7 kHz:
windowLength_points = 1024
spectralEnvelope = soundgen::getSpectralEnvelope(
  nr = windowLength_points / 2, nc = 1, samplingRate = samplingRate,
  formants = list('f1' = data.frame(time = 0, freq = 7000,
    amp = 50, width = 2000)))
noise = generateNoise(len = samplingRate,
  samplingRate = samplingRate, spectralEnvelope = as.numeric(spectralEnvelope),
  play = playback)
# plot(spectralEnvelope, type = 'l')

# Low-frequency, wind-like noise
spectralEnvelope = soundgen::getSpectralEnvelope(
  nr = windowLength_points / 2, nc = 1, lipRad = 0,
  samplingRate = samplingRate, formants = list('f1' = data.frame(
    time = 0, freq = 150, amp = 30, width = 90)))
noise = generateNoise(len = samplingRate,
  samplingRate = samplingRate, spectralEnvelope = as.numeric(spectralEnvelope),
  play = playback)

# Manual filter, e.g. for a kettle-like whistle (narrow-band noise)
spectralEnvelope = c(rep(0, 100), 120, rep(0, 100)) # any length is fine
# plot(spectralEnvelope, type = 'b') # notch filter at Nyquist / 2, here 4 kHz
noise = generateNoise(len = samplingRate, spectralEnvelope = spectralEnvelope,
  samplingRate = samplingRate, play = playback)

# Compare to a similar sound created with soundgen()
# (unvoiced only, a single formant at 4 kHz)
noise_s = soundgen(pitch = NULL,
  noise = data.frame(time = c(0, 1000), value = c(0, 0)),
  formants = list(f1 = data.frame(freq = 4000, amp = 80, width = 20)),
  play = playback)

# Use the spectral envelope of an existing recording (bleating of a sheep)
# (see also the same example with tonal source in ?addFormants)

```

```

data(sheep, package = 'seewave') # import a recording from seewave
sound_orig = as.numeric(sheep@left)
samplingRate = sheep@samp.rate
# playme(sound_orig, samplingRate)

# extract the original spectrogram
windowLength = c(5, 10, 50, 100)[1] # try both narrow-band (eg 100 ms)
# to get "harmonics" and wide-band (5 ms) to get only formants
spectralEnvelope = spectrogram(sound_orig, windowLength = windowLength,
  samplingRate = samplingRate, output = 'original', padWithSilence = FALSE)
sound_noise = generateNoise(len = length(sound_orig),
  spectralEnvelope = spectralEnvelope, rolloffNoise = 0,
  samplingRate = samplingRate, play = playback)
# playme(sound_noise, samplingRate)

# The spectral envelope is similar to the original recording. Compare:
par(mfrow = c(1, 2))
seewave::meanspec(sound_orig, f = samplingRate, dB = 'max0')
seewave::meanspec(sound_noise, f = samplingRate, dB = 'max0')
par(mfrow = c(1, 1))
# However, the excitation source is now white noise
# (which sounds like noise if windowLength is ~5-10 ms,
# but becomes more and more like the original at longer window lengths)

## End(Not run)

```

getDuration

Get duration

Description

Returns the duration of one or more audio files (mostly useful for running on an entire folder). If threshold is set, it also removes the leading and trailing silences or near-silences, thus returning the duration of relatively loud central fragments of each sound. Silences are located based on the amplitude of root mean square (RMS) amplitude with [getRMS](#). Note that the threshold is set relative to the observed maximum RMS, just as in [analyze](#). This means that even very quiet sounds are not treated as nothing but silence.

Usage

```

getDuration(
  x,
  samplingRate = NULL,
  silence = 0.01,
  rms = list(windowLength = 20, step = 5),
  reportEvery = NULL,
  cores = 1
)

```


Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
silence	leading and trailing sections quieter than this proportion of maximum RMS amplitude are removed when calculating duration_noSilence (NULL = don't calculate duration_noSilence to save time)
rms	a list of control parameters passed to getRMS
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing

Value

Returns duration (s) and duration_noSilence (duration without leading and trailing silences).

See Also

[analyze](#) [getLoudness](#)

Examples

```
s = c(rep(0, 550), runif(400, -1, 1), rep(0, 50))
osc(s, samplingRate = 1000)
# true duration_noSilence is 400 ms
getDuration(s, samplingRate = 1000, silence = .01)
getDuration(s, samplingRate = 1000, silence = .1,
            rms = list(windowLength = 5, step = 1))

## Not run:
d = getDuration('~Downloads/temp')
hist(d$duration - d$duration_noSilence)

## End(Not run)
```

getEntropy

Entropy

Description

Returns Weiner or Shannon entropy of an input vector such as the spectrum of a sound. Non-positive input values are converted to a small positive number (`convertNonPositive`). If all elements are zero, returns NA.

Usage

```
getEntropy(
  x,
  type = c("weiner", "shannon")[1],
  normalize = FALSE,
  convertNonPositive = 1e-10
)
```

Arguments

x vector of positive floats

type 'shannon' for Shannon (information) entropy, 'weiner' for Weiner entropy

normalize if TRUE, Shannon entropy is normalized by the length of input vector to range from 0 to 1. It has no affect on Weiner entropy

convertNonPositive all non-positive values are converted to convertNonPositive

Examples

```
# Here are four simplified power spectra, each with 9 frequency bins:
s = list(
  c(rep(0, 4), 1, rep(0, 4)),      # a single peak in spectrum
  c(0, 0, 1, 0, 0, .75, 0, 0, .5), # perfectly periodic, with 3 harmonics
  rep(0, 9),                      # a silent frame
  rep(1, 9)                       # white noise
)

# Weiner entropy is ~0 for periodic, NA for silent, 1 for white noise
sapply(s, function(x) round(getEntropy(x), 2))

# Shannon entropy is ~0 for periodic with a single harmonic, moderate for
# periodic with multiple harmonics, NA for silent, highest for white noise
sapply(s, function(x) round(getEntropy(x, type = 'shannon'), 2))

# Normalized Shannon entropy - same but forced to be 0 to 1
sapply(s, function(x) round(getEntropy(x,
  type = 'shannon', normalize = TRUE), 2))
```

getEnv

Get amplitude envelope

Description

Returns the smoothed amplitude envelope of a waveform on the original scale. Unlike `seewave::env`, this function always returns an envelope of the same length as the original sound, regardless of the amount of smoothing.

Usage

```
getEnv(
  sound,
  windowLength_points,
  method = c("rms", "hil", "peak", "raw", "mean")[1]
)
```

Arguments

sound numeric vector
windowLength_points the length of smoothing window, in points
method 'peak' for peak amplitude per window, 'rms' for root mean square amplitude, 'mean' for mean (for DC offset removal), 'hil' for Hilbert, 'raw' for low-pass filtering the actual sound

Examples

```
a = rnorm(500) * seq(1, 0, length.out = 500)
windowLength_points = 50
scale = max(abs(a))
plot(a, type = 'l', ylim = c(-scale, scale))
points(soundgen:::getEnv(a, windowLength_points, 'rms'),
       type = 'l', col = 'red')
points(soundgen:::getEnv(a, windowLength_points, 'peak'),
       type = 'l', col = 'green')
points(soundgen:::getEnv(a, windowLength_points, 'hil'),
       type = 'l', col = 'blue')
points(soundgen:::getEnv(a, windowLength_points, 'mean'),
       type = 'l', lty = 3, lwd = 3)
```

getIntegerRandomWalk *Discrete random walk*

Description

Takes a continuous random walk and converts it to continuous epochs of repeated values 0/1/2, each at least minLength points long. 0/1/2 correspond to different noise regimes: 0 = no noise, 1 = subharmonics, 2 = subharmonics and jitter/shimmer.

Usage

```
getIntegerRandomWalk(
  rw,
  nonlinBalance = 50,
  minLength = 50,
  q1 = NULL,
```

```

    q2 = NULL,
    plot = FALSE
)

```

Arguments

rw	a random walk generated by getRandomWalk (expected range 0 to 100)
nonlinBalance	a number between 0 to 100: 0 = returns all zeros; 100 = returns all twos
minLength	the minimum length of each epoch
q1, q2	cutoff points for transitioning from regime 0 to 1 (q1) or from regime 1 to 2 (q2). See noiseThresholdsDict for defaults
plot	if TRUE, plots the random walk underlying nonlinear regimes

Value

Returns a vector of integers (0/1/2) of the same length as rw.

Examples

```

rw = getRandomWalk(len = 100, rw_range = 100, rw_smoothing = .2)
r = getIntegerRandomWalk(rw, nonlinBalance = 75,
                        minLength = 10, plot = TRUE)
r = getIntegerRandomWalk(rw, nonlinBalance = 15,
                        q1 = 30, q2 = 70,
                        minLength = 10, plot = TRUE)

```

getLoudness

Get loudness

Description

Estimates subjective loudness per frame, in sone. Based on EMBSD speech quality measure, particularly the matlab code in Yang (1999) and Timoney et al. (2004). Note that there are many ways to estimate loudness and many other factors, ignored by this model, that could influence subjectively experienced loudness. Please treat the output with a healthy dose of skepticism! Also note that the absolute value of calculated loudness critically depends on the chosen "measured" sound pressure level (SPL). `getLoudness` estimates how loud a sound will be experienced if it is played back at an SPL of `SPL_measured` dB. The most meaningful way to use the output is to compare the loudness of several sounds analyzed with identical settings or of different segments within the same recording.

Usage

```

getLoudness(
  x,
  samplingRate = NULL,
  scale = NULL,
  from = NULL,
  to = NULL,
  windowLength = 50,
  step = NULL,
  overlap = 50,
  SPL_measured = 70,
  Pref = 2e-05,
  spreadSpectrum = TRUE,
  summaryFun = c("mean", "median", "sd"),
  reportEvery = NULL,
  cores = 1,
  plot = TRUE,
  savePlots = NULL,
  main = NULL,
  ylim = NULL,
  width = 900,
  height = 500,
  units = "px",
  res = NA,
  mar = c(5.1, 4.1, 4.1, 4.1),
  ...
)

```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>scale</code>	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
<code>from, to</code>	if <code>NULL</code> (default), analyzes the whole sound, otherwise from...to (s)
<code>windowLength</code>	length of FFT window, ms
<code>step</code>	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
<code>overlap</code>	overlap between successive FFT frames, %
<code>SPL_measured</code>	sound pressure level at which the sound is presented, dB
<code>Pref</code>	reference pressure, Pa (currently has no effect on the estimate)
<code>spreadSpectrum</code>	if <code>TRUE</code> , applies a spreading function to account for frequency masking

summaryFun	functions used to summarize each acoustic characteristic, eg "c('mean', 'sd')"; user-defined functions are fine (see examples); NAs are omitted automatically for mean/median/sd/min/max/range/sum, otherwise take care of NAs yourself
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
plot	should a spectrogram be plotted? TRUE / FALSE
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
main	plot title
ylim	frequency range to plot, kHz (defaults to 0 to Nyquist frequency). NB: still in kHz, even if yScale = bark, mel, or ERB
width, height, units, res	graphical parameters for saving plots passed to png
mar	margins of the spectrogram
...	other plotting parameters passed to spectrogram

Details

Algorithm: calibrates the sound to the desired SPL (Timoney et al., 2004), extracts a spectrogram with [powspec](#), converts to bark scale with ([audspec](#)), spreads the spectrum to account for frequency masking across the critical bands (Yang, 1999), converts dB to phon by using standard equal loudness curves (ISO 226), converts phon to sone (Timoney et al., 2004), sums across all critical bands, and applies a correction coefficient to standardize output. Calibrated so as to return a loudness of 1 sone for a 1 kHz pure tone with SPL of 40 dB.

Value

Returns a list:

specSone spectrum in bark-sone (one per file): a matrix of loudness values in sone, with frequency on the bark scale in rows and time (STFT frames) in columns

loudness a vector of loudness in sone per STFT frame (one per file)

summary a dataframe of summary loudness measures (one row per file)

References

- ISO 226 as implemented by Jeff Tackett (2005) on <https://www.mathworks.com/matlabcentral/fileexchange/7028-iso-226-equal-loudness-level-contour-signal>
- Timoney, J., Lysaght, T., Schoenwiesner, M., & MacManus, L. (2004). Implementing loudness models in matlab.
- Yang, W. (1999). Enhanced Modified Bark Spectral Distortion (EMBSD): An Objective Speech Quality Measure Based on Audible Distortion and Cognitive Model. Temple University.

See Also[getRMS](#) [analyze](#)**Examples**

```

sounds = list(
  white_noise = runif(8000, -1, 1),
  white_noise2 = runif(8000, -1, 1) / 2, # ~6 dB quieter
  pure_tone_1KHz = sin(2*pi*1000/16000*(1:8000)) # pure tone at 1 kHz
)
l = getLoudness(
  x = sounds, samplingRate = 16000, scale = 1,
  windowLength = 20, step = NULL,
  overlap = 50, SPL_measured = 40,
  Pref = 2e-5, plot = FALSE)
l$summary
# white noise (sound 1) is twice as loud as pure tone at 1 KHz (sound 3),
# and note that the same white noise with lower amplitude has lower loudness
# (provided that "scale" is specified)
# compare: lapply(sounds, range)

## Not run:
s = soundgen()
# playme(s)
l1 = getLoudness(s, samplingRate = 16000, SPL_measured = 70)
l1$summary
# The estimated loudness in sone depends on target SPL
l2 = getLoudness(s, samplingRate = 16000, SPL_measured = 40)
l2$summary

# ...but not (much) on windowLength and samplingRate
l3 = getLoudness(s, samplingRate = 16000, SPL_measured = 40, windowLength = 50)
l3$summary

# input can be an audio file...
getLoudness('~Downloads/temp/032_ut_anger_30-m-roar-curse.wav')

...or a folder with multiple audio files
getLoudness('~Downloads/temp2', plot = FALSE)$summary
# Compare:
analyze('~Downloads/temp2', pitchMethods = NULL,
  plot = FALSE, silence = 0)$summary$loudness_mean
# (per STFT frame; should be similar if silence = 0, because
# otherwise analyze() discards frames considered silent)

# custom summaryFun
ran = function(x) diff(range(x))
getLoudness('~Downloads/temp2', plot = FALSE,
  summaryFun = c('mean', 'ran'))$summary

## End(Not run)

```

getPitchZc

*Zero-crossing rate***Description**

A less precise, but very quick method of pitch tracking based on measuring zero-crossing rate in low-pass-filtered audio. Recommended for processing long recordings with typical pitch values well below the first formant frequency, such as speech. Calling this function is considerably faster than using the same pitch-tracking method in [analyze](#). Note that, unlike `analyze()`, it returns the times of individual zero crossings (hopefully corresponding to glottal cycles) instead of pitch values at fixed time intervals.

Usage

```
getPitchZc(
  x,
  samplingRate = NULL,
  scale = NULL,
  from = NULL,
  to = NULL,
  pitchFloor = 50,
  pitchCeiling = 400,
  zcThres = 0.1,
  zcWin = 5,
  silence = 0.04,
  envWin = 5,
  summaryFun = c("mean", "sd"),
  reportEvery = NULL
)
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>scale</code>	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
<code>from</code> , <code>to</code>	if <code>NULL</code> (default), analyzes the whole sound, otherwise from...to (s)
<code>pitchFloor</code> , <code>pitchCeiling</code>	absolute bounds for pitch candidates (Hz)
<code>zcThres</code>	pitch candidates with certainty below this value are treated as noise and set to NA (0 = anything goes, 1 = pitch must be perfectly stable over <code>zcWin</code>)
<code>zcWin</code>	certainty in pitch candidates depends on how stable pitch is over <code>zcWin</code> glottal cycles (odd integer > 3)

silence	minimum root mean square (RMS) amplitude, below which pitch candidates are set to NA (NULL = don't consider RMS amplitude)
envWin	window length for calculating RMS envelope, ms
summaryFun	functions used to summarize each acoustic characteristic; see analyze
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)

Details

Algorithm: the audio is bandpass-filtered from `pitchFloor` to `pitchCeiling`, and the timing of all zero crossings is saved. This is not enough, however, because unvoiced sounds like white noise also have plenty of zero crossings. Accordingly, an attempt is made to detect voiced segments (or steady musical tones, etc.) by looking for stable regions, with several zero-crossings at relatively regular intervals (see parameters `zcThres` and `zcWin`). Very quiet parts of audio are also treated as not having a pitch.

Value

Returns a dataframe containing

time time stamps of all zero crossings except the last one, after bandpass-filtering

pitch pitch calculated from the time between consecutive zero crossings

cert certainty in each pitch candidate calculated from local pitch stability, 0 to 1

See Also

[analyze](#)

Examples

```
data(sheep, package = 'seewave')
# spectrogram(sheep)
zc = getPitchZc(sheep, pitchCeiling = 250)
plot(zc$detailed[, c('time', 'pitch')], type = 'b')

# Convert to a standard pitch contour sampled at regular time intervals:
pitch = getSmoothContour(
  anchors = data.frame(time = zc$detailed$time, value = zc$detailed$pitch),
  len = 1000, NA_to_zero = FALSE, discontThres = 0)
spectrogram(sheep, extraContour = pitch, ylim = c(0, 2))

## Not run:
# process all files in a folder
zc = getPitchZc('~Downloads/temp')
zc$summary

## End(Not run)
```

 getPrior

Get prior for pitch candidates

Description

Prior for adjusting the estimated pitch certainties in [analyze](#). For ex., if primarily working with speech, we could prioritize pitch candidates in the expected pitch range (100-1000 Hz) and decrease our confidence in candidates with very high or very low frequency as unlikely but still remotely possible. You can think of this as a "soft" alternative to setting absolute pitch floor and ceiling. Algorithm: the multiplier for each pitch candidate is the density of prior distribution with mean = priorMean (Hz) and sd = priorSD (semitones) normalized so max = 1 over [pitchFloor, pitchCeiling]. Useful for previewing the prior given to [analyze](#).

Usage

```
getPrior(
  priorMean,
  priorSD,
  distribution = c("normal", "gamma")[1],
  pitchFloor = 75,
  pitchCeiling = 3000,
  len = 100,
  plot = TRUE,
  pitchCands = NULL,
  ...
)
```

Arguments

priorMean, priorSD	specifies the mean (Hz) and standard deviation (semitones) of gamma distribution describing our prior knowledge about the most likely pitch values for this file. For ex., priorMean = 300, priorSD = 6 gives a prior with mean = 300 Hz and SD = 6 semitones (half an octave). To avoid using any priors, set priorMean = NA, priorAdapt = FALSE
distribution	the shape of prior distribution on the musical scale: 'normal' (mode = priorMean) or 'gamma' (skewed to lower frequencies)
pitchFloor, pitchCeiling	absolute bounds for pitch candidates (Hz)
len	the required length of output vector (resolution)
plot	if TRUE, plots the prior
pitchCands	a matrix of pitch candidate frequencies (for internal soundgen use)
...	additional graphical parameters passed on to plot()

Value

Returns a numeric vector of certainties of length `len` if `pitchCands` is `NULL` and a numeric matrix of the same dimensions as `pitchCands` otherwise.

See Also

[analyze_pitch_app](#)

Examples

```
soundgen:::getPrior(priorMean = 150, # Hz
                   priorSD = 2)    # semitones
soundgen:::getPrior(150, 6)
s = soundgen:::getPrior(450, 24, pitchCeiling = 6000)
plot(s, type = 'l')
```

getRandomWalk	<i>Random walk</i>
---------------	--------------------

Description

Generates a random walk with flexible control over its range, trend, and smoothness. It works by calling `stats::rnorm` at each step and taking a cumulative sum of the generated values. Smoothness is controlled by initially generating a shorter random walk and upsampling.

Usage

```
getRandomWalk(
  len,
  rw_range = 1,
  rw_smoothing = 0.2,
  method = c("linear", "spline")[2],
  trend = 0
)
```

Arguments

<code>len</code>	an integer specifying the required length of random walk. If <code>len</code> is 1, returns a single draw from a gamma distribution with <code>mean=1</code> and <code>sd=rw_range</code>
<code>rw_range</code>	the upper bound of the generated random walk (the lower bound is set to 0)
<code>rw_smoothing</code>	specifies the amount of smoothing, basically the number of points used to construct the <code>rw</code> as a proportion of <code>len</code> , from 0 (no smoothing) to 1 (maximum smoothing to a straight line)
<code>method</code>	specifies the method of smoothing: either linear interpolation (<code>'linear'</code> , see <code>stats::approx</code>) or cubic splines (<code>'spline'</code> , see <code>stats::spline</code>)

trend mean of generated normal distribution (vectors are also acceptable, as long as their length is an integer multiple of len). If positive, the random walk has an overall upwards trend (good values are between 0 and 0.5 or -0.5). Trend = c(1,-1) gives a roughly bell-shaped rw with an upward and a downward curve. Larger absolute values of trend produce less and less random behavior

Value

Returns a numeric vector of length len and range from 0 to rw_range.

Examples

```
plot(getRandomWalk(len = 1000, rw_range = 5, rw_smoothing = 0))
plot(getRandomWalk(len = 1000, rw_range = 5, rw_smoothing = .2))
plot(getRandomWalk(len = 1000, rw_range = 5, rw_smoothing = .95))
plot(getRandomWalk(len = 1000, rw_range = 5, rw_smoothing = .99))
plot(getRandomWalk(len = 1000, rw_range = 5, rw_smoothing = 1))
plot(getRandomWalk(len = 1000, rw_range = 15,
  rw_smoothing = .2, trend = c(.1, -.1)))
plot(getRandomWalk(len = 1000, rw_range = 15,
  rw_smoothing = .2, trend = c(15, -1)))
```

getRMS

RMS amplitude

Description

Calculates root mean square (RMS) amplitude in overlapping windows, providing an envelope of RMS amplitude - a measure of sound intensity. Longer windows provide smoother, more robust estimates; shorter windows and more overlap improve temporal resolution, but they also increase processing time and make the contour less smooth.

Usage

```
getRMS(
  x,
  samplingRate = NULL,
  scale = NULL,
  from = NULL,
  to = NULL,
  windowLength = 50,
  step = NULL,
  overlap = 70,
  stereo = c("left", "right", "average", "both")[1],
  killDC = FALSE,
  normalize = TRUE,
  windowDC = 200,
  summaryFun = "mean",
```

```

    reportEvery = NULL,
    cores = 1,
    plot = FALSE,
    savePlots = NULL,
    main = NULL,
    xlab = "",
    ylab = "",
    type = "b",
    col = "green",
    lwd = 2,
    width = 900,
    height = 500,
    units = "px",
    res = NA,
    ...
)

```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
scale	maximum possible amplitude of input used for normalization of input vector (only needed if x is a numeric vector)
from, to	if NULL (default), analyzes the whole sound, otherwise from...to (s)
windowLength	length of FFT window, ms
step	you can override overlap by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of 1/samplingRate, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
overlap	overlap between successive FFT frames, %
stereo	'left' = only left channel, 'right' = only right channel, 'average' = take the mean of the two channels, 'both' = return RMS for both channels separately
killDC	if TRUE, removed DC offset (see also flatEnv)
normalize	if TRUE, the RMS amplitude is returned as proportion of the maximum possible amplitude as given by scale
windowDC	the window for calculating DC offset, ms
summaryFun	functions used to summarize each acoustic characteristic; see analyze
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
plot	if TRUE, plot a contour of RMS amplitude
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)

xlab, ylab, main
 general graphical parameters
 type, col, lwd graphical parameters pertaining to the RMS envelope
 width, height, units, res
 graphical parameters for saving plots passed to [png](#)
 ... other graphical parameters

Details

Note that you can also get similar estimates per frame from [analyze](#) on a normalized scale of 0 to 1, but `getRMS` is much faster, operates on the original scale, and plots the amplitude contour. If you need RMS for the entire sound instead of per frame, you can simply calculate it as $\sqrt{\text{mean}(x^2)}$, where x is your waveform. Having RMS estimates per frame gives more flexibility: RMS per sound can be calculated as the mean / median / max of RMS values per frame.

Value

Returns a list containing:

\$detailed: a list of RMS amplitudes per frame for each sound, on the scale of input; names give time stamps for the center of each frame, in ms.

\$summary: a dataframe with summary measures, one row per sound

See Also

[analyze](#) [getLoudness](#)

Examples

```
s = soundgen() + .25 # with added DC offset
# osc(s)
r = getRMS(s, samplingRate = 16000, from = .05,
  windowLength = 40, overlap = 50, killDC = TRUE,
  plot = TRUE, type = 'l', lty = 2, main = 'RMS envelope')
r
# short window = jagged envelope
r = getRMS(s, samplingRate = 16000,
  windowLength = 5, overlap = 0, killDC = TRUE,
  plot = TRUE, col = 'blue', pch = 13, main = 'RMS envelope')

# stereo
wave_stereo = tuneR::Wave(
  left = runif(1000, -1, 1) * 16000,
  right = runif(1000, -1, 1) / 3 * 16000,
  bit = 16, samp.rate = 4000)
getRMS(wave_stereo)$summary
getRMS(wave_stereo, stereo = 'right')$summary
getRMS(wave_stereo, stereo = 'average')$summary
getRMS(wave_stereo, from = .05,
  stereo = 'both', plot = TRUE)$summary
```

```

## Not run:
r = getRMS('~Downloads/temp', savePlots = '~Downloads/temp/plots')
r$summary

# Compare:
analyze('~Downloads/temp', pitchMethods = NULL,
        plot = FALSE)$ampl_mean
# (per STFT frame, but should be very similar)

User-defined summary functions:
ran = function(x) diff(range(x))
meanSD = function(x) {
  paste0('mean = ', round(mean(x), 2), '; sd = ', round(sd(x), 2))
}
getRMS('~Downloads/temp', summaryFun = c('mean', 'ran', 'meanSD'))$summary

## End(Not run)

```

getRolloff

Control rolloff of harmonics

Description

Harmonics are generated as separate sine waves. But we don't want each harmonic to be equally strong, so we normally specify some rolloff function that describes the loss of energy in upper harmonics relative to the fundamental frequency (f_0). `getRolloff` provides flexible control over this rolloff function, going beyond simple exponential decay (`rolloff`). Use quadratic terms to modify the behavior of a few lower harmonics, `rolloffOct` to adjust the rate of decay per octave, and `rolloffKHz` for rolloff correction depending on f_0 . Plot the output with different parameter values and see examples below and the vignette to get a feel for how to use `getRolloff` effectively.

Usage

```

getRolloff(
  pitch_per_gc = c(440),
  nHarmonics = NULL,
  rolloff = -6,
  rolloffOct = 0,
  rolloffParab = 0,
  rolloffParabHarm = 3,
  rolloffParabCeiling = NULL,
  rolloffKHz = 0,
  baseline = 200,
  dynamicRange = 80,
  samplingRate = 16000,
  plot = FALSE
)

```

Arguments

pitch_per_gc	a vector of f0 per glottal cycle, Hz
nHarmonics	maximum number of harmonics to generate (very weak harmonics with amplitude < -dynamicRange will be discarded)
rolloff	basic rolloff from lower to upper harmonics, db/octave (exponential decay). All rolloff parameters are in anchor format. See getRolloff for more details
rolloffOct	basic rolloff changes from lower to upper harmonics (regardless of f0) by rolloffOct dB/oct. For example, we can get steeper rolloff in the upper part of the spectrum
rolloffParab	an optional quadratic term affecting only the first rolloffParabHarm harmonics. The middle harmonic of the first rolloffParabHarm harmonics is amplified or dampened by rolloffParab dB relative to the basic exponential decay
rolloffParabHarm	the number of harmonics affected by rolloffParab
rolloffParabCeiling	quadratic adjustment is applied only up to rolloffParabCeiling, Hz. If not NULL, it overrides rolloffParabHarm
rolloffKHz	rolloff changes linearly with f0 by rolloffKHz dB/kHz. For ex., -6 dB/kHz gives a 6 dB steeper basic rolloff as f0 goes up by 1000 Hz
baseline	The "neutral" f0, at which no adjustment of rolloff takes place regardless of rolloffKHz
dynamicRange	dynamic range, dB. Harmonics and noise more than dynamicRange under maximum amplitude are discarded to save computational resources
samplingRate	sampling rate (needed to stop at Nyquist frequency and for plotting purposes)
plot	if TRUE, produces a plot

Value

Returns a matrix of amplitude multiplication factors for adjusting the amplitude of harmonics relative to f0 (1 = no adjustment, 0 = silent). Each row of output contains one harmonic, and each column contains one glottal cycle.

See Also

[soundgen](#)

Examples

```
# steady exponential rolloff of -12 dB per octave
rolloff = getRolloff(pitch_per_gc = 150, rolloff = -12,
  rolloffOct = 0, rolloffKHz = 0, plot = TRUE)
# the rate of rolloff slows down by 1 dB each octave
rolloff = getRolloff(pitch_per_gc = 150, rolloff = -12,
  rolloffOct = 1, rolloffKHz = 0, plot = TRUE)

# rolloff can be made to depend on f0 using rolloffKHz
rolloff = getRolloff(pitch_per_gc = c(150, 400, 800),
```



```

    rolloffOct = 0, rolloffKHz = -3, plot = TRUE)
# without the correction for f0 (rolloffKHz),
# high-pitched sounds have the same rolloff as low-pitched sounds,
# producing unnaturally strong high-frequency harmonics
rolloff = getRolloff(pitch_per_gc = c(150, 400, 800),
    rolloffOct = 0, rolloffKHz = 0, plot = TRUE)

# parabolic adjustment of lower harmonics
rolloff = getRolloff(pitch_per_gc = 350, rolloffParab = 0,
    rolloffParabHarm = 2, plot = TRUE)
# rolloffParabHarm = 1 affects only f0
rolloff = getRolloff(pitch_per_gc = 150, rolloffParab = 30,
    rolloffParabHarm = 1, plot = TRUE)
# rolloffParabHarm=2 or 3 affects only h1
rolloff = getRolloff(pitch_per_gc = 150, rolloffParab = 30,
    rolloffParabHarm = 2, plot = TRUE)
# rolloffParabHarm = 4 affects h1 and h2, etc
rolloff = getRolloff(pitch_per_gc = 150, rolloffParab = 30,
    rolloffParabHarm = 4, plot = TRUE)
# negative rolloffParab weakens lower harmonics
rolloff = getRolloff(pitch_per_gc = 150, rolloffParab = -20,
    rolloffParabHarm = 7, plot = TRUE)
# only harmonics below 2000 Hz are affected
rolloff = getRolloff(pitch_per_gc = c(150, 600),
    rolloffParab = -20, rolloffParabCeiling = 2000,
    plot = TRUE)

# dynamic rolloff (varies over time)
rolloff = getRolloff(pitch_per_gc = c(150, 250),
    rolloff = c(-12, -18, -24), plot = TRUE)
rolloff = getRolloff(pitch_per_gc = c(150, 250), rolloffParab = 40,
    rolloffParabHarm = 1:5, plot = TRUE)

## Not run:
# Note: getRolloff() is called internally by soundgen()
# using the data.frame format for all vectorized parameters
# Compare:
s1 = soundgen(syllLen = 1000, pitch = 250,
    rolloff = c(-24, -2, -18), plot = TRUE)
s2 = soundgen(syllLen = 1000, pitch = 250,
    rolloff = data.frame(time = c(0, .2, 1),
        value = c(-24, -2, -18)),
    plot = TRUE)

# Also works for rolloffOct, rolloffParab, etc:
s3 = soundgen(syllLen = 1000, pitch = 250,
    rolloffParab = 20, rolloffParabHarm = 1:15, plot = TRUE)

## End(Not run)

```

Description

Returns a smooth contour based on an arbitrary number of anchors. Used by [soundgen](#) for generating intonation contour, mouth opening, etc. This function is mostly intended to be used internally by [soundgen](#), more precisely to construct (upsample) smooth curves from a number of anchors. For general upsampling or downsampling of audio, use [resample](#). Note that pitch contours are treated as a special case: values are log-transformed prior to smoothing, so that with 2 anchors we get a linear transition on a log scale (as if we were operating with musical notes rather than frequencies in Hz). Pitch plots have two Y axes: one showing Hz and the other showing musical notation.

Usage

```
getSmoothContour(
  anchors = data.frame(time = c(0, 1), value = c(0, 1)),
  len = NULL,
  thisIsPitch = FALSE,
  normalizeTime = TRUE,
  interpol = c("approx", "spline", "loess")[3],
  loessSpan = NULL,
  discontinThres = 0.05,
  jumpThres = 0.01,
  valueFloor = NULL,
  valueCeiling = NULL,
  plot = FALSE,
  xlim = NULL,
  ylim = NULL,
  xlab = "Time, ms",
  ylab = ifelse(thisIsPitch, "Frequency, Hz", "Amplitude"),
  main = ifelse(thisIsPitch, "Pitch contour", ""),
  samplingRate = 16000,
  voiced = NULL,
  contourLabel = NULL,
  NA_to_zero = TRUE,
  ...
)
```

Arguments

anchors	a numeric vector of values or a list/dataframe with one column (value) or two columns (time and value). anchors\$time can be in ms (with len=NULL) or in arbitrary units, eg 0 to 1 (with duration determined by len, which must then be provided in ms). So anchors\$time is assumed to be in ms if len=NULL and relative if len is specified. anchors\$value can be on any scale.
len	the required length of the output contour. If NULL, it will be calculated based on the maximum time value (in ms) and samplingRate
thisIsPitch	(boolean) is this a pitch contour? If true, log-transforms before smoothing and plots in both Hz and musical notation
normalizeTime	if TRUE, normalizes anchors\$time values to range from 0 to 1

interpol	method of interpolation between anchors: "approx" = linear with approx , "spline" = cubic splines with spline , "loess" = local polynomial regression with loess
loessSpan	controls the amount of smoothing when interpolating between anchors with loess , so only has an effect if interpol = 'loess' (1 = strong, 0.5 = weak smoothing)
discontThres	if two anchors are closer in time than discontThres (on a 0-1 scale, ie specified as proportion of total length), the contour is broken into segments with a linear transition between these segments
jumpThres	if anchors are closer than jumpThres, a new section starts with no transition at all (e.g. for adding pitch jumps)
valueFloor, valueCeiling	lower/upper bounds for the contour
plot	(boolean) produce a plot?
xlim, ylim, xlab, ylab, main	plotting options
samplingRate	sampling rate used to convert time values to points (Hz)
voiced, contourLabel	graphical pars for plotting breathing contours (see examples below)
NA_to_zero	if TRUE, all NAs are replaced with zero
...	other plotting options passed to plot()

Value

Returns a numeric vector of length len.

Examples

```
# long format: anchors are a dataframe
a = getSmoothContour(anchors = data.frame(
  time = c(50, 137, 300), value = c(0.03, 0.78, 0.5)),
  normalizeTime = FALSE,
  voiced = 200, valueFloor = 0, plot = TRUE, main = '',
  samplingRate = 16000) # breathing

# short format: anchors are a vector (equal time steps assumed)
a = getSmoothContour(anchors = c(350, 800, 600),
  len = 5500, thisIsPitch = TRUE, plot = TRUE,
  samplingRate = 3500) # pitch

# a single anchor gives constant value
a = getSmoothContour(anchors = 800,
  len = 500, thisIsPitch = TRUE, plot = TRUE, samplingRate = 500)

# two pitch anchors give loglinear F0 change
a = getSmoothContour(anchors = c(220, 440),
  len = 500, thisIsPitch = TRUE, plot = TRUE, samplingRate = 500)

## Two closely spaced anchors produce a pitch jump
```

```

# one loess for the entire contour
a1 = getSmoothContour(anchors = list(time = c(0, .15, .2, .7, 1),
  value = c(360, 116, 550, 700, 610)), len = 500, thisIsPitch = TRUE,
  plot = TRUE, samplingRate = 500)
# two segments with a linear transition
a2 = getSmoothContour(anchors = list(time = c(0, .15, .17, .7, 1),
  value = c(360, 116, 550, 700, 610)), len = 500, thisIsPitch = TRUE,
  plot = TRUE, samplingRate = 500)
# two segments with an abrupt jump
a3 = getSmoothContour(anchors = list(time = c(0, .15, .155, .7, 1),
  value = c(360, 116, 550, 700, 610)), len = 500, thisIsPitch = TRUE,
  plot = TRUE, samplingRate = 500)
# compare:
plot(a2)
plot(a3) # NB: the segment before the jump is upsampled to compensate

## Control the amount of smoothing
getSmoothContour(c(1, 3, 9, 10, 9, 9, 2), len = 100, plot = TRUE,
  loessSpan = NULL) # default amount of smoothing (depends on dur)
getSmoothContour(c(1, 3, 9, 10, 9, 9, 2), len = 100, plot = TRUE,
  loessSpan = .85) # more smoothing than default
getSmoothContour(c(1, 3, 9, 10, 9, 9, 2), len = 100, plot = TRUE,
  loessSpan = .5) # less smoothing
getSmoothContour(c(1, 3, 9, 10, 9, 9, 2), len = 100, plot = TRUE,
  interpol = 'approx') # linear interpolation (no smoothing)

## Upsample preserving leading and trailing NAs
anchors = data.frame(time = c(1, 4, 5, 7, 10, 20, 23, 25, 30),
  value = c(NA, NA, 10, 15, 12, NA, 17, 15, NA))
plot(anchors, type = 'b')
anchors_ups = getSmoothContour(
  anchors, len = 200,
  interpol = 'approx', # only approx can propagate NAs
  NA_to_zero = FALSE, # preserve NAs
  discontinThres = 0) # don't break into sub-contours
plot(anchors_ups, type = 'b')

```

getSpectralEnvelope *Spectral envelope*

Description

Prepares a spectral envelope for filtering a sound to add formants, lip radiation, and some stochastic component regulated by temperature. Formants are specified as a list containing time, frequency, amplitude, and width values for each formant (see examples). See vignette('sound_generation', package = 'soundgen') for more information.

Usage

```
getSpectralEnvelope(
```

```

nr,
nc,
formants = NA,
formantDep = 1,
formantWidth = 1,
lipRad = 6,
noseRad = 4,
mouth = NA,
mouthOpenThres = 0.2,
openMouthBoost = 0,
vocalTract = NULL,
temperature = 0.05,
formDrift = 0.3,
formDisp = 0.2,
formantDepStoch = 1,
smoothLinearFactor = 1,
formantCeiling = 2,
samplingRate = 16000,
speedSound = 35400,
smoothing = list(),
output = c("simple", "detailed")[1],
plot = FALSE,
duration = NULL,
colorTheme = c("bw", "seewave", "...")[1],
col = NULL,
xlab = "Time",
ylab = "Frequency, kHz",
...
)

```

Arguments

nr	the number of frequency bins = $\text{windowLength_points}/2$, where <code>windowLength_points</code> is the size of window for Fourier transform
nc	the number of time steps for Fourier transform
formants	a character string like "aau" referring to default presets for speaker "M1"; a vector of formant frequencies; or a list of formant times, frequencies, amplitudes, and bandwidths, with a single value of each for static or multiple values of each for moving formants. <code>formants = NA</code> defaults to <code>schwa</code> . Time stamps for formants and <code>mouthOpening</code> can be specified in ms or an any other arbitrary scale.
formantDep	scale factor of formant amplitude (1 = no change relative to amplitudes in formants)
formantWidth	scale factor of formant bandwidth (1 = no change)
lipRad	the effect of lip radiation on source spectrum, dB/oct (the default of +6 dB/oct produces a high-frequency boost when the mouth is open)
noseRad	the effect of radiation through the nose on source spectrum, dB/oct (the alternative to <code>lipRad</code> when the mouth is closed)

mouth	mouth opening (0 to 1, 0.5 = neutral, i.e. no modification) (anchor format)
mouthOpenThres	open the lips (switch from nose radiation to lip radiation) when the mouth is open >mouthOpenThres, 0 to 1
openMouthBoost	amplify the voice when the mouth is open by openMouthBoost dB
vocalTract	the length of vocal tract, cm. Used for calculating formant dispersion (for adding extra formants) and formant transitions as the mouth opens and closes. If NULL or NA, the length is estimated based on specified formant frequencies, if any (anchor format)
temperature	hyperparameter for regulating the amount of stochasticity in sound generation
formDrift	scale factor regulating the effect of temperature on the depth of random drift of all formants (user-defined and stochastic): the higher, the more formants drift at a given temperature
formDisp	scale factor regulating the effect of temperature on the irregularity of the dispersion of stochastic formants: the higher, the more unevenly stochastic formants are spaced at a given temperature
formantDepStoch	multiplication factor for the amplitude of additional formants added above the highest specified formant (0 = none, 1 = default)
smoothLinearFactor	regulates smoothing of formant anchors (0 to +Inf) as they are upsampled to the number of fft steps nc. This is necessary because the input formants normally contains fewer sets of formant values than the number of fft steps. smoothLinearFactor = 0: close to default spline; >3: approaches linear extrapolation
formantCeiling	frequency to which stochastic formants are calculated, in multiples of the Nyquist frequency; increase up to ~10 for long vocal tracts to avoid losing energy in the upper part of the spectrum
samplingRate	sampling frequency, Hz
speedSound	speed of sound in warm air, cm/s. Stevens (2000) "Acoustic phonetics", p. 138
smoothing	a list of parameters passed to getSmoothContour to control the interpolation and smoothing of contours: <code>interpol</code> (approx / spline / loess), <code>loessSpan</code> , <code>discontThres</code> , <code>jumpThres</code>
output	"simple" returns just the spectral filter, while "detailed" also returns a data.frame of formant frequencies over time (needed for internal purposes such as formant locking)
plot	if TRUE, produces a plot of the spectral envelope
duration	duration of the sound, ms (for plotting purposes only)
colorTheme	black and white ('bw'), as in seewave package ('seewave'), or another color theme (e.g. 'heat.colors')
col	actual colors, eg <code>rev(rainbow(100))</code> - see <code>?hcl.colors</code> for colors in base R (overrides colorTheme)
xlab, ylab	labels of axes
...	other graphical parameters passed on to <code>image()</code>

Value

Returns a spectral filter: a matrix with frequency bins in rows and time steps in columns. Accordingly, rownames of the output give central frequency of each bin (in kHz), while colnames give time stamps (in ms if duration is specified, otherwise 0 to 1).

Examples

```
# [a] with only F1-F3 visible, with no stochasticity
e = getSpectralEnvelope(nr = 512, nc = 50, duration = 300,
  formants = soundgen:::convertStringToFormants('a'),
  temperature = 0, plot = TRUE, col = heat.colors(150))
# image(t(e)) # to plot the output on a linear scale instead of dB

# some "wiggling" of specified formants plus extra formants on top
e = getSpectralEnvelope(nr = 512, nc = 50,
  formants = c(860, 1430, 2900),
  temperature = 0.1, formantDepStoch = 1, plot = TRUE)

# a schwa based on variable length of vocal tract
e = getSpectralEnvelope(nr = 512, nc = 100, formants = NA,
  vocalTract = list(time = c(0, .4, 1), value = c(13, 18, 17)),
  temperature = .1, plot = TRUE)

# no formants at all, only lip radiation
e = getSpectralEnvelope(nr = 512, nc = 50, lipRad = 6,
  formants = NA, temperature = 0, plot = FALSE)
plot(e[, 1], type = 'l') # linear scale
plot(20 * log10(e[, 1]), type = 'l') # dB scale - 6 dB/oct

# mouth opening
e = getSpectralEnvelope(nr = 512, nc = 50,
  vocalTract = 16, plot = TRUE, lipRad = 6, noseRad = 4,
  mouth = data.frame(time = c(0, .5, 1), value = c(0, 0, .5)))

# scale formant amplitude and/or bandwidth
e1 = getSpectralEnvelope(nr = 512, nc = 50,
  formants = soundgen:::convertStringToFormants('a'),
  formantWidth = 1, formantDep = 1) # defaults
e2 = getSpectralEnvelope(nr = 512, nc = 50,
  formants = soundgen:::convertStringToFormants('a'),
  formantWidth = 1.5, formantDep = 1.5)
plot(as.numeric(rownames(e2)), 20 * log10(e2[, 1]),
  type = 'l', xlab = 'KHz', ylab = 'dB', col = 'red', lty = 2)
points(as.numeric(rownames(e1)), 20 * log10(e1[, 1]), type = 'l')

# manual specification of formants
e3 = getSpectralEnvelope(
  nr = 512, nc = 50, samplingRate = 16000, plot = TRUE,
  formants = list(
    f1 = list(freq = c(900, 500), amp = c(30, 35), width = c(80, 50)),
    f2 = list(freq = c(1900, 2500), amp = c(25, 30), width = 100),
    f3 = list(freq = 3400, amp = 30, width = 120)
```

```

))

# extra zero-pole pair (doesn't affect estimated VTL and thus the extra
# formants added on top)
e4 = getSpectralEnvelope(
  nr = 512, nc = 50, samplingRate = 16000, plot = TRUE,
  formants = list(
    f1 = list(freq = c(900, 500), amp = c(30, 35), width = c(80, 50)),
    f1.5 = list(freq = 1300, amp = -15),
    f1.7 = list(freq = 1500, amp = 15),
    f2 = list(freq = c(1900, 2500), amp = c(25, 30), width = 100),
    f3 = list(freq = 3400, amp = 30, width = 120)
  )
)
plot(as.numeric(rownames(e4)), 20 * log10(e3[, ncol(e3)]),
     type = 'l', xlab = 'KHz', ylab = 'dB')
points(as.numeric(rownames(e4)), 20 * log10(e4[, ncol(e4)]),
       type = 'l', col = 'red', lty = 2)

```

getSurprisal

Get surprisal

Description

Tracks the (un)predictability of spectral changes in a sound over time, returning a continuous contour of "surprisal". This is an attempt to track auditory salience over time - that is, to identify parts of a sound that are likely to involuntarily attract the listeners' attention. The function returns surprisal proper ('\$surprisal') and its product with increases in loudness ('\$surprisalLoudness'). Because `getSurprisal()` is slow and experimental, it is not called by `analyze()`.

Usage

```

getSurprisal(
  x,
  samplingRate = NULL,
  scale = NULL,
  from = NULL,
  to = NULL,
  step = 20,
  winSurp = 2000,
  method = c("acf", "np")[1],
  yScale = c("bark", "mel", "log")[1],
  nFilters = 64,
  dynamicRange = 80,
  minFreq = 20,
  maxFreq = samplingRate/2,
  summaryFun = "mean",
  reportEvery = NULL,
  cores = 1,

```



```

plot = TRUE,
savePlots = NULL,
osc = c("none", "linear", "dB")[2],
heights = c(3, 1),
ylim = NULL,
contrast = 0.2,
brightness = 0,
maxPoints = c(1e+05, 5e+05),
padWithSilence = TRUE,
colorTheme = c("bw", "seewave", "heat.colors", "...")[1],
col = NULL,
extraContour = NULL,
xlab = NULL,
ylab = NULL,
xaxp = NULL,
mar = c(5.1, 4.1, 4.1, 2),
main = NULL,
grid = NULL,
width = 900,
height = 500,
units = "px",
res = NA,
...
)

```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>scale</code>	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
<code>from, to</code>	if <code>NULL</code> (default), analyzes the whole sound, otherwise <code>from...to</code> (s)
<code>step</code>	step, ms (determines time resolution). <code>step = NULL</code> means no downsampling at all (ncol of output = length of input audio)
<code>winSurp</code>	surprisal analysis window, ms (<code>Inf</code> = from sound onset to each point)
<code>method</code>	<code>acf</code> = change in maximum autocorrelation after adding the final point, <code>np</code> = non-linear prediction (see nonlinPred)
<code>yScale</code>	scale of the frequency axis: <code>'linear'</code> = linear, <code>'log'</code> = logarithmic (musical), <code>'bark'</code> = bark with hz2bark , <code>'mel'</code> = mel with hz2mel , <code>'ERB'</code> = Equivalent Rectangular Bandwidths with HzToERB
<code>nFilters</code>	the number of filters (determines frequency resolution)
<code>dynamicRange</code>	dynamic range, dB. All values more than one <code>dynamicRange</code> under maximum are treated as zero
<code>minFreq, maxFreq</code>	the range of frequencies to analyze

summaryFun	functions used to summarize each acoustic characteristic, eg "c('mean', 'sd')"; user-defined functions are fine (see examples); NAs are omitted automatically for mean/median/sd/min/max/range/sum, otherwise take care of NAs yourself
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
plot	if TRUE, plots the auditory spectrogram and the surprisalLoudness contour
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
osc	"none" = no oscillogram; "linear" = on the original scale; "dB" = in decibels
heights	a vector of length two specifying the relative height of the spectrogram and the oscillogram (including time axes labels)
ylim	frequency range to plot, kHz (defaults to 0 to Nyquist frequency). NB: still in kHz, even if yScale = bark, mel, or ERB
contrast	spectrum is exponentiated by contrast (any real number, recommended -1 to +1). Contrast >0 increases sharpness, <0 decreases sharpness
brightness	how much to "lighten" the image (>0 = lighter, <0 = darker)
maxPoints	the maximum number of "pixels" in the oscillogram (if any) and spectrogram; good for quickly plotting long audio files; defaults to c(1e5, 5e5)
padWithSilence	if TRUE, pads the sound with just enough silence to resolve the edges properly (only the original region is plotted, so the apparent duration doesn't change)
colorTheme	black and white ('bw'), as in seewave package ('seewave'), or any palette from palette such as 'heat.colors', 'cm.colors', etc
col	actual colors, eg rev(rainbow(100)) - see ?hcl.colors for colors in base R (overrides colorTheme)
extraContour	a vector of arbitrary length scaled in Hz (regardless of yScale!) that will be plotted over the spectrogram (eg pitch contour); can also be a list with extra graphical parameters such as lwd, col, etc. (see examples)
xlab, ylab, main, mar, xaxp	graphical parameters for plotting
grid	if numeric, adds n = grid dotted lines per kHz
width, height, units, res	graphical parameters for saving plots passed to png
...	other graphical parameters

Details

Algorithm: we start with an auditory spectrogram produced by applying a bank of bandpass filters to the signal, by default with central frequencies equally spaced on the bark scale (see [audSpectrogram](#)). For each frequency channel, a sliding window is analyzed to compare the actually observed final value with its expected value. There are many ways to extrapolate / predict time series and thus perform this comparison such as autocorrelation (method = 'acf') or nonlinear prediction (method = 'np'). The resulting per-channel surprisal contours are aggregated by taking their mean weighted by the average amplitude of each frequency channel across the analysis window. Because increases in loudness are known to be important predictors of auditory salience, loudness per frame is also returned, as well as the square root of the product of its derivative and surprisal.

Value

Returns a list with \$detailed per-frame and \$summary per-file results (see [analyze](#) for more information). Three measures are reported: loudness (in sone, as per [getLoudness](#)), the first derivative of loudness with respect to time (dLoudness), surprisal (non-negative), and surprisalLoudness (geometric mean of surprisal and dLoudness, treating negative values of dLoudness as zero).

Examples

```
# A quick example
s = soundgen(nSyl = 2, syllLen = 50, pauseLen = 25, addSilence = 15)
surp = getSurprisal(s, samplingRate = 16000)
surp

## Not run:
# A more meaningful example
sound = soundgen(nSyl = 5, syllLen = 150,
  pauseLen = c(50, 50, 50, 130), pitch = c(200, 150),
  noise = list(time = c(-300, 200), value = -20), plot = TRUE)
# playme(sound)
surp = getSurprisal(sound, samplingRate = 16000,
  yScale = 'bark', method = 'acf')
surp = getSurprisal(sound, samplingRate = 16000,
  yScale = 'bark', method = 'np') # very slow

# short window = amnesia (every even is equally surprising)
getSurprisal(sound, samplingRate = 16000, winSurp = 250)
# long window - remembers further into the past, Inf = from the beginning
surp = getSurprisal(sound, samplingRate = 16000, winSurp = Inf)

# plot "pure" surprisal, without weighting by loudness
spectrogram(sound, 16000, extraContour = surp$detailed$surprisal /
  max(surp$detailed$surprisal, na.rm = TRUE) * 8000)

# NB: surprisalLoudness contour is also log-transformed if yScale = 'log',
# so zeros become NAs
surp = getSurprisal(sound, samplingRate = 16000, yScale = 'log')

# add bells and whistles
surp = getSurprisal(sound, samplingRate = 16000,
  yScale = 'mel',
  osc = 'dB', # plot oscillogram in dB
  heights = c(2, 1), # spectro/osc height ratio
  brightness = -.1, # reduce brightness
  # colorTheme = 'heat.colors', # pick color theme...
  col = rev(hcl.colors(30, palette = 'Viridis')), # ...or specify the colors
  cex.lab = .75, cex.axis = .75, # text size and other base graphics pars
  ylim = c(0, 5), # always in kHz
  main = 'Audiogram with surprisal contour', # title
  extraContour = list(col = 'blue', lty = 2, lwd = 2)
# + axis labels, etc
)
```

```
surp = getSurprisal('~Downloads/temp/', savePlots = '~Downloads/temp/surp')
surp$summary

## End(Not run)
```

 hillenbrand

Formants in American vowels

Description

Typical relative frequencies of the first four formants measured in dF units (average spacing between formants, or formant dispersion) above or below schwa based on estimated VTL in American English, from Hillenbrand (1995), who measured F1-F4 in ~1.5K recordings (139 speakers, 12 vowels from each). Audio and formant measurements are freely available online: <https://homepages.wmich.edu/~hillenbr/voweldata>. The dataset below is the result of modeling Hillenbrand's data with brms: `mvbind(F1rel, F2rel) ~ vowel + (vowelspeaker)`. It shows the most credible location of each vowel centroid in the F1Rel-F2Rel space.

Usage

```
hillenbrand
```

Format

An object of class `data.frame` with 12 rows and 5 columns.

Details

A dataframe of 12 observations and 5 columns: "vowel" = vowel (American English), "F1Rel" to "F4Rel" = formant frequencies in dF relative to their neutral, equidistant positions in a perfectly cylindrical vocal tract. See [schwa](#) - this is what `schwa()` returns as `$ff_relative_dF`

References

Hillenbrand, J., Getty, L. A., Clark, M. J., & Wheeler, K. (1995). Acoustic characteristics of American English vowels. *The Journal of the Acoustical society of America*, 97(5), 3099-3111.

Examples

```
plot(hillenbrand$F1Rel, hillenbrand$F2Rel, type = 'n')
text(hillenbrand$F1Rel, hillenbrand$F2Rel, labels = hillenbrand$vowel)
```

HzToERB	<i>Convert Hz to ERB rate</i>
---------	-------------------------------

Description

Converts from Hz to the number of Equivalent Rectangular Bandwidths (ERBs) below input frequency. See <https://www2.ling.su.se/staff/hartmut/bark.htm> and https://en.wikipedia.org/wiki/Equivalent_rectangular_bandwidth

Usage

```
HzToERB(h, method = c("linear", "quadratic")[1])
```

Arguments

h	vector or matrix of frequencies (Hz)
method	approximation to use

See Also

[ERBToHz](#) [HzToSemitones](#) [HzToNotes](#)

Examples

```
HzToERB(c(-20, 20, 100, 440, 1000, NA))

f = 20:20000
erb_lin = HzToERB(f, 'linear')
erb_quadratic = HzToERB(f, 'quadratic')
plot(f, erb_lin, log = 'x', type = 'l')
points(f, erb_quadratic, col = 'blue', type = 'l')

# compare with the bark scale:
barks = tuneR::hz2bark(f)
points(f, barks / max(barks) * max(erb_lin),
      col = 'red', type = 'l', lty = 2)
```

HzToNotes	<i>Convert Hz to notes</i>
-----------	----------------------------

Description

Converts from Hz to musical notation like A4 - note A of the fourth octave above C0 (16.35 Hz).

Usage

```
HzToNotes(h, showCents = FALSE, A4 = 440)
```

Arguments

h	vector or matrix of frequencies (Hz)
showCents	if TRUE, show cents to the nearest notes (cent = 1/100 of a semitone)
A4	frequency of note A in the fourth octave (modern standard ISO 16 or concert pitch = 440 Hz)

See Also

[notesToHz](#) [HzToSemitones](#)

Examples

```
HzToNotes(c(440, 293, 115, 16.35, 4))

HzToNotes(c(440, 415, 80, 81), showCents = TRUE)
# 80 Hz is almost exactly midway (+49 cents) between D#2 and E2

# Baroque tuning A415, half a semitone flat relative to concert pitch A440
HzToNotes(c(440, 415, 16.35), A4 = 415)
```

HzToSemitones	<i>Convert Hz to semitones</i>
---------------	--------------------------------

Description

Converts from Hz to semitones above C-5 (~0.5109875 Hz) or another reference frequency. This may not seem very useful, but note that this gives us a nice logarithmic scale for generating natural pitch transitions.

Usage

```
HzToSemitones(h, ref = 0.5109875)
```

Arguments

h	vector or matrix of frequencies (Hz)
ref	frequency of the reference value (defaults to C-5, 0.51 Hz)

See Also

[semitonesToHz](#) [HzToNotes](#)

Examples

```

s = HzToSemitones(c(440, 293, 115))
# to convert to musical notation
notesDict$note[1 + round(s)]
# note the "1 +": semitones ABOVE C-5, i.e. notesDict[1, ] is C-5

# Any reference tone can be specified. For ex., for semitones above C0, use:
HzToSemitones(440, ref = 16.35)
# TIP: see notesDict for a table of Hz frequencies to musical notation

```

```
invertSpectrogram      Invert spectrogram
```

Description

Transforms a spectrogram into a time series with inverse STFT. The problem is that an ordinary spectrogram preserves only the magnitude (modulus) of the complex STFT, while the phase is lost, and without phase it is impossible to reconstruct the original audio accurately. So there are a number of algorithms for "guessing" the phase that would produce an audio whose magnitude spectrogram is very similar to the target spectrogram. Useful for certain filtering operations that modify the magnitude spectrogram followed by inverse STFT, such as filtering in the spectrotemporal modulation domain.

Usage

```

invertSpectrogram(
  spec,
  samplingRate,
  windowLength,
  overlap,
  step = NULL,
  wn = "hanning",
  specType = c("abs", "log", "dB")[1],
  initialPhase = c("zero", "random", "spsi")[3],
  nIter = 50,
  normalize = TRUE,
  play = TRUE,
  verbose = FALSE,
  plotError = TRUE
)

```

Arguments

spec	the spectrogram that is to be transform to a time series: numeric matrix with frequency bins in rows and time frames in columns
samplingRate	sampling rate of x (only needed if x is a numeric vector)
windowLength	length of FFT window, ms

<code>overlap</code>	overlap between successive FFT frames, %
<code>step</code>	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
<code>wn</code>	window type accepted by <code>ftwindow</code> , currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop
<code>specType</code>	the scale of target spectrogram: 'abs' = absolute, 'log' = log-transformed, 'dB' = in decibels
<code>initialPhase</code>	initial phase estimate: "zero" = set all phases to zero; "random" = Gaussian noise; "spsi" (default) = single-pass spectrogram inversion (Beauregard et al., 2015)
<code>nIter</code>	the number of iterations of the GL algorithm (Griffin & Lim, 1984), 0 = don't run
<code>normalize</code>	if TRUE, normalizes the output to range from -1 to +1
<code>play</code>	if TRUE, plays back the reconstructed audio
<code>verbose</code>	if TRUE, prints estimated time left every 10% of GL iterations
<code>plotError</code>	if TRUE, produces a scree plot of squared error over GL iterations (useful for choosing 'nIter')

Details

Algorithm: takes the spectrogram, makes an initial guess at the phase (zero, noise, or a more intelligent estimate by the SPSI algorithm), fine-tunes over 'nIter' iterations with the GL algorithm, reconstructs the complex spectrogram using the best phase estimate, and performs inverse STFT. The single-pass spectrogram inversion (SPSI) algorithm is implemented as described in Beauregard et al. (2015) following the python code at https://github.com/lonce/SPSI_Python. The Griffin-Lim (GL) algorithm is based on Griffin & Lim (1984).

Value

Returns the reconstructed audio as a numeric vector.

References

- Griffin, D., & Lim, J. (1984). Signal estimation from modified short-time Fourier transform. *IEEE Transactions on Acoustics, Speech, and Signal Processing*, 32(2), 236-243.
- Beauregard, G. T., Harish, M., & Wyse, L. (2015, July). Single pass spectrogram inversion. In *2015 IEEE International Conference on Digital Signal Processing (DSP)* (pp. 427-431). IEEE.

See Also

[spectrogram](#) [filterSoundByMS](#)

Examples

```

# Create a spectrogram
samplingRate = 16000
windowLength = 40
overlap = 75
wn = 'hanning'

s = soundgen(samplingRate = samplingRate, addSilence = 100)
spec = spectrogram(s, samplingRate = samplingRate,
  wn = wn, windowLength = windowLength, overlap = overlap,
  padWithSilence = FALSE, output = 'original')

# Invert the spectrogram, attempting to guess the phase
# Note that samplingRate, wn, windowLength, and overlap must be the same as
# in the original (ie you have to know how the spectrogram was created)
s_new = invertSpectrogram(spec, samplingRate = samplingRate,
  windowLength = windowLength, overlap = overlap, wn = wn,
  initialPhase = 'spsi', nIter = 10, specType = 'abs', play = FALSE)

## Not run:
# Verify the quality of audio reconstruction
# playme(s, samplingRate); playme(s_new, samplingRate)
spectrogram(s, samplingRate, osc = TRUE)
spectrogram(s_new, samplingRate, osc = TRUE)

## End(Not run)

```

matchPars

Match soundgen pars (experimental)

Description

Attempts to find settings for [soundgen](#) that will reproduce an existing sound. The principle is to mutate control parameters, trying to improve fit to target. The currently implemented optimization algorithm is simple hill climbing. Disclaimer: this function is experimental and may or may not work for particular tasks. It is intended as a supplement to - not replacement of - manual optimization. See vignette('sound_generation', package = 'soundgen') for more information.

Usage

```

matchPars(
  target,
  samplingRate = NULL,
  pars = NULL,
  init = NULL,
  probMutation = 0.25,
  stepVariance = 0.1,
  maxIter = 50,

```

```

    minExpectedDelta = 0.001,
    compareSoundsPars = list(),
    verbose = TRUE
  )

```

Arguments

target	the sound we want to reproduce using soundgen: path to a .wav file or numeric vector
samplingRate	sampling rate of target (only needed if target is a numeric vector, rather than a .wav file)
pars	arguments to soundgen that we are attempting to optimize
init	a list of initial values for the optimized parameters pars and the values of other arguments to soundgen that are fixed at non-default values (if any)
probMutation	the probability of a parameter mutating per iteration
stepVariance	scale factor for calculating the size of mutations
maxIter	maximum number of mutated sounds produced without improving the fit to target
minExpectedDelta	minimum improvement in fit to target required to accept the new sound candidate
compareSoundsPars	a list of control parameters passed to compareSounds
verbose	if TRUE, plays back the accepted candidate at each iteration and reports the outcome

Value

Returns a list of length 2: \$history contains the tried parameter values together with their fit to target (\$history\$sim), and \$pars contains a list of the final - hopefully the best - parameter settings.

Examples

```

## Not run:
target = soundgen(syllLen = 600, pitch = c(300, 200),
                 rolloff = -15, play = TRUE, plot = TRUE)
# we hope to reproduce this sound

# Match pars based on acoustic analysis alone, without any optimization.
# This *MAY* match temporal structure, pitch, and stationary formants
m1 = matchPars(target = target,
              samplingRate = 16000,
              maxIter = 0, # no optimization, only acoustic analysis
              verbose = TRUE)
cand1 = do.call(soundgen, c(m1$pars, list(
  temperature = 0.001, play = TRUE, plot = TRUE)))

```

```

# Try to improve the match by optimizing rolloff
# (this may take a few minutes to run, and the results may vary)
m2 = matchPars(target = target,
               samplingRate = 16000,
               pars = 'rolloff',
               maxIter = 100,
               verbose = TRUE)
# rolloff should be moving from default (-9) to target (-15):
sapply(m2$history, function(x) x$pars$rolloff)
cand2 = do.call(soundgen, c(m2$pars, list(play = TRUE, plot = TRUE)))

## End(Not run)

```

modulationSpectrum	<i>Modulation spectrum</i>
--------------------	----------------------------

Description

Produces a modulation spectrum of waveform(s) or audio file(s), with temporal modulation along the X axis (Hz) and spectral modulation (1/KHz) along the Y axis. A good visual analogy is decomposing the spectrogram into a sum of ripples of various frequencies and directions. Roughness is calculated as the proportion of energy / amplitude of the modulation spectrum within roughRange of temporal modulation frequencies. The frequency of amplitude modulation (amMsFreq, Hz) is calculated as the highest peak in the smoothed AM function, and its purity (amMsPurity, dB) as the ratio of this peak to the median AM over amRange. For relatively short and steady sounds, set amRes = NULL and analyze the entire sound. For longer sounds and when roughness or AM vary over time, set amRes to get multiple measurements over time (see examples).

Usage

```

modulationSpectrum(
  x,
  samplingRate = NULL,
  scale = NULL,
  from = NULL,
  to = NULL,
  amRes = 5,
  maxDur = 5,
  logSpec = FALSE,
  windowLength = 15,
  step = NULL,
  overlap = 80,
  wn = "hanning",
  zp = 0,
  power = 1,
  normalize = TRUE,
  roughRange = c(30, 150),
  amRange = c(10, 200),

```

```

returnMS = TRUE,
returnComplex = FALSE,
summaryFun = c("mean", "median", "sd"),
averageMS = FALSE,
reportEvery = NULL,
cores = 1,
plot = TRUE,
savePlots = NULL,
logWarpX = NULL,
logWarpY = NULL,
quantiles = c(0.5, 0.8, 0.9),
kernelSize = 5,
kernelSD = 0.5,
colorTheme = c("bw", "seewave", "heat.colors", "...")[1],
col = NULL,
main = NULL,
xlab = "Hz",
ylab = "1/KHz",
xlim = NULL,
ylim = NULL,
width = 900,
height = 500,
units = "px",
res = NA,
...
)

```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>scale</code>	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
<code>from, to</code>	if <code>NULL</code> (default), analyzes the whole sound, otherwise <code>from...to</code> (s)
<code>amRes</code>	target resolution of amplitude modulation, Hz. If <code>NULL</code> , the entire sound is analyzed at once, resulting in a single roughness value (unless it is longer than <code>maxDur</code> , in which case it is analyzed in chunks <code>maxDur</code> s long). If <code>amRes</code> is set, roughness is calculated for windows $\sim 1000/\text{amRes}$ ms long (but at least 3 STFT frames). <code>amRes</code> also affects the amount of smoothing when calculating <code>amMsFreq</code> and <code>amMsPurity</code>
<code>maxDur</code>	sounds longer than <code>maxDur</code> s are split into fragments, and the modulation spectra of all fragments are averaged
<code>logSpec</code>	if <code>TRUE</code> , the spectrogram is log-transformed prior to taking 2D FFT
<code>windowLength</code>	length of FFT window, ms
<code>step</code>	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and

	thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
overlap	overlap between successive FFT frames, %
wn	window type accepted by <code>ftwindow</code> , currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop
zp	window length after zero padding, points
power	raise modulation spectrum to this power (eg power = 2 for $\wedge 2$, or "power spectrum")
normalize	if TRUE, the modulation spectrum of each analyzed fragment <code>maxDur</code> in duration is separately normalized to have <code>max = 1</code>
roughRange	the range of temporal modulation frequencies that constitute the "roughness" zone, Hz
amRange	the range of temporal modulation frequencies that we are interested in as "amplitude modulation" (AM), Hz
returnMS	if FALSE, only roughness is returned (much faster)
returnComplex	if TRUE, returns a complex modulation spectrum (without normalization and warping)
summaryFun	functions used to summarize each acoustic characteristic, eg <code>c('mean', 'sd')</code> ; user-defined functions are fine (see examples); NAs are omitted automatically for mean/median/sd/min/max/range/sum, otherwise take care of NAs yourself
averageMS	if TRUE, the modulation spectra of all inputs are averaged into a single output; if FALSE, a separate MS is returned for each input
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
plot	if TRUE, plots the modulation spectrum of each sound
savePlots	if a valid path is specified, a plot is saved in this folder (defaults to NA)
logWarpX, logWarpY	numeric vector of length 2: <code>c(sigma, base)</code> of pseudolog-warping the modulation spectrum, as in function <code>pseudo_log_trans()</code> from the "scales" package
quantiles	labeled contour values, % (e.g., "50" marks regions that contain 50% of the sum total of the entire modulation spectrum)
kernelSize	the size of Gaussian kernel used for smoothing (1 = no smoothing)
kernelSD	the SD of Gaussian kernel used for smoothing, relative to its size
colorTheme	black and white (<code>'bw'</code>), as in <code>seewave</code> package (<code>'seewave'</code>), or any palette from palette such as <code>'heat.colors'</code> , <code>'cm.colors'</code> , etc
col	actual colors, eg <code>rev(rainbow(100))</code> - see <code>?hcl.colors</code> for colors in base R (overrides <code>colorTheme</code>)
xlab, ylab, main, xlim, ylim	graphical parameters
width, height, units, res	parameters passed to <code>png</code> if the plot is saved
...	other graphical parameters passed on to <code>filled.contour.mod</code> and <code>contour</code> (see spectrogram)

Details

Algorithm: prepare a spectrogram, take its logarithm (if `logSpec = TRUE`), center, perform a 2D Fourier transform (see also `spectral::spec.fft()`), take the upper half of the resulting symmetric matrix, and raise it to power. The result is returned as `$original`. For plotting purposes, the modulation matrix can be smoothed with Gaussian blur (see [gaussianSmooth2D](#)) and log-warped (if `logWarp` is a positive number). This processed modulation spectrum is returned as `$processed`. If the audio is long enough, multiple windows are analyzed, resulting in a vector of roughness values. For multiple inputs, such as a list of waveforms or path to a folder with audio files, the ensemble of modulation spectra can be interpolated to the same spectral and temporal resolution and averaged (if `averageMS`).

Value

Returns a list with the following components:

- `$original` modulation spectrum prior to blurring and log-warping, but after squaring if `power = TRUE`, a matrix of nonnegative values. Rownames are spectral modulation frequencies (cycles/KHz), and colnames are temporal modulation frequencies (Hz).
- `$processed` modulation spectrum after blurring and log-warping
- `$complex` untransformed complex modulation spectrum (returned only if `returnComplex = TRUE`)
- `$roughness` proportion of energy / amplitude of the modulation spectrum within `roughRange` of temporal modulation frequencies, % - a vector if `amRes` is numeric and the sound is long enough, a single number otherwise
- `$amMsFreq` frequency of the highest peak, within `amRange`, of the folded AM function (average AM across all FM bins for both negative and positive AM frequencies), where a peak is a local maximum over `amRes` Hz. Like `roughness`, `amMsFreq` and `amMsPurity` can be single numbers or vectors, depending on whether the sound is analyzed as a whole or in chunks
- `$amMsPurity` ratio of the peak at `amMsFreq` to the median AM over `amRange`, dB
- `$summary` dataframe with summaries of `roughness`, `amMsFreq`, and `amMsPurity`

References

- Singh, N. C., & Theunissen, F. E. (2003). Modulation spectra of natural sounds and ethological theories of auditory processing. *The Journal of the Acoustical Society of America*, 114(6), 3394-3411.

See Also

[plotMS spectrogram analyze](#)

Examples

```
# White noise
ms = modulationSpectrum(runif(16000), samplingRate = 16000,
  logSpec = FALSE, power = TRUE,
  amRes = NULL) # analyze the entire sound, giving a single roughness value
str(ms)
```

```

# Harmonic sound
s = soundgen(amFreq = 25, amDep = 50)
ms = modulationSpectrum(s, samplingRate = 16000, amRes = NULL)
ms[c('roughness', 'amMsFreq', 'amMsPurity')] # a single value for each
ms1 = modulationSpectrum(s, samplingRate = 16000, amRes = 5)
ms1[c('roughness', 'amMsFreq', 'amMsPurity')]
# measured over time (low values of amRes mean more precision, so we analyze
# longer segments and get fewer values per sound)

# Embellish
ms = modulationSpectrum(s, samplingRate = 16000,
  xlab = 'Temporal modulation, Hz', ylab = 'Spectral modulation, 1/KHz',
  colorTheme = 'heat.colors', main = 'Modulation spectrum', lty = 3)

## Not run:
# A long sound with varying AM and a bit of chaos at the end
s_long = soundgen(syllLen = 3500, pitch = c(250, 320, 280),
  amFreq = c(30, 55), amDep = c(20, 60, 40),
  jitterDep = c(0, 0, 2))
playme(s_long)
ms = modulationSpectrum(s_long, 16000)
# plot AM over time
plot(x = seq(1, 1500, length.out = length(ms$amMsFreq)), y = ms$amMsFreq,
  cex = 10^(ms$amMsPurity/20) * 10, xlab = 'Time, ms', ylab = 'AM frequency, Hz')
# plot roughness over time
spectrogram(s_long, 16000, ylim = c(0, 4),
  extraContour = list(ms$roughness / max(ms$roughness) * 4000, col = 'blue'))

# As with spectrograms, there is a tradeoff in time-frequency resolution
s = soundgen(pitch = 500, amFreq = 50, amDep = 100,
  samplingRate = 44100, plot = TRUE)
# playme(s, samplingRate = 44100)
ms = modulationSpectrum(s, samplingRate = 44100,
  windowLength = 50, step = 50, amRes = NULL) # poor temporal resolution
ms = modulationSpectrum(s, samplingRate = 44100,
  windowLength = 5, step = 1, amRes = NULL) # poor frequency resolution
ms = modulationSpectrum(s, samplingRate = 44100,
  windowLength = 15, step = 3, amRes = NULL) # a reasonable compromise

# customize the plot
ms = modulationSpectrum(s, samplingRate = 44100,
  windowLength = 15, overlap = 80, amRes = NULL,
  kernelSize = 17, # more smoothing
  xlim = c(-70, 70), ylim = c(0, 4), # zoom in on the central region
  quantiles = c(.25, .5, .8), # customize contour lines
  col = rev(rainbow(100)), # alternative palette
  power = 2) # ^2
# Note the peaks at FM = 2/KHz (from "pitch = 500") and AM = 50 Hz (from
# "amFreq = 50")

# Input can be a wav/mp3 file
ms = modulationSpectrum('~Downloads/temp/16002_Faking_It_Large_clear.wav')

```

```

# Input can be path to folder with audio files. Each file is processed
# separately, and the output can contain an MS per file...
ms1 = modulationSpectrum('~Downloads/temp', kernelSize = 11,
                        plot = FALSE, averageMS = FALSE)

ms1$summary
names(ms1$original) # a separate MS per file
# ...or a single MS can be calculated:
ms2 = modulationSpectrum('~Downloads/temp', kernelSize = 11,
                        plot = FALSE, averageMS = TRUE)

plotMS(ms2$original)
ms2$summary

# Input can also be a list of waveforms (numeric vectors)
ss = vector('list', 10)
for (i in 1:length(ss)) {
  ss[[i]] = soundgen(syllLen = runif(1, 100, 1000), temperature = .4,
                    pitch = runif(3, 400, 600))
}
# lapply(ss, playme)
# MS of the first sound
ms1 = modulationSpectrum(ss[[1]], samplingRate = 16000, scale = 1)
# average MS of all 10 sounds
ms2 = modulationSpectrum(ss, samplingRate = 16000, scale = 1, averageMS = TRUE, plot = FALSE)
plotMS(ms2$original)

# A sound with ~3 syllables per second and only downsweeps in F0 contour
s = soundgen(nSyl = 8, syllLen = 200, pauseLen = 100, pitch = c(300, 200))
# playme(s)
ms = modulationSpectrum(s, samplingRate = 16000, maxDur = .5,
                        xlim = c(-25, 25), colorTheme = 'seewave',
                        power = 2)
# note the asymmetry b/c of downsweeps

# "power = 2" returns squared modulation spectrum - note that this affects
# the roughness measure!
ms$roughness
# compare:
modulationSpectrum(s, samplingRate = 16000, maxDur = .5,
                  xlim = c(-25, 25), colorTheme = 'seewave',
                  power = 1)$roughness # much higher roughness

# Plotting with or without log-warping the modulation spectrum:
ms = modulationSpectrum(soundgen(), samplingRate = 16000, plot = TRUE)
ms = modulationSpectrum(soundgen(), samplingRate = 16000,
                        logWarpX = c(2, 2), plot = TRUE)

# logWarp and kernelSize have no effect on roughness
# because it is calculated before these transforms:
modulationSpectrum(s, samplingRate = 16000, logWarpX = c(1, 10))$roughness
modulationSpectrum(s, samplingRate = 16000, logWarpX = NA)$roughness
modulationSpectrum(s, samplingRate = 16000, kernelSize = 17)$roughness

```



```

# Log-transform the spectrogram prior to 2D FFT (affects roughness):
modulationSpectrum(s, samplingRate = 16000, logSpec = FALSE)$roughness
modulationSpectrum(s, samplingRate = 16000, logSpec = TRUE)$roughness

# Complex modulation spectrum with phase preserved
ms = modulationSpectrum(soundgen(), samplingRate = 16000,
                        returnComplex = TRUE)
plotMS(abs(ms$complex)) # note the symmetry
# compare:
plotMS(ms$original)

## End(Not run)

```

morph

Morph sounds

Description

Takes two formulas for synthesizing two target sounds with [soundgen](#) and produces a number of intermediate forms (morphs), attempting to go from one target sound to the other in a specified number of equal steps. Normally you will want to set temperature very low; the `tempEffects` argument is not supported.

Usage

```

morph(
  formula1,
  formula2,
  nMorphs,
  playMorphs = TRUE,
  savePath = NA,
  samplingRate = 16000
)

```

Arguments

<code>formula1, formula2</code>	lists of parameters for calling soundgen that produce the two target sounds between which morphing will occur. Character strings containing the full call to <code>soundgen</code> are also accepted (see examples)
<code>nMorphs</code>	the number of morphs to produce, including target sounds
<code>playMorphs</code>	if TRUE, the morphs will be played
<code>savePath</code>	if it is the path to an existing directory, morphs will be saved there as individual .wav files (defaults to NA)
<code>samplingRate</code>	sampling rate of output, Hz. NB: overrides the values in <code>formula1</code> and <code>formula2</code>

Value

A list of two sublists (`$formulas` and `$sounds`), each of length `nMorphs`. For ex., the formula for the second hybrid is `m$formulas[[2]]`, and the waveform is `m$sounds[[2]]`

See Also

[soundgen](#)

Examples

```
## Not run:
# write two formulas or copy-paste them from soundgen_app() or presets:
playback = c(TRUE, FALSE)[2]
# [a] to barking
m = morph(formula1 = list(repeatBout = 2),
          # equivalently: formula1 = 'soundgen(repeatBout = 2)',
          formula2 = presets$Misc$Dog_bark,
          nMorphs = 5, playMorphs = playback)
# use $formulas to access formulas for each morph, $sounds for waveforms
# m$formulas[[4]]
# playme(m$sounds[[3]])

# morph intonation and vowel quality
m = morph(
  'soundgen(pitch = c(300, 250, 400),
            formants = c(350, 2900, 3600, 4700))',
  'soundgen(pitch = c(300, 700, 500, 300),
            formants = c(800, 1250, 3100, 4500))',
  nMorphs = 5, playMorphs = playback
)

# from a grunt of disgust to a moan of pleasure
m = morph(
  formula1 = 'soundgen(syllLen = 180, pitch = c(160, 160, 120), rolloff = -12,
                      nonlinBalance = 70, subDep = 15, jitterDep = 2,
                      formants = c(550, 1200, 2100, 4300, 4700, 6500, 7300),
                      noise = data.frame(time = c(0, 180, 270), value = c(-25, -25, -40)),
                      rolloffNoise = 0)',
  formula2 = 'soundgen(syllLen = 320, pitch = c(340, 330, 300),
                      rolloff = c(-18, -16, -30), ampl = c(0, -10), formants = c(950, 1700, 3700),
                      noise = data.frame(time = c(0, 300, 440), value = c(-35, -25, -65)),
                      mouth = c(.4, .5), rolloffNoise = -5, attackLen = 30)',
  nMorphs = 8, playMorphs = playback
)

# from scream_010 to moan_515b
# (see online demos at http://cogsci.se/soundgen/humans/humans.html)
m = morph(
  formula1 = "soundgen(
    syllLen = 490,
    pitch = list(time = c(0, 80, 250, 370, 490),
                value = c(1000, 2900, 3200, 2900, 1000)),
```

```

    rolloff = c(-5, 0, -25), rolloffKHz = 0,
    temperature = 0.001,
    jitterDep = c(.5, 1, 0), shimmerDep = c(5, 15, 0),
    formants = c(1100, 2300, 3100, 4000, 5300, 6200),
    mouth = c(.3, .5, .6, .5, .3))",
formula2 = "soundgen(syllLen = 520,
    pitch = c(300, 310, 300),
    ampl = c(0, -30),
    temperature = 0.001, rolloff = c(-18, -25),
    jitterDep = .05, shimmerDep = 2,
    formants = list(f1 = c(700, 900),
        f2 = c(1600, 1400),
        f3 = c(3600, 3500), f4 = c(4300, 4200)),
    mouth = c(.5, .3),
    noise = data.frame(time = c(0, 400, 660),
        value = c(-20, -10, -60)),
    rolloffNoise = c(-5, -15))",
nMorphs = 5, playMorphs = playback
)

## End(Not run)

```

msToSpec

Modulation spectrum to spectrogram

Description

Takes a complex MS and transforms it to a complex spectrogram with proper row (frequency) and column (time) labels.

Usage

```
msToSpec(ms, windowLength = NULL, step = NULL)
```

Arguments

ms	target modulation spectrum (matrix of complex numbers)
windowLength	length of FFT window, ms
step	you can override overlap by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of 1/samplingRate, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)

Value

Returns a spectrogram - a numeric matrix of complex numbers of the same dimensions as ms.

Examples

```
s = soundgen(syllLen = 250, amFreq = 25, amDep = 50,
             pitch = 250, samplingRate = 16000)
spec = spectrogram(s, samplingRate = 16000, windowLength = 25, step = 5)
ms = specToMS(spec)
image(x = as.numeric(colnames(ms)), y = as.numeric(rownames(ms)),
      z = t(log(abs(ms))), xlab = 'Amplitude modulation, Hz',
      ylab = 'Frequency modulation, cycles/kHz')
spec_new = msToSpec(ms)
image(x = as.numeric(colnames(spec_new)), y = as.numeric(rownames(spec_new)),
      z = t(log(abs(spec_new))), xlab = 'Time, ms',
      ylab = 'Frequency, kHz')
```

naiveBayes

*Naive Bayes***Description**

An implementation of a Naive Bayes classifier adapted to autocorrelated time series such as the type of nonlinear vocal phenomena in consecutive audio frames. All predictors must be continuous, and the outcome must be categorical. Cases with missing values are not deleted because the posterior probabilities of each outcome class can be calculated from different combinations of predictors on a case-by-case basis. Two optional modifications of a standard Naive Bayes algorithm can be made: (1) classifications can be "clumped" at the final stage, ensuring that every run or "epoch" of a particular predicted class is at least `minLength` steps long, and (2) priors can be continuously adapted based on the likelihood function of the preceding `wlPrior` observations if `prior = 'dynamic'`.

Usage

```
naiveBayes(
  formula,
  train,
  test = train,
  prior = c("flat", "static", "dynamic")[2],
  wlPrior = 3,
  wlClumper = NULL,
  runBack = TRUE,
  plot = FALSE
)
```

Arguments

<code>formula</code>	model formula of the type <code>outcome ~ predictor1 + predictor2 + ...</code> (no interactions)
<code>train</code>	either the training dataframe or the output of <code>naiveBayes_train</code> . This data is used to calculate class-specific distributions of the predictors and prior class probabilities

test	the test dataframe. This data is used to make predictions - that is, outcome class probabilities given the values of predictors
prior	"flat" = all classes are equally likely a prior, "static" = use class probabilities in the training dataset, "dynamic" = update prior probabilities from weighted likelihoods of wLPrior preceding observations
wLPrior	the length of a Gaussian window used for updating dynamic priors
wLClumper	the minimum expected number of observations of the same class before the class can change
runBack	if TRUE, the dynamic prior is calculated both forward and backward and averaged (only has an effect if prior = 'dynamic')
plot	if TRUE, produces diagnostic plots

Value

Returns the test dataframe with new columns: "pr" = the predicted class membership, "[class]" = posterior probabilities per class, "like_[class]" = log-likelihoods, "prior_[class]" = log-priors, "priorF_[class]" / "priorB_[class]" = forward / backward log-priors per class.

Examples

```
set.seed(151)
## create some fake data
df = data.frame(group = rep(c(
  rep('A', 150), rep('B', 50), rep('A', 120),
  rep('A', 100), rep('B', 30), rep('A', 90)
), 3))
df$group = as.factor(df$group)
df$x1 = rnorm(nrow(df), mean = ifelse(df$group == 'A', 3, 6), sd = 2)
df$x2 = rnorm(nrow(df), mean = ifelse(df$group == 'A', 2, -1), sd = 2)
boxplot(x1 ~ group, df)
boxplot(x2 ~ group, df)

## train the classifier
mod_train = naiveBayes_train(group ~ x1 + x2, data = df)
mod_train

## test on new data generated by the same process
test = data.frame(group = rep(c(
  rep('A', 90), rep('B', 40), rep('A', 150),
  rep('B', 40), rep('A', 130), rep('B', 30)
), 2))
test$group = as.factor(test$group)
test$x1 = rnorm(nrow(test), mean = ifelse(test$group == 'A', 3, 6), sd = 2)
test$x2 = rnorm(nrow(test), mean = ifelse(test$group == 'A', 2, -1), sd = 2)

# flat priors (same prior probability for each class)
nb_flat = naiveBayes(group ~ x1 + x2, train = mod_train, test = test,
  prior = 'flat', plot = TRUE)
# same as passing 'train' directly to the model, w/o calling naiveBayes_train():
nb_flat = naiveBayes(group ~ x1 + x2, train = df, test = test, prior = 'flat')
```

```

table(nb_flat$group, nb_flat$pr)
mean(nb_flat$group == nb_flat$pr) # 84% correct

# static priors (use original class proportions as prior class probabilities)
nb_static = naiveBayes(group ~ x1 + x2, train = mod_train, test = test,
  prior = 'static', w1Clumper = NULL, plot = TRUE)
table(nb_static$group, nb_static$pr)
mean(nb_static$group == nb_static$pr) # 87% correct

# specify custom static priors
mod_train2 = mod_train
mod_train$table
mod_train2$table = list(A = .1, B = .9) # sum to 1
nb_static2 = naiveBayes(group ~ x1 + x2, train = mod_train2, test = test,
  prior = 'static', w1Clumper = NULL, plot = TRUE)
mean(nb_static2$group == nb_static2$pr) # 61% correct

# if we expect autocorrelation, ie class X is more likely a priori if the
# last few observations were also likely to be class X, we can use dynamic
# priors and/or clumper the predicted classes (the latter imposes strong
# constraints on the predictions, but may be worth it if the data is known to
# be strongly "clumpered", ie if we know classes occur in long'ish runs)
nb1 = naiveBayes(group ~ x1 + x2, train = mod_train, test = test,
  prior = 'dynamic', w1Prior = 10, plot = TRUE)
table(nb1$group, nb1$pr)
mean(nb1$group == nb1$pr) # 94% correct

nb2 = naiveBayes(group ~ x1 + x2, train = mod_train, test = test,
  prior = 'static', w1Clumper = 10, plot = TRUE)
table(nb2$group, nb2$pr)
mean(nb2$group == nb2$pr) # 89% correct

nb3 = naiveBayes(group ~ x1 + x2, train = mod_train, test = test,
  prior = 'dynamic', w1Prior = 10, w1Clumper = 10, plot = TRUE)
table(nb3$group, nb3$pr)
mean(nb3$group == nb3$pr) # 98% correct

# NAs in the data are not a problem
test1 = test
test1$x1[sample(1:nrow(test1), 100)] = NA
test1$x2[sample(1:nrow(test1), 10)] = NA
summary(test1)

nb4 = naiveBayes(group ~ x1 + x2, train = mod_train, test = test,
  prior = 'dynamic', w1Prior = 10, plot = TRUE)
table(nb4$group, nb4$pr)
mean(nb4$group == nb4$pr) # still 94% correct

```

Description

Returns conditional means and standard deviations per class as well as a table with the global proportions of each class in the dataset. This is mostly useful because the output can be passed on to [naiveBayes](#) to save time if `naiveBayes()` is called in a loop with the same training dataset.

Usage

```
naiveBayes_train(formula, data)
```

Arguments

formula	outcome ~ predictor1 + predictor1 + ...
data	training dataset

 nonlinPred

Nonlinear prediction

Description

Predicts new points in a time series. The functionality is provided by [nonLinearPrediction](#). This function is just a simple wrapper "for dummies" that reconstructs the phase space under the hood, including the choice of time lag, embedding dimensions, etc. It can also predict not one but many points in a single step.

Usage

```
nonlinPred(
  x,
  nPoints = 1,
  time.lag = NULL,
  embedding.dim = NULL,
  max.embedding.dim = 15,
  threshold = 0.95,
  max.relative.change = 0.1,
  radius = NULL,
  radius.increment = NULL,
  plot = FALSE
)
```

Arguments

x	numeric vector
nPoints	number of points to predict, ideally not more than $\text{length}(x) / 2$ (the function is called recursively to predict longer sequences, but don't expect miracles)
time.lag	time lag for constructing Takens vectors. Defaults to the time to the first exponential decay of mutual information. See timeLag

embedding.dim the number of dimensions of the phase space. Defaults to an estimate based on [estimateEmbeddingDim](#)

max.embedding.dim, threshold, max.relative.change parameters used to estimate the optimal number of embedding dimensions - see [estimateEmbeddingDim](#)

radius, radius.increment the radius used for detecting neighbors in the phase space and its increment in case no neighbors are found - see [nonLinearPrediction](#)

plot if TRUE, plots the original time series and the predictions

Value

Returns a numeric vector on the same scale as input x.

Examples

```
x = c(rep(1, 3), rep(0, 4), rep(1, 3), rep(0, 4), rep(1, 3), 0, 0)
nonlinPred(x, 5, plot = TRUE)

nonlinPred(sin(1:25), 22, plot = TRUE)

x = soundgen(syllLen = 50, addSilence = 0)[250:450]
nonlinPred(x, 100, plot = TRUE)

nonlinPred(c(rnorm(5), NA, rnorm(3)))
nonlinPred(1:4)
nonlinPred(1:6)

## Not run:
s1 = soundgen(syllLen = 500, pitch = rnorm(5, 200, 20),
              addSilence = 0, plot = TRUE)

playme(s1)
length(s1)
# we can predict output that is longer than the original time series by
# predicting a bit at a time and using the output as the new input
s2 = nonlinPred(s1, 16000)
spectrogram(c(s1, s2))
playme(c(s1, s2))

## End(Not run)
```

normalizeFolder

Normalize folder

Description

Normalizes the amplitude of all wav/mp3 files in a folder based on their peak or RMS amplitude or subjective loudness. This is good for playback experiments, which require that all sounds should have similar intensity or loudness.

Usage

```
normalizeFolder(
  myfolder,
  type = c("peak", "rms", "loudness")[1],
  maxAmp = 0,
  summaryFun = "mean",
  windowLength = 50,
  step = NULL,
  overlap = 70,
  killDC = FALSE,
  windowDC = 200,
  saveAudio = NULL,
  reportEvery = NULL
)
```

Arguments

myfolder	full path to folder containing input audio files
type	normalize so the output files has the same peak amplitude ('peak'), root mean square amplitude ('rms'), or subjective loudness in sone ('loudness')
maxAmp	maximum amplitude in dB (0 = max possible, -10 = 10 dB below max possible, etc.)
summaryFun	should the output files have the same mean / median / max etc rms amplitude or loudness? (summaryFun has no effect if type = 'peak')
windowLength	length of FFT window, ms
step	you can override overlap by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of 1/samplingRate, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
overlap	overlap between successive FFT frames, %
killDC	if TRUE, removed DC offset (see also flatEnv)
windowDC	the window for calculating DC offset, ms
saveAudio	full path to where the normalized files should be saved (defaults to 'myfolder/normalized')
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)

Details

Algorithm: first all files are rescaled to have the same peak amplitude of maxAmp dB. If type = 'peak', the process ends here. If type = 'rms', there are two additional steps. First the original RMS amplitude of all files is calculated per frame by [getRMS](#). The "quietest" sound with the lowest summary RMS value is not modified, so its peak amplitude remains maxAmp dB. All the remaining sounds are rescaled linearly, so that their summary RMS values becomes the same as that of the "quietest" sound, and their peak amplitudes become smaller, <maxAmp. Finally, if type = 'loudness', the subjective loudness of each sound is estimated by [getLoudness](#), which assumes frequency sensitivity typical of human hearing. The following normalization procedure is similar to that for type = 'rms'.

See Also

[getRMS](#) [analyze](#) [getLoudness](#)

Examples

```
## Not run:
# put a few short audio files in a folder, eg '~/Downloads/temp'
getRMS '~/Downloads/temp2', summaryFun = 'mean')$summary # different
normalizeFolder '~/Downloads/temp2', type = 'rms', summaryFun = 'mean',
  saveAudio = '~/Downloads/temp2/normalized')
getRMS '~/Downloads/temp2/normalized', summaryFun = 'mean')$summary # same
# If the saved audio files are treated as stereo with one channel missing,
# try reconverting with ffmpeg (saving is handled by tuneR::writeWave)

## End(Not run)
```

notesDict	<i>Conversion table from Hz to musical notation</i>
-----------	---

Description

A dataframe of 192 rows and 2 columns: "note" and "freq" (Hz). Range: C-5 (0.51 Hz) to B10 (31608.53 Hz)

Usage

```
notesDict
```

Format

An object of class `data.frame` with 192 rows and 2 columns.

notesToHz	<i>Convert notes to Hz</i>
-----------	----------------------------

Description

Converts to Hz from musical notation like A4 - note A of the fourth octave above C0 (16.35 Hz).

Usage

```
notesToHz(n, A4 = 440)
```

Arguments

n	vector or matrix of notes
A4	frequency of note A in the fourth octave (modern standard ISO 16 or concert pitch = 440 Hz)

See Also

[HzToNotes](#) [HzToSemitones](#)

Examples

```
notesToHz(c("A4", "D4", "A#2", "C0", "C-2"))

# Baroque tuning A415, half a semitone flat relative to concert pitch A440
notesToHz(c("A4", "D4", "A#2", "C0", "C-2"), A4 = 415)
```

optimizePars

Optimize parameters for acoustic analysis

Description

This customized wrapper for `optim` attempts to optimize the parameters of `segment` or `analyze` by comparing the results with a manually annotated "key". This optimization function uses a single measurement per audio file (e.g., median pitch or the number of syllables). For other purposes, you may want to adapt the optimization function so that the key specifies the exact timing of syllables, their median length, frame-by-frame pitch values, or any other characteristic that you want to optimize for. The general idea remains the same, however: we want to tune function parameters to fit our type of audio and research priorities. The default settings of `segment` and `analyze` have been optimized for human non-linguistic vocalizations.

Usage

```
optimizePars(
  myfolder,
  key,
  myfun,
  pars,
  bounds = NULL,
  fitnessPar,
  fitnessFun = function(x) 1 - cor(x, key, use = "pairwise.complete.obs"),
  nIter = 10,
  init = NULL,
  initSD = 0.2,
  control = list(maxit = 50, reltol = 0.01, trace = 0),
  otherPars = list(plot = FALSE),
  mygrid = NULL,
  verbose = TRUE
)
```

Arguments

`myfolder` path to where the .wav files live
`key` a vector containing the "correct" measurement that we are aiming to reproduce

myfun	the function being optimized: either 'segment' or 'analyze' (in quotes)
pars	names of arguments to myfun that should be optimized
bounds	a list setting the lower and upper boundaries for possible values of optimized parameters. For ex., if we optimize smooth and smoothOverlap, reasonable bounds might be list(low = c(5, 0), high = c(500, 95))
fitnessPar	the name of output variable that we are comparing with the key, e.g. 'nBursts' or 'pitch_median'
fitnessFun	the function used to evaluate how well the output of myfun fits the key. Defaults to 1 - Pearson's correlation (i.e. 0 is perfect fit, 1 is awful fit). For pitch, log scale is more meaningful, so a good fitness criterion is "function(x) 1 - cor(log(x), log(key), use = 'pairwise.complete.obs')"
nIter	repeat the optimization several times to check convergence
init	initial values of optimized parameters (if NULL, the default values are taken from the definition of myfun)
initSD	each optimization begins with a random seed, and initSD specifies the SD of normal distribution used to generate random deviation of initial values from the defaults
control	a list of control parameters passed on to <code>optim</code> . The method used is "Nelder-Mead"
otherPars	a list of additional arguments to myfun
mygrid	a dataframe with one column per parameter to optimize, with each row specifying the values to try. If not NULL, optimizePars simply evaluates each combination of parameter values, without calling <code>optim</code> (see examples)
verbose	if TRUE, reports the values of parameters evaluated and fitness

Details

If your sounds are very different from human non-linguistic vocalizations, you may want to change the default values of other arguments to speed up convergence. Adapt the code to enforce suitable constraints, depending on your data.

Value

Returns a matrix with one row per iteration with fitness in the first column and the best values of each of the optimized parameters in the remaining columns.

Examples

```
## Not run:
# Download 260 sounds from the supplements in Anikin & Persson (2017)
# - see http://cogsci.se/publications.html
# Unzip them into a folder, say '~/Downloads/temp'
myfolder = '~/Downloads/temp260' # 260 .wav files live here

# Optimization of SEGMENTATION
# Import manual counts of syllables in 260 sounds from
```

```

# Anikin & Persson (2017) (our "key")
key = segmentManual # a vector of 260 integers

# Run optimization loop several times with random initial values
# to check convergence
# NB: with 260 sounds and default settings, this might take ~20 min per iteration!
res = optimizePars(myfolder = myfolder, myfun = 'segment', key = key,
  pars = c('shortestSyl', 'shortestPause'),
  fitnessPar = 'nBursts', otherPars = list(method = 'env'),
  nIter = 3, control = list(maxit = 50, reltol = .01, trace = 0))

# Examine the results
print(res)
for (c in 2:ncol(res)) {
  plot(res[, c], res[, 1], main = colnames(res)[c])
}
pars = as.list(res[1, 2:ncol(res)]) # top candidate (best pars)
s = do.call(segment, c(myfolder, pars)) # segment with best pars
cor(key, as.numeric(s[, fitnessPar]))
boxplot(as.numeric(s[, fitnessPar]) ~ as.integer(key), xlab='key')
abline(a=0, b=1, col='red')

# Try a grid with particular parameter values instead of formal optimization
res = optimizePars(myfolder = myfolder, myfun = 'segment', key = segmentManual,
  pars = c('shortestSyl', 'shortestPause'),
  fitnessPar = 'nBursts', otherPars = list(method = 'env'),
  mygrid = expand.grid(shortestSyl = c(30, 40),
    shortestPause = c(30, 40, 50)))
1 - res$fit # correlations with key

# Optimization of PITCH TRACKING (takes several hours!)
key = as.numeric(log(pitchManual))
res = optimizePars(
  myfolder = myfolder,
  myfun = 'analyze',
  key = key, # log-scale better for pitch
  pars = c('windowLength', 'silence'),
  bounds = list(low = c(5, 0), high = c(200, .2)),
  fitnessPar = 'pitch_median',
  nIter = 2,
  otherPars = list(plot = FALSE, loudness = NULL, novelty = NULL,
    roughness = NULL, nFormants = 0),
  fitnessFun = function(x) {
    1 - cor(log(x), key, use = 'pairwise.complete.obs') *
    (1 - mean(is.na(x) & is.finite(key))) # penalize failing to detect f0
  })

## End(Not run)

```

Description

Plots the oscillogram (waveform) of a sound on a linear or logarithmic scale (in dB). To get a dB scale, centers and normalizes the sound, then takes a logarithm of the positive part and a flipped negative part, which is analogous to "Waveform (dB)" view in Audacity. For more plotting options, check [oscillo](#).

Usage

```
osc(
  x,
  samplingRate = NULL,
  scale = NULL,
  from = NULL,
  to = NULL,
  dynamicRange = 80,
  dB = FALSE,
  returnWave = FALSE,
  reportEvery = NULL,
  cores = 1,
  plot = TRUE,
  savePlots = NULL,
  main = NULL,
  xlab = NULL,
  ylab = NULL,
  ylim = NULL,
  bty = "n",
  midline = TRUE,
  maxPoints = 10000,
  width = 900,
  height = 500,
  units = "px",
  res = NA,
  ...
)
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>scale</code>	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
<code>from</code> , <code>to</code>	if <code>NULL</code> (default), analyzes the whole sound, otherwise from...to (s)
<code>dynamicRange</code>	dynamic range, dB. All values more than one <code>dynamicRange</code> under maximum are treated as zero
<code>dB</code>	if <code>TRUE</code> , plots on a dB instead of linear scale
<code>returnWave</code>	if <code>TRUE</code> , returns a log-transformed waveform as a numeric vector

reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
plot	if TRUE, plots the oscillogram
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
main	plot title
xlab, ylab	axis labels
ylim	override default amplitude scale for non-centered sounds
bty	box type (see '?par')
midline	if TRUE, draws a line at 0 dB
maxPoints	the maximum number of points to plot (speeds up the plotting of long audio files, but beware of antialiasing)
width, height, units, res	graphical parameters for saving plots passed to png
...	Other graphical parameters passed on to 'plot()'

Value

If `returnWave = TRUE`, returns the input waveform on the original or dB scale: a vector with range from `'-dynamicRange'` to `'dynamicRange'`.

Examples

```

sound = sin(1:2000/10) *
  getSmoothContour(anchors = c(1, .01, .5), len = 2000)

# Oscillogram on a linear scale without bells and whistles, just base R
plot(sound, type = 'l')

# Oscillogram options with soundgen
osc(sound)           # linear
osc(sound, dB = TRUE) # dB

# For numeric vectors, indicate samplingRate and scale (max amplitude)
osc(sound, samplingRate = 1000, scale = 100, dB = TRUE)

# Embellish and customize the plot
o = osc(sound, samplingRate = 1000, dB = TRUE, midline = FALSE,
        main = 'My waveform', col = 'blue', returnWave = TRUE)
abline(h = -80, col = 'orange', lty = 3)
o[1:10] # the waveform in dB

## Not run:
# Wave object
data(sheep, package = 'seewave')
osc(sheep, dB = TRUE)

```

```

# Plot a section
osc(sheep, from = .5, to = 1.2)

# for long files, reduce the resolution to plot quickly (careful: if the
# resolution is too low, antialiasing may cause artifacts)
osc(sheep, dB = TRUE, maxPoints = 2500)
osc(sheep, samplingRate = 5000, maxPoints = 100)

# files several minutes long can be plotted in under a second
osc('~Downloads/speechEx.wav', maxPoints = 20000)

# saves oscillograms of all audio files in a folder
osc('~Downloads/temp2', savePlots = '')

## End(Not run)

```

permittedValues

Defaults and ranges for soundgen()

Description

A dataset containing defaults and ranges of key variables for `soundgen()` and `soundgen_app()`. Adjust as needed.

Usage

```
permittedValues
```

Format

A matrix with 58 rows and 4 columns:

default default value

low lowest permitted value

high highest permitted value

step increment for adjustment ...

`phasegram`*Phasegram*

Description

Produces a phasegram of a sound or another time series, which is a collection of Poincare sections cut through phase portraits of consecutive frames. The x axis is time, just as in a spectrogram, the y axis is a slice through the phase portrait, and the color shows the density of trajectories at each point of the phase portrait.

Usage

```
phasegram(  
  x,  
  samplingRate = NULL,  
  from = NULL,  
  to = NULL,  
  windowLength = 10,  
  step = windowLength/2,  
  timeLag = NULL,  
  theilerWindow = NULL,  
  nonlinStats = c("ed", "d2", "ml", "sur"),  
  pars_ed = list(max.embedding.dim = 15),  
  pars_d2 = list(min.embedding.dim = 2, min.radius = 0.001, n.points.radius = 20),  
  pars_ml = list(min.embedding.dim = 2, radius = 0.001),  
  pars_sur = list(FUN = nonlinearTseries::timeAsymmetry, K = 1),  
  bw = 0.01,  
  bins = 5/bw,  
  reportEvery = NULL,  
  cores = 1,  
  rasterize = FALSE,  
  plot = TRUE,  
  savePlots = NULL,  
  colorTheme = c("bw", "seewave", "heat.colors", "...")[1],  
  col = NULL,  
  xlab = "Time",  
  ylab = "",  
  main = NULL,  
  width = 900,  
  height = 500,  
  units = "px",  
  res = NA,  
  ...  
)
```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
from, to	if NULL (default), analyzes the whole sound, otherwise from...to (s)
windowLength	the length of each frame analyzed separately (ms)
step	time step between consecutive frames (ms)
timeLag	time lag between the original and time-shifted version of each frame that together represent the phase portrait (ms). Defaults to the number of steps beyond which the mutual information function reaches its minimum or, if that fails, the steps until mutual information experiences the first exponential decay - see timeLag
theilerWindow	time lag between two points that are considered locally independent and can be treated as neighbors in the reconstructed phase space. defaults to the first minimum or, if unavailable, the first zero of the autocorrelation function (or, failing that, to timeLag * 2)
nonlinStats	nonlinear statistics to report: "ed" = the optimal number of embedding dimensions, "d2" = correlation dimension D2, "ml" = maximum Lyapunov exponent, "sur" = the results of surrogate data testing for stochasticity. These are calculated using the functionality of the package nonlinearTseries, which is seriously slow, so the default is just to get the phasegram itself
pars_ed	a list of control parameters passed to estimateEmbeddingDim
pars_d2	a list of control parameters passed to corrDim
pars_ml	a list of control parameters passed to maxLyapunov
pars_sur	a list of control parameters passed to surrogateTest
bw	standard deviation of the smoothing kernel, as in density
bins	the number of bins along the Y axis after rasterizing (has no effect if rasterize = FALSE)
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
rasterize	if FALSE, only plots and returns Poincare sections on the original scale (most graphical parameters will then have no effect); if TRUE, rasterizes the phasegram matrix and plots it with more graphical parameters
plot	should a spectrogram be plotted? TRUE / FALSE
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
colorTheme	black and white ('bw'), as in seewave package ('seewave'), or any palette from palette such as 'heat.colors', 'cm.colors', etc
col	actual colors, eg rev(rainbow(100)) - see ?hcl.colors for colors in base R (overrides colorTheme)

```

xlab, ylab, main
    graphical parameters passed to soundgen:::filled.contour.mod (if rasterize =
    TRUE) or plot (if rasterize = FALSE)
width, height, units, res
    graphical parameters for saving plots passed to png
...
    other graphical parameters passed to soundgen:::filled.contour.mod (if rasterize
    = TRUE) or plot (if rasterize = FALSE)

```

Details

Algorithm: the input sound is normalized to $[-1, 1]$ and divided into consecutive frames `windowLength` ms long without multiplying by any windowing function (unlike in STFT). For each frame, a phase portrait is obtained by time-shifting the frame by `timeLag` ms. A Poincare section is taken through the phase portrait (currently at a fixed angle, namely the default in `poincareMap`), giving the intersection points of trajectories with this bisecting line. The density of intersections is estimated with a smoothing kernel of bandwidth `bw` (as an alternative to using histogram bins). The density distributions per frame are stacked together into a phasegram (output: "orig"). The ranges of phase portraits depend on the amplitude of signal in each frame. The resulting phasegram can optionally be rasterized to smooth it for plotting (output: "rasterized").

Value

Returns a list of three components: "orig" = the full phasegram; "rasterized" = a rasterized version. For both, `$time` is the middle of each frame (ms), `$x` is the coordinate along a Poincare section (since the audio is normalized, the scale is $[-1, 1]$), and `$y` is the density of intersections of system trajectories with the Poincare section. The third component is `$descriptives`, which gives the result of nonlinear analysis per frame. Currently implemented: `shannon` = Shannon entropy of Poincare sections, `nPeaks` = log-number of peaks in the density distribution of Poincare sections, `ml` = maximum Lyapunov exponent (positive values suggest chaos), `ed` = optimal number of embedding dimensions (shows the complexity of the reconstructed attractor), `d2` = correlation dimension, `sur` = probability of stochasticity according to surrogate data testing (0 = deterministic, 1 = stochastic).

References

- Herbst, C. T., Herzel, H., Švec, J. G., Wyman, M. T., & Fitch, W. T. (2013). Visualization of system dynamics using phasegrams. *Journal of the Royal Society Interface*, 10(85), 20130288.
- Huffaker, R., Huffaker, R. G., Bittelli, M., & Rosa, R. (2017). *Nonlinear time series analysis with R*. Oxford University Press.

Examples

```

target = soundgen(syllLen = 300, pitch = c(350, 420, 420, 410, 340) * 3,
  subDep = c(0, 0, 60, 50, 0, 0) / 2, addSilence = 0, plot = TRUE)
# Nonlinear statistics are also returned (slow - disable by setting
# nonlinStats = NULL if these are not needed)
ph = phasegram(target, 16000, nonlinStats = NULL)

## Not run:
ph = phasegram(target, 16000, windowLength = 20, step = 20,

```

```

    rasterize = TRUE, bw = .01, bins = 150)
ph$descriptives

# Unfortunately, phasegrams are greatly affected by noise. Compare:
target2 = soundgen(syllLen = 300, pitch = c(350, 420, 420, 410, 340) * 3,
  subDep = c(0, 0, 60, 50, 0, 0)/2, noise = -10, addSilence = 0, plot = TRUE)
ph2 = phasegram(target2, 16000)

s2 = soundgen(syllLen = 3000, addSilence = 0, temperature = 1e-6,
  pitch = c(380, 550, 500, 220), subDep = c(0, 0, 40, 0, 0, 0, 0, 0),
  amDep = c(0, 0, 0, 0, 80, 0, 0, 0), amFreq = 80,
  jitterDep = c(0, 0, 0, 0, 0, 3))
spectrogram(s2, 16000, yScale = 'bark')
phasegram(s2, 16000, windowLength = 10, nonlinStats = NULL, bw = .001)
phasegram(s2, 16000, windowLength = 10, nonlinStats = NULL, bw = .02)

## End(Not run)

```

pitchContour

Manually corrected pitch contours in 260 sounds

Description

A dataframe of 260 rows and two columns: "file" for filename in the corpus (Anikin & Persson, 2017) and "pitch" for pitch values per frame. The corpus can be downloaded from <http://cogsci.se/publications.html>

Usage

```
pitchContour
```

Format

An object of class `data.frame` with 260 rows and 2 columns.

pitchDescriptives

Pitch descriptives

Description

Provides common descriptives of time series such as pitch contours, including measures of average / range / variability / slope / inflections etc. Several degrees of smoothing can be applied consecutively. The summaries are produced on the original and log-transformed scales, so this is meant to be used on frequency-related variables in Hz.

Usage

```
pitchDescriptives(
  x,
  step = NULL,
  timeUnit,
  smoothBW = c(NA, 10, 1),
  inflThres = 0.2,
  extraSummaryFun = c(),
  ref = 16.35,
  plot = FALSE
)
```

Arguments

x	input: numeric vector, a list of time stamps and values in rows, a dataframe with one row per file and time/pitch values stored as characters (as exported by pitch_app), or path to csv file containing the output of pitch_app or analyze
step	distance between values in s (only needed if input is a vector)
timeUnit	specify whether the time stamps (if any) are in ms or s
smoothBW	a vector of bandwidths (Hz) for consecutive smoothing of input using pitchSmoothPraat ; NA = no smoothing
inflThres	minimum difference (in semitones) between consecutive extrema to consider them inflections; to apply a different threshold at each smoothing level, provide inflThres as a vector of the same length as smoothBW; NA = no threshold
extraSummaryFun	additional summary function(s) that take a numeric vector with some NAs and return a single number, eg c('myFun1', 'myFun2')
ref	reference value for transforming Hz to semitones, defaults to C0 (16.35 Hz)
plot	if TRUE, plots the inflections for manual verification

Value

Returns a dataframe with columns containing summaries of one or multiple inputs (one input per row). The descriptives are as follows:

duration total duration, s

durDefined duration after omitting leading and trailing NAs

propDefined proportion of input with non-NA value, eg proportion of voiced frames if the input is pitch

start, start_oct, end, end_oct the first and last values on the original scale and in octaves above C0 (16.3516 Hz)

mean, median, max, min average and extreme values on the original scale

mean_oct, median_oct, min_oct, max_oct same in octaves above C0

time_max, time_min the location of minimum and maximum relative to durDefined, 0 to 1

range, range_sem, sd, sd_sem range and standard deviation on the original scale and in semitones

CV coefficient of variation = $sd/mean$ (provided for historical reasons)

meanSlope, meanSlope_sem mean slope in Hz/s or semitones/s (NB: does not depend on duration or missing values)

meanAbsSlope, meanAbsSlope_sem mean absolute slope (modulus, ie rising and falling sections no longer cancel out)

maxAbsSlope, maxAbsSlope_sem the steepest slope

Examples

```
x = c(NA, NA, 405, 441, 459, 459, 460, 462, 462, 458, 458, 445, 458, 451,
444, 444, 430, 416, 409, 403, 403, 389, 375, NA, NA, NA, NA, NA, NA, NA,
NA, 183, 677, 677, 846, 883, 886, 924, 938, 883, 946, 846, 911, 826, 826,
788, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, 307,
307, 368, 377, 383, 383, 383, 380, 377, 377, 377, 374, 374, 375, 375, 375,
375, 368, 371, 374, 375, 361, 375, 389, 375, 375, 375, 375, 375, 314, 169,
NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, 238, 285, 361, 374, 375, 375,
375, 375, 375, 389, 403, 389, 389, 375, 375, 389, 375, 348, 361, 375, 348,
348, 361, 348, 342, 361, 361, 361, 365, 365, 361, 966, 966, 966, 959, 959,
946, 1021, 1021, 1026, 1086, 1131, 1131, 1146, 1130, 1172, 1240, 1172, 1117,
1103, 1026, 1026, 966, 919, 946, 882, 832, NA, NA, NA, NA, NA, NA, NA, NA,
NA, NA)
plot(x, type = 'b')
ci95 = function(x) diff(quantile(na.omit(x), probs = c(.025, .975)))
pd = pitchDescriptives(
  x, step = .025, timeUnit = 's',
  smoothBW = c(NA, 10, 1), # original + smoothed at 10 Hz and 1 Hz
  inflThres = c(NA, .2, .2), # different for each level of smoothing
  extraSummaryFun = 'ci95', # user-defined, here 95% coverage interval
  plot = TRUE
)
pd

## Not run:
# a single file
data(sheep, package = 'seewave')
a = analyze(sheep)
pd1 = pitchDescriptives(a$detailed[, c('time', 'pitch')],
  timeUnit = 'ms', inflThres = NA, plot = TRUE)
pd2 = pitchDescriptives(a$detailed[, c('time', 'pitch')],
  timeUnit = 'ms', inflThres = c(0.1, 0.1, .5), plot = TRUE)

# multiple files returned by analyze()
an = analyze('~Downloads/temp')
pd = pitchDescriptives(an$detailed, timeUnit = 'ms')
pd

# multiple files returned by pitch_app()
pd = pitchDescriptives(
  '~Downloads/pitch_manual_1708.csv',
  timeUnit = 'ms', smoothBW = c(NA, 2), inflThres = .25)
```

```
# a single file, exported from Praat
par(mfrow = c(3, 1))
pd = pitchDescriptives(
  '~/Downloads/F-Hin-0m_jana.wav_F0contour.txt',
  timeUnit = 's', smoothBW = c(NA, 25, 2), inflThres = .25, plot = TRUE)
par(mfrow = c(1, 1))

## End(Not run)
```

pitchManual

*Manual pitch estimation in 260 sounds***Description**

A vector of manually verified pitch values per sound in the corpus of 590 human non-linguistic emotional vocalizations from Anikin & Persson (2017). The corpus can be downloaded from <http://cogsci.se/publications.html>

Usage

```
pitchManual
```

Format

An object of class numeric of length 260.

pitchSmoothPraat

*Pitch smoothing as in Praat***Description**

Smooths an intonation (pitch) contour with a low-pass filter, as in Praat (<http://www.fon.hum.uva.nl/praat/>). Algorithm: interpolates missing values (unvoiced frames), performs FFT to obtain the spectrum, multiplies by a Gaussian filter, performs an inverse FFT, and fills the missing values back in. The bandwidth parameter is about half the cutoff frequency (ie some frequencies will still be present up to $\sim 2 * \text{bandwidth}$)

Usage

```
pitchSmoothPraat(pitch, bandwidth, samplingRate, plot = FALSE)
```

Arguments

pitch	numeric vector of pitch values (NA = unvoiced)
bandwidth	the bandwidth of low-pass filter, Hz (high = less smoothing, close to zero = more smoothing)
samplingRate	the number of pitch values per second
plot	if TRUE, plots the original and smoothed pitch contours

See Also[analyze](#)**Examples**

```
pitch = c(NA, NA, 405, 441, 459, 459, 460, 462, 462, 458, 458, 445, 458, 451,
444, 444, 430, 416, 409, 403, 403, 389, 375, NA, NA, NA, NA, NA, NA, NA, NA,
NA, 183, 677, 677, 846, 883, 886, 924, 938, 883, 946, 846, 911, 826, 826,
788, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, 307,
307, 368, 377, 383, 383, 383, 380, 377, 377, 377, 374, 374, 375, 375, 375,
375, 368, 371, 374, 375, 361, 375, 389, 375, 375, 375, 375, 375, 314, 169,
NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, 238, 285, 361, 374, 375, 375,
375, 375, 375, 389, 403, 389, 389, 389, 375, 375, 389, 375, 348, 361, 375, 348,
348, 361, 348, 342, 361, 361, 361, 365, 365, 361, 966, 966, 966, 959, 959,
946, 1021, 1021, 1026, 1086, 1131, 1131, 1146, 1130, 1172, 1240, 1172, 1117,
1103, 1026, 1026, 966, 919, 946, 882, 832, NA, NA, NA, NA, NA, NA, NA, NA,
NA, NA)
pitchSmoothPraat(pitch, bandwidth = 10, samplingRate = 40, plot = TRUE)
pitchSmoothPraat(pitch, bandwidth = 2, samplingRate = 40, plot = TRUE)
```

pitch_app

*Interactive pitch tracker***Description**

Starts a shiny app for manually editing pitch contours. The settings in the panels on the left correspond to arguments to [analyze](#) - see `?analyze` and the vignette on acoustic analysis for help and examples. You can verify the pitch contours first, and then feed them back into `analyze` (see examples).

Usage

```
pitch_app()
```

Value

When proceeding to the next file in the cue, two types of backups are created. (1) A global object called "my_pitch" is created or updated. This list becomes visible when the app is terminated, and it contains the usual outputs of `analyze()` (`$detailed` and `$summary`) plus lists of manually corrected voiced and unvoiced frames. (2) The app saves to disk a .csv file with one row per audio file. Apart from the usual descriptives from `analyze()`, there are two additional columns: "time" with time stamps (the midpoint of each STFT frame, ms) and "pitch" with the manually corrected pitch values for each frame (Hz). When the orange "Download results" button is clicked, a context menu pops up offering to terminate the app - if that happens, the results are also returned directly into R. To process pitch contours further in R, work directly with `my_pitch[[myfile]]$time` and `my_pitch[[myfile]]$pitch` or, if loading the csv file, do something like:


```
a = read.csv('~Downloads/output.csv', stringsAsFactors = FALSE)
pitch = as.numeric(unlist(strsplit(a$pitch, ',')))
mean(pitch, na.rm = TRUE); sd(pitch, na.rm = TRUE)
```

Suggested workflow

Start by setting the basic analysis settings such as pitchFloor, pitchCeiling, silence, etc. Then click "Load audio" to upload one or several audio files (wav/mp3). Long files will be very slow, so please cut your audio into manageable chunks (ideally <10 s). If Shiny complains that maximum upload size is exceeded, you can increase it, say to 30 MB, with 'options(shiny.maxRequestSize = 30 * 1024^2)'. Once the audio has been uploaded to the browser, fine-tune the analysis settings as needed, edit the pitch contour in the first file to your satisfaction, then click "Next" to proceed to the next file, etc. Remember that setting a reasonable prior is often faster than adjusting the contour one anchor at a time. When done, click "Save results". If working with many files, you might want to save the results occasionally in case the app crashes (although you should still be able to recover your data if it does - see below).

How to edit pitch contours

Left-click to add a new anchor, double-click to remove it or unvoice the frame. Each time you make a change, the entire pitch contour is re-fit, so making a change in one frame can affect the path through candidates in adjacent frames. You can control this behavior by changing the settings in Out/Path and Out/Smoothing. If correctly configured, the app corrects the contour with only a few manual values - you shouldn't need to manually edit every single frame. For longer files, you can zoom in/out and navigate within the file. You can also select a region to voice/unvoice or shift it as a whole or to set a prior based on selected frequency range.

Recovering lost data

Every time you click "next" or "last" to move in between files in the queue, the output you've got so far is saved in a backup file called "temp.csv", and the "my_pitch" global object is updated. If the app crashes or is closed without saving the results, this backup file preserves your data. To recover it, access this file manually on disk or simply restart pitch_app() - a dialog box will pop up and ask whether you want to append the old data to the new one. Path to backup file: "[R_installation_folder]/soundgen/shiny/pitch_app/www/temp.csv", for example, "/home/allgoodguys/R/x86_64-pc-linux-gnu-library/3.6/soundgen/shiny/pitch_app/www/temp.csv"

See Also

[formant_app](#)

Examples

```
## Not run:
# Recommended workflow for analyzing a lot of short audio files
path_to_audio = '~/Downloads/temp' # our audio lives here

# STEP 1: extract manually corrected pitch contours
my_pitch = pitch_app() # runs in default browser such as Firefox or Chrome
# To change system default browser, run something like:
options('browser' = '/usr/bin/firefox') # path to the executable on Linux

# STEP 2: run analyze() with manually corrected pitch contours to obtain
```

```

# accurate descriptives like the proportion of energy in harmonics above f0,
# etc. This also gives you formants and loudness estimates (disabled in
# pitch_app to speed things up)
df2 = analyze(path_to_audio,
  pitchMethods = 'autocor', # only needed for HNR
  nFormants = 5,           # now we can measure formants as well
  pitchManual = my_pitch
  # or, if loading the output of pitch_app() from the disk:
  # pitchManual = '~/Downloads/output.csv'
  # pitchManual = '~/path_to_some_folder/my_pitch_contours.rds
  # etc
)

# STEP 3: add other acoustic descriptors, for ex.
df3 = segment(path_to_audio)

# STEP 4: merge df2, df3, df4, ... in R or a spreadsheet editor to have all
# acoustic descriptives together

# To verify your pitch contours and/or edit them later, copy output.csv to
# the folder with your audio, run pitch_app(), and load the audio + csv
# together. The saved pitch contours are treated as manual anchors

## End(Not run)

```

playme

Play audio

Description

Plays one or more sounds: wav/mp3 file(s), Wave objects, or numeric vectors. This is a simple wrapper for the functionality provided by [play](#). Recommended players on Linux: "play" from the "vox" library (default), "aplay".

Usage

```
playme(x, samplingRate = 16000, player = NULL, from = NULL, to = NULL)
```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
player	the name of player to use, eg "aplay", "play", "vlc", etc. Defaults to "play" on Linux, "afplay" on MacOS, and tuneR default on Windows. In case of errors, try setting another default player for play
from, to	play a selected time range (s)

Examples

```
## Not run:
# Play an audio file:
playme('pathToMyAudio/audio.wav')

# Create and play a numeric vector:
f0_Hz = 440
sound = sin(2 * pi * f0_Hz * (1:16000) / 16000)
playme(sound, 16000)
playme(sound, 16000, from = .1, to = .5) # play from 100 to 500 ms

# In case of errors, look into tuneR::play(). For ex., you might need to
# specify which player to use:
playme(sound, 16000, player = 'aplay')

# To avoid doing it all the time, set the default player:
tuneR::setWavPlayer('aplay')
playme(sound, 16000) # should now work without specifying the player

## End(Not run)
```

plotMS

Plot modulation spectrum

Description

Plots a single modulation spectrum returned by `modulationSpectrum`. The result is the same as the plot produced by `modulationSpectrum`, but calling `plotMS` is handy for processed modulation spectra - for instance, for plotting the difference between the modulation spectra of two sounds or groups of sounds.

Usage

```
plotMS(
  ms,
  X = NULL,
  Y = NULL,
  quantiles = c(0.5, 0.8, 0.9),
  colorTheme = c("bw", "seewave", "heat.colors", "...")[1],
  col = NULL,
  logWarpX = NULL,
  logWarpY = NULL,
  main = NULL,
  xlab = "Hz",
  ylab = "1/KHz",
  xlim = NULL,
  ylim = NULL,
  audio = NULL,
```

```
    ...
  )
```

Arguments

<code>ms</code>	modulation spectrum - a matrix with temporal modulation in columns and spectral modulation in rows, as returned by modulationSpectrum
<code>X, Y</code>	rownames and colnames of <code>ms</code> , respectively
<code>quantiles</code>	labeled contour values, % (e.g., "50" marks regions that contain 50% of the sum total of the entire modulation spectrum)
<code>colorTheme</code>	black and white ('bw'), as in <code>seewave</code> package ('seewave'), or any palette from palette such as 'heat.colors', 'cm.colors', etc
<code>col</code>	actual colors, eg <code>rev(rainbow(100))</code> - see <code>?hcl.colors</code> for colors in base R (overrides <code>colorTheme</code>)
<code>logWarpX, logWarpY</code>	numeric vector of length 2: <code>c(sigma, base)</code> of pseudolog-warping the modulation spectrum, as in function <code>pseudo_log_trans()</code> from the "scales" package
<code>xlab, ylab, main, xlim, ylim</code>	graphical parameters
<code>audio</code>	(internal) a list of audio attributes
<code>...</code>	other graphical parameters passed on to <code>filled.contour.mod</code> and contour (see spectrogram)

Examples

```
ms1 = modulationSpectrum(runif(4000), samplingRate = 16000, plot = TRUE)
plotMS(ms1$processed) # identical to above

# compare two modulation spectra
ms2 = modulationSpectrum(soundgen(syllLen = 100, addSilence = 0),
                          samplingRate = 16000)
# ensure the two matrices have the same dimensions
ms2_resized = soundgen::interpolMatrix(ms2$original,
                                       nr = nrow(ms1$original), nc = ncol(ms1$original))
# plot the difference
plotMS(log(ms1$original / ms2_resized), quantile = NULL,
       col = colorRampPalette(c('blue', 'yellow')) (50))
```

presets

Presets

Description

A library of presets for easy generation of a few nice sounds.

Usage

```
presets
```

Format

A list of length 4.

prosody	<i>Prosody</i>
---------	----------------

Description

Exaggerates or flattens the intonation by performing a dynamic pitch shift, changing pitch excursion from its original median value without changing the formants. This is a particular case of pitch shifting, which is performed with `shiftPitch`. The result is likely to be improved if manually corrected pitch contours are provided. Depending on the nature of audio, the settings that control pitch shifting may also need to be fine-tuned with the `shiftPitch_pars` argument.

Usage

```
prosody(
  x,
  samplingRate = NULL,
  multProsody,
  analyze_pars = list(),
  shiftPitch_pars = list(),
  pitchManual = NULL,
  play = FALSE,
  saveAudio = NULL,
  reportEvery = NULL,
  cores = 1,
  plot = FALSE,
  savePlots = NULL,
  width = 900,
  height = 500,
  units = "px",
  res = NA,
  ...
)
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)

<code>multProsody</code>	multiplier of pitch excursion from median (on a logarithmic or musical scale): >1 = exaggerate intonation, 1 = no change, <1 = flatten, 0 = completely flat at the original median pitch
<code>analyze_pars</code>	a list of parameters to pass to analyze (only needed if <code>pitchManual</code> is NULL - that is, if we attempt to track pitch automatically)
<code>shiftPitch_pars</code>	a list of parameters to pass to shiftPitch to fine-tune the pitch-shifting algorithm
<code>pitchManual</code>	manually corrected pitch contour. For a single sound, provide a numeric vector of any length. For multiple sounds, provide a dataframe with columns "file" and "pitch" (or path to a csv file) as returned by pitch_app , ideally with the same <code>windowLength</code> and <code>step</code> as in current call to <code>analyze</code> . A named list with pitch vectors per file is also OK (eg as returned by <code>pitch_app</code>)
<code>play</code>	if TRUE, plays the processed audio
<code>saveAudio</code>	full (!) path to folder for saving the processed audio; NULL = don't save, "" = same as input folder (NB: overwrites the originals!)
<code>reportEvery</code>	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
<code>cores</code>	number of cores for parallel processing
<code>plot</code>	should a spectrogram be plotted? TRUE / FALSE
<code>savePlots</code>	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
<code>width, height, units, res</code>	graphical parameters for saving plots passed to png
<code>...</code>	other graphical parameters

Value

If the input is a single audio (file, Wave, or numeric vector), returns the processed waveform as a numeric vector with the original sampling rate and scale. If the input is a folder with several audio files, returns a list of processed waveforms, one for each file.

See Also

[shiftPitch](#)

Examples

```
s = soundgen(syllLen = 200, pitch = c(150, 220), addSilence = 50,
             plot = TRUE, yScale = 'log')
# playme(s)
s1 = prosody(s, 16000, multProsody = 2,
            analyze_pars = list(windowLength = 30, step = 15),
            shiftPitch_pars = list(windowLength = 20, step = 5, freqWindow = 300),
            plot = TRUE)
# playme(s1)
# spectrogram(s1, 16000, yScale = 'log')
```

```

## Not run:
# Flat intonation - remove all frequency modulation
s2 = prosody(s, 16000, multProsody = 0,
  analyze_pars = list(windowLength = 30, step = 15),
  shiftPitch_pars = list(windowLength = 20, step = 5, freqWindow = 300),
  plot = TRUE)
playme(s2)
spectrogram(s2, 16000, yScale = 'log')

# Download an example - a bit of speech (sampled at 16000 Hz)
download.file('http://cogsci.se/soundgen/audio/speechEx.wav',
  destfile = '~/Downloads/temp1/speechEx.wav')
target = '~/Downloads/temp1/speechEx.wav'
samplingRate = tuneR::readWave(target)@samp.rate
spectrogram(target, yScale = 'log')
playme(target)

s3 = prosody(target, multProsody = 1.5,
  analyze_pars = list(windowLength = 30, step = 15),
  shiftPitch_pars = list(freqWindow = 400, propagation = 'adaptive'))
spectrogram(s3, tuneR::readWave(target)@samp.rate, yScale = 'log')
playme(s3)

# process all audio files in a folder
s4 = prosody '~/Downloads/temp', multProsody = 2, savePlots = '',
  saveAudio = '~/Downloads/temp/prosody')
str(s4) # returns a list with audio (+ saves it to disk)

## End(Not run)

```

reportTime

Report time

Description

Provides a nicely formatted "estimated time left" in loops plus a summary upon completion.

Usage

```

reportTime(
  i,
  time_start,
  nIter = NULL,
  reportEvery = NULL,
  jobs = NULL,
  prefix = ""
)

```

Arguments

<code>i</code>	current iteration
<code>time_start</code>	time when the loop started running
<code>nIter</code>	total number of iterations
<code>reportEvery</code>	report progress every <code>n</code> iterations
<code>jobs</code>	vector of length <code>nIter</code> specifying the relative difficulty of each iteration. If not NULL, estimated time left takes into account whether the jobs ahead will take more or less time than the jobs already completed
<code>prefix</code>	a string to print before "Done...", eg "Chain 1: "

Examples

```

time_start = proc.time()
for (i in 1:100) {
  Sys.sleep(i ^ 1.02 / 10000)
  reportTime(i = i, time_start = time_start, nIter = 100,
    jobs = (1:100) ^ 1.02, prefix = 'Chain 1: ')
}
## Not run:
# Unknown number of iterations:
time_start = proc.time()
for (i in 1:20) {
  Sys.sleep(i ^ 2 / 10000)
  reportTime(i = i, time_start = time_start,
    jobs = (1:20) ^ 2, reportEvery = 5)
}

# when analyzing a bunch of audio files, their size is a good estimate
# of how long each will take to process
time_start = proc.time()
filenames = list.files('~Downloads/temp', pattern = "*.wav|.mp3",
  full.names = TRUE)
fileSizes = file.info(filenames)$size
for (i in 1:length(filenames)) {
  # ...do what you have to do with each file...
  reportTime(i = i, nIter = length(filenames),
    time_start = time_start, jobs = fileSizes)
}

## End(Not run)

```


Description

Changes the sampling rate without introducing artefacts like aliasing. Algorithm: to downsample, applies a low-pass filter, then decimates with `approx`; to upsample, performs linear interpolation with `approx`, then applies a low-pass filter. NAs can be interpolated or preserved in the output. The length of output is determined, in order of precedence, by `len / mult / samplingRate_new`. For simple vector operations, this is very similar to `approx`, but the leading and trailing NAs are also preserved (see examples).

Usage

```
resample(
  x,
  samplingRate = NULL,
  samplingRate_new = NULL,
  mult = NULL,
  len = NULL,
  lowPass = TRUE,
  na.rm = FALSE,
  reportEvery = NULL,
  cores = 1,
  saveAudio = NULL,
  plot = FALSE,
  savePlots = NULL,
  width = 900,
  height = 500,
  units = "px",
  res = NA,
  ...
)
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>samplingRate_new</code>	an alternative to <code>mult</code> provided that the old <code>samplingRate</code> is known (NB: <code>mult</code> takes precedence)
<code>mult</code>	multiplier of sampling rate: new sampling rate = old sampling rate \times <code>mult</code> , so 1 = no effect, >1 = upsample, <1 = downsample
<code>len</code>	if specified, overrides <code>mult</code> and <code>samplingRate_new</code> and simply returns a vector of length <code>len</code>
<code>lowPass</code>	if TRUE, applies a low-pass filter before decimating or after upsampling to avoid aliasing
<code>na.rm</code>	if TRUE, NAs are interpolated, otherwise they are preserved in the output
<code>reportEvery</code>	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)

cores	number of cores for parallel processing
saveAudio	full path to the folder in which to save audio files (one per detected syllable)
plot	should a spectrogram be plotted? TRUE / FALSE
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
width, height, units, res	graphical parameters for saving plots passed to png
...	other graphical parameters

Examples

```
## Example 1: a short vector with NAs
x = c(NA, 1, 2, 3, NA, NA, 6, 7, 8, NA)

# upsample
resample(x, mult = 3.5, lowPass = FALSE, plot = TRUE) # just approx
resample(x, mult = 3.5, lowPass = TRUE, plot = TRUE) # low-pass + approx
resample(x, mult = 3.5, lowPass = FALSE, na.rm = TRUE, plot = TRUE)

# downsample
resample(x, mult = 0.5, lowPass = TRUE, plot = TRUE)
resample(x, mult = 0.5, na.rm = TRUE, plot = TRUE)
resample(x, len = 5, na.rm = TRUE, plot = TRUE) # same

# The most important TIP: use resample() for audio files and the internal
# soundgen::.resample(list(sound = ...)) for simple vector operations because
# it's >1000 times faster. For example:
soundgen::.resample(list(sound = x), mult = 3.5, lowPass = FALSE)

## Example 2: a sound
silence = rep(0, 10)
samplingRate = 1000
fr = seq(100, 300, length.out = 400)
x = c(silence, sin(cumsum(fr) * 2 * pi / samplingRate), silence)
spectrogram(x, samplingRate)

# downsample
x1 = resample(x, mult = 1 / 2.5)
spectrogram(x1, samplingRate / 2.5) # no aliasing
# cf:
x1bad = resample(x, mult = 1 / 2.5, lowPass = FALSE)
spectrogram(x1bad, samplingRate / 2.5) # aliasing

# upsample
x2 = resample(x, mult = 3)
spectrogram(x2, samplingRate * 3) # nothing above the old Nyquist
# cf:
x2bad = resample(x, mult = 3, lowPass = FALSE)
spectrogram(x2bad, samplingRate * 3) # high-frequency artefacts

## Not run:
```

```
# Example 3: resample all audio files in a folder to 8000 Hz
resample('~Downloads/temp', saveAudio = '~Downloads/temp/sr8000/',
         samplingRate_new = 8000, savePlots = '~Downloads/temp/sr8000/')

## End(Not run)
```

reverb

*Reverb & echo***Description**

Adds reverberation and/or echo to a sound. Algorithm for reverb: adds time-shifted copies of the signal weighted by a decay function, which is analogous to convoluting the input with a parametric model of some hypothetical impulse response function. In simple terms: we specify how much and when the sound rebounds back (as from a wall) and add these time-shifted copies to the original, optionally with some spectral filtering.

Usage

```
reverb(
  x,
  samplingRate = NULL,
  echoDelay = 200,
  echoLevel = -20,
  reverbDelay = 70,
  reverbSpread = 130,
  reverbLevel = -25,
  reverbDensity = 50,
  reverbType = "gaussian",
  filter = list(),
  dynamicRange = 80,
  output = c("audio", "detailed")[1],
  play = FALSE,
  reportEvery = NULL,
  cores = 1,
  saveAudio = NULL
)
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>echoDelay</code>	the delay at which the echo appears, ms
<code>echoLevel</code>	the rate at which the echo weakens at each repetition, dB
<code>reverbDelay</code>	the time of maximum reverb density, ms

reverbSpread	standard deviation of reverb spread around time reverbDelay, ms
reverbLevel	the maximum amplitude of reverb, dB below input
reverbDensity	the number of echos or "voices" added
reverbType	so far only "gaussian" has been implemented
filter	(optional) a spectral filter to apply to the created reverb and echo (see addFormants for acceptable formats)
dynamicRange	the precision with which the reverb and echo are calculated, dB
output	"audio" = returns just the processed audio, "detailed" = returns a list with reverb window, the added reverb/echo, etc.
play	if TRUE, plays the processed audio
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
saveAudio	full (!) path to folder for saving the processed audio; NULL = don't save, "" = same as input folder (NB: overwrites the originals!)

Examples

```

s = soundgen()
s_rev = reverb(s, 16000)
# playme(s_rev)

## Not run:
# double echo, no reverb
s1 = reverb(s, samplingRate = 16000, reverbLevel = NULL,
            echoDelay = c(250, 800), echoLevel = c(-15, -25))
# playme(s1)
# spectrogram(s1, 16000, osc = TRUE, ylim = c(0, 4))

# only reverb (indoors)
s2 = reverb(s, samplingRate = 16000, echoDelay = NULL,
            reverbDelay = 70, reverbSpread = 130,
            reverbLevel = -20, reverbDensity = 20)
# playme(s2)
# spectrogram(s2, 16000, osc = TRUE, ylim = c(0, 4))

# reverb (caves)
s3 = reverb(s, samplingRate = 16000, echoDelay = NULL,
            reverbDelay = 600, reverbSpread = 1500,
            reverbLevel = -10, reverbDensity = 100)
# playme(s3)
# spectrogram(s3, 16000, osc = TRUE, ylim = c(0, 4))

# both echo and reverb with high frequencies emphasized
s4 = reverb(s, samplingRate = 16000,
            echoDelay = 250, echoLevel = -20,
            reverbDelay = 70, reverbSpread = 120,
            reverbLevel = -25, reverbDensity = 50,

```

```

        filter = list(formants = NULL, lipRad = 3))
# playme(s4)
# spectrogram(s4, 16000, osc = TRUE, ylim = c(0, 4))

# add reverb to a recording
s5 = reverb('~Downloads/temp260/ut_fear_57-m-tone.wav',
            echoDelay = 850, echoLevel = -40)
# playme(s5, 44100)

# add reverb to all files in a folder, save the result
reverb('~Downloads/temp2', saveAudio = '~Downloads/temp2/rvb')

## End(Not run)

```

schwa

Schwa-related formant conversion

Description

This function performs several conceptually related types of conversion of formant frequencies in relation to the neutral schwa sound based on the one-tube model of the vocal tract. This is useful for speaker normalization because absolute formant frequencies measured in Hz depend strongly on overall vocal tract length (VTL). For example, adult men vs. children or grizzly bears vs. dog puppies have very different formant spaces in Hz, but it is possible to define a VTL-normalized formant space that is applicable to all species and sizes. Case 1: if we know vocal tract length (VTL) but not formant frequencies, `schwa()` estimates formants corresponding to a neutral schwa sound in this vocal tract, assuming that it is perfectly cylindrical. Case 2: if we know the frequencies of a few lower formants, `schwa()` estimates the deviation of observed formant frequencies from the neutral values expected in a perfectly cylindrical vocal tract (based on the VTL as specified or as estimated from formant dispersion). Case 3: if we want to generate a sound with particular relative formant frequencies (e.g. high F1 and low F2 relative to the schwa for this vocal tract), `schwa()` calculates the corresponding formant frequencies in Hz. See examples below for an illustration of these three suggested uses.

Usage

```

schwa(
  formants = NULL,
  vocalTract = NULL,
  formants_relative = NULL,
  nForm = 8,
  interceptZero = TRUE,
  tube = c("closed-open", "open-open")[1],
  speedSound = 35400,
  plot = FALSE
)

```

Arguments

<code>formants</code>	a numeric vector of observed (measured) formant frequencies, Hz
<code>vocalTract</code>	the length of vocal tract, cm
<code>formants_relative</code>	a numeric vector of target relative formant frequencies, % deviation from schwa (see examples)
<code>nForm</code>	the number of formants to estimate (integer)
<code>interceptZero</code>	if TRUE, forces the regression curve to pass through the origin. This reduces the influence of highly variable lower formants, but we have to commit to a particular model of the vocal tract: closed-open or open-open/closed-closed (method = "regression" only)
<code>tube</code>	the vocal tract is assumed to be a cylindrical tube that is either "closed-open" or "open-open" (same as closed-closed)
<code>speedSound</code>	speed of sound in warm air, cm/s. Stevens (2000) "Acoustic phonetics", p. 138
<code>plot</code>	if TRUE, plots vowel quality in speaker-normalized F1-F2 space

Details

Algorithm: the expected formant dispersion is given by $(2 * \text{formant_number} - 1) * \text{speedSound} / (4 * \text{formant_frequency})$ for a closed-open tube (mouth open) and $\text{formant_number} * \text{speedSound} / (2 * \text{formant_frequency})$ for an open-open or closed-closed tube. F1 is schwa is expected at half the value of formant dispersion. See e.g. Stevens (2000) "Acoustic phonetics", p. 139. Basically, we estimate vocal tract length and see if each formant is higher or lower than expected for this vocal tract. For this to work, we have to know either the frequencies of enough formants (not just the first two) or the true length of the vocal tract. See also [estimateVTL](#) on the algorithm for estimating formant dispersion if VTL is not known (note that schwa calls [estimateVTL](#) with the option `method = 'regression'`).

Value

Returns a list with the following components:

- `vtl_measured`** VTL as provided by the user, cm
- `vocalTract_apparent`** VTL estimated based on formants frequencies provided by the user, cm
- `formantDispersion`** average distance between formants, Hz
- `ff_measured`** formant frequencies as provided by the user, Hz
- `ff_schwa`** formant frequencies corresponding to a neutral schwa sound in this vocal tract, Hz
- `ff_theoretical`** formant frequencies corresponding to the user-provided relative formant frequencies, Hz
- `ff_relative`** deviation of formant frequencies from those expected for a schwa, % (e.g. if the first `ff_relative` is -25, it means that F1 is 25% lower than expected for a schwa in this vocal tract)
- `ff_relative_semitones`** deviation of formant frequencies from those expected for a schwa, semitones. Like `ff_relative`, this metric is invariant to vocal tract length, but the variance tends to be greater for lower vs. higher formants

ff_relative_dF deviation of formant frequencies from those expected for a schwa, proportion of formant spacing (dF). Unlike `ff_relative` and `ff_relative_semitones`, this metric has similar variance for lower and higher formants

See Also

[estimateVTL](#)

Examples

```
## CASE 1: known VTL
# If vocal tract length is known, we calculate expected formant frequencies
schwa(vocalTract = 17.5)
schwa(vocalTract = 13, nForm = 5)
schwa(vocalTract = 13, nForm = 5, tube = 'open-open')

## CASE 2: known (observed) formant frequencies
# Let's take formant frequencies in four vocalizations, namely
# (/a/, /i/, /mmm/, /roar/) by the same male speaker:
formants_a = c(860, 1430, 2900, NA, 5200) # NAs are OK - here F4 is unknown
s_a = schwa(formants = formants_a, plot = TRUE)
s_a
# We get an estimate of VTL (s_a$vtl_apparent),
# same as with estimateVTL(formants_a)
# We also get theoretical schwa formants: s_a$ff_schwa
# And we get the difference (% and semitones) in observed vs expected
# formant frequencies: s_a[c('ff_relative', 'ff_relative_semitones')]
# [a]: F1 much higher than expected, F2 slightly lower (see plot)

formants_i = c(300, 2700, 3400, 4400, 5300, 6400)
s_i = schwa(formants = formants_i, plot = TRUE)
s_i
# The apparent VTL is slightly smaller (14.5 cm)
# [i]: very low F1, very high F2

formants_mmm = c(1200, 2000, 2800, 3800, 5400, 6400)
schwa(formants_mmm, tube = 'closed-closed', plot = TRUE)
# ~schwa, but with a closed mouth

formants_roar = c(550, 1000, 1460, 2280, 3350,
                 4300, 4900, 5800, 6900, 7900)
s_roar = schwa(formants = formants_roar, plot = TRUE)
s_roar
# Note the enormous apparent VTL (22.5 cm!)
# (lowered larynx and rounded lips exaggerate the apparent size)
# s_roar$ff_relative: high F1 and low F2-F4

schwa(formants = formants_roar[1:4], plot = TRUE)
# based on F1-F4, apparent VTL is almost 28 cm!
# Since the lowest formants are the most salient,
# the apparent size is exaggerated even further
```

```

# If you know VTL, a few lower formants are enough to get
# a good estimate of the relative formant values:
schwa(formants = formants_roar[1:4], vocalTract = 19, plot = TRUE)
# NB: in this case theoretical and relative formants are calculated
# based on user-provided VTL (vtl_measured) rather than vtl_apparent

## CASE 3: from relative to absolute formant frequencies
# Say we want to generate a vowel sound with F1 20% below schwa
# and F2 40% above schwa, with VTL = 15 cm
s = schwa(formants_relative = c(-20, 40), vocalTract = 15, plot = TRUE)
# s$ff_schwa gives formant frequencies for a schwa, while
# s$ff_theoretical gives formant frequencies for a sound with
# target relative formant values (low F1, high F2)
schwa(formants = s$ff_theoretical)

```

segment

Segment a sound

Description

Finds syllables and bursts separated by background noise in long recordings (up to 1-2 hours of audio per file). Syllables are defined as continuous segments that seem to be different from noise based on amplitude and/or spectral similarity thresholds. Bursts are defined as local maxima in signal envelope that are high enough both in absolute terms (relative to the global maximum) and with respect to the surrounding region (relative to local minima). See vignette('acoustic_analysis', package = 'soundgen') for details.

Usage

```

segment(
  x,
  samplingRate = NULL,
  from = NULL,
  to = NULL,
  shortestSyl = 40,
  shortestPause = 40,
  method = c("env", "spec", "mel")[3],
  propNoise = NULL,
  SNR = NULL,
  noiseLevelStabWeight = c(1, 0.25),
  windowLength = 40,
  step = NULL,
  overlap = 80,
  reverbPars = list(reverbDelay = 70, reverbSpread = 130, reverbLevel = -35,
    reverbDensity = 50),
  interburst = NULL,
  peakToTrough = SNR + 3,
  troughLocation = c("left", "right", "both", "either")[4],

```



```

summaryFun = c("median", "sd"),
maxDur = 30,
reportEvery = NULL,
cores = 1,
plot = FALSE,
savePlots = NULL,
saveAudio = NULL,
addSilence = 50,
main = NULL,
xlab = "",
ylab = "Signal, dB",
showLegend = FALSE,
width = 900,
height = 500,
units = "px",
res = NA,
maxPoints = c(1e+05, 5e+05),
specPlot = list(colorTheme = "bw"),
contourPlot = list(lty = 1, lwd = 2, col = "green"),
sylPlot = list(lty = 1, lwd = 2, col = "blue"),
burstPlot = list(pch = 8, cex = 3, col = "red"),
...
)

```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
from, to	if NULL (default), analyzes the whole sound, otherwise from...to (s)
shortestSyl	minimum acceptable length of syllables, ms
shortestPause	minimum acceptable break between syllables, ms (syllables separated by shorter pauses are merged)
method	the signal used to search for syllables: 'env' = Hilbert-transformed amplitude envelope, 'spec' = spectrogram, 'mel' = mel-transformed spectrogram (see tuneR::melfcc)
propNoise	the proportion of non-zero sound assumed to represent background noise (note that complete silence is not considered, so padding with silence won't affect the algorithm)
SNR	expected signal-to-noise ratio (dB above noise), which determines the threshold for syllable detection. The meaning of "dB" here is approximate since the "signal" may be different from sound intensity
noiseLevelStabWeight	a vector of length 2 specifying the relative weights of the overall signal level vs. stability when attempting to automatically locate the regions that represent noise. Increasing the weight of stability tends to accentuate the beginning and end of each syllable.

windowLength	length of FFT window, ms
step	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
overlap	overlap between successive FFT frames, %
reverbPars	parameters passed on to <code>reverb</code> to attempt to cancel the effects of reverberation or echo, which otherwise tend to merge short and loud segments like rapid barks
interburst	minimum time between two consecutive bursts (ms). Defaults to the average detected (<code>syllable + pause</code>) / 2
peakToTrough	to qualify as a burst, a local maximum has to be at least <code>peakToTrough</code> dB above the left and/or right local trough(s) (controlled by <code>troughLocation</code>) over the analysis window (controlled by <code>interburst</code>). Defaults to SNR + 3 dB
troughLocation	should local maxima be compared to the trough on the left and/or right of it? Values: 'left', 'right', 'both', 'either'
summaryFun	functions used to summarize each acoustic characteristic; see <code>analyze</code>
maxDur	long files are split into chunks <code>maxDur</code> s in duration to avoid running out of RAM; the outputs for all fragments are glued together, but plotting is switched off. Note that noise profile is estimated in each chunk separately, so set it low if the background noise is highly variable
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing
plot	if TRUE, produces a segmentation plot
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
saveAudio	full path to the folder in which to save audio files (one per detected syllable)
addSilence	if syllables are saved as separate audio files, they can be padded with some silence (ms)
xlab, ylab, main	main plotting parameters
showLegend	if TRUE, shows a legend for thresholds
width, height, units, res	parameters passed to <code>png</code> if the plot is saved
maxPoints	the maximum number of "pixels" in the oscillogram (if any) and spectrogram; good for quickly plotting long audio files; defaults to <code>c(1e5, 5e5)</code>
specPlot	a list of graphical parameters for displaying the spectrogram (if <code>method = 'spec'</code> or <code>'mel'</code>); set to NULL to hide the spectrogram
contourPlot	a list of graphical parameters for displaying the signal contour used to detect syllables (see details)
sylPlot	a list of graphical parameters for displaying the syllables
burstPlot	a list of graphical parameters for displaying the bursts
...	other graphical parameters passed to <code>graphics::plot</code>

Details

Algorithm: for each chunk at most `maxDur` long, first the audio recording is partitioned into signal and noise regions: the quietest and most stable regions are located, and noise threshold is defined from a user-specified proportion of noise in the recording (`propNoise`) or, if `propNoise = NULL`, from the lowest local maximum in the density function of a weighted product of amplitude and stability (that is, we assume that quiet and stable regions are likely to represent noise). Once we know what the noise looks like - in terms of its typical amplitude and/or spectrum - we derive signal contour as its difference from noise at each time point. If `method = 'env'`, this is Hilbert transform minus noise, and if `method = 'spec'` or `'mel'`, this is the inverse of cosine similarity between the spectrum of each frame and the estimated spectrum of noise weighted by amplitude. By default, signal-to-noise ratio (SNR) is estimated as half-median of above-noise signal, but it is recommended that this parameter is adjusted by hand to suit the purposes of segmentation, as it is the key setting that controls the balance between false negatives (missing faint signals) and false positives (hallucinating signals that are actually noise). Note also that effects of echo or reverberation can be taken into account: syllable detection threshold may be raised following powerful acoustic bursts with the help of the `reverbPars` argument. At the final stage, continuous "islands" SNR dB above noise level are detected as syllables, and "peaks" on the islands are detected as bursts. The algorithm is very flexible, but the parameters may be hard to optimize by hand. If you have an annotated sample of the sort of audio you are planning to analyze, with syllables and/or bursts counted manually, you can use it for automatic optimization of control parameters (see [optimizePars](#)).

Value

If `summaryFun = NULL`, returns returns a list containing full stats on each syllable and burst (one row per syllable and per burst), otherwise returns only a dataframe with one row per file - a summary of the number and spacing of syllables and vocal bursts.

See Also

[analyze ssm](#)

Examples

```
sound = soundgen(nSyl = 4, sylLen = 100, pauseLen = 70,
                 attackLen = 20, amplGlobal = c(0, -20),
                 pitch = c(368, 284), temperature = .001)
# add noise so SNR decreases from 20 to 0 dB from syl1 to syl4
sound = sound + runif(length(sound), -10 ^ (-20 / 20), 10 ^ (-20 / 20))
# osc(sound, samplingRate = 16000, dB = TRUE)
# spectrogram(sound, samplingRate = 16000, osc = TRUE)
# playme(sound, samplingRate = 16000)

s = segment(sound, samplingRate = 16000, plot = TRUE)
s

# customizing the plot
segment(sound, samplingRate = 16000, plot = TRUE,
        sylPlot = list(lty = 2, col = 'gray20'),
        burstPlot = list(pch = 16, col = 'blue'),
        specPlot = list(col = rev(heat.colors(50))),
```

```

        xlab = 'Some custom label', cex.lab = 1.2,
        showLegend = TRUE,
        main = 'My awesome plot')
## Not run:
# set SNR manually to control detection threshold
s = segment(sound, samplingRate = 16000, SNR = 1, plot = TRUE)

# Download 260 sounds from the supplements to Anikin & Persson (2017) at
# http://cogsci.se/publications.html
# unzip them into a folder, say '~/Downloads/temp'
myfolder = '~/Downloads/temp260' # 260 .wav files live here
s = segment(myfolder, propNoise = .05, SNR = 3)

# Check accuracy: import a manual count of syllables (our "key")
key = segmentManual # a vector of 260 integers
trial = as.numeric(s$summary$nBursts)
cor(key, trial, use = 'pairwise.complete.obs')
boxplot(trial ~ as.integer(key), xlab='key')
abline(a=0, b=1, col='red')

# or look at the detected syllables instead of bursts:
cor(key, s$summary$nSyl, use = 'pairwise.complete.obs')

## End(Not run)

```

segmentManual

Manual counts of syllables in 260 sounds

Description

A vector of the number of syllables in the corpus of 260 human non-linguistic emotional vocalizations from Anikin & Persson (2017). The corpus can be downloaded from <http://cogsci.se/publications.html>

Usage

```
segmentManual
```

Format

An object of class `numeric` of length 260.

semitonesToHz	<i>Convert semitones to Hz</i>
---------------	--------------------------------

Description

Converts from semitones above C-5 (~0.5109875 Hz) or another reference frequency to Hz. See [HzToSemitones](#)

Usage

```
semitonesToHz(s, ref = 0.5109875)
```

Arguments

s	vector or matrix of frequencies (semitones above C0)
ref	frequency of the reference value (defaults to C-5, 0.51 Hz)

See Also

[HzToSemitones](#)

shiftFormants	<i>Shift formants</i>
---------------	-----------------------

Description

Raises or lowers formants (resonance frequencies), changing the voice quality or timbre of the sound without changing its pitch, statically or dynamically. Note that this is only possible when the fundamental frequency f_0 is lower than the formant frequencies. For best results, `freqWindow` should be no lower than f_0 and no higher than formant bandwidths. Obviously, this is impossible for many signals, so just try a few reasonable values, like ~200 Hz for speech. If `freqWindow` is not specified, `soundgen` sets it to the average detected f_0 , which is slow.

Usage

```
shiftFormants(  
  x,  
  multFormants,  
  samplingRate = NULL,  
  freqWindow = NULL,  
  dynamicRange = 80,  
  windowLength = 50,  
  step = NULL,  
  overlap = 75,  
  wn = "gaussian",
```

```

    interpol = c("approx", "spline")[1],
    normalize = c("max", "orig", "none")[2],
    play = FALSE,
    saveAudio = NULL,
    reportEvery = NULL,
    cores = 1,
    ...
)

```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>multFormants</code>	1 = no change, >1 = raise formants (eg 1.1 = 10% up, 2 = one octave up), <1 = lower formants. Anchor format accepted (see soundgen)
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>freqWindow</code>	the width of spectral smoothing window, Hz. Defaults to detected <code>f0</code>
<code>dynamicRange</code>	dynamic range, dB. All values more than one <code>dynamicRange</code> under maximum are treated as zero
<code>windowLength</code>	length of FFT window, ms
<code>step</code>	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
<code>overlap</code>	overlap between successive FFT frames, %
<code>wn</code>	window type accepted by <code>ftwindow</code> , currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop
<code>interpol</code>	the method for interpolating scaled spectra
<code>normalize</code>	"orig" = same as input (default), "max" = maximum possible peak amplitude, "none" = no normalization
<code>play</code>	if TRUE, plays the synthesized sound using the default player on your system. If character, passed to <code>play</code> as the name of player to use, eg "aplay", "play", "vlc", etc. In case of errors, try setting another default player for <code>play</code>
<code>saveAudio</code>	full path to the folder in which to save audio files (one per detected syllable)
<code>reportEvery</code>	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
<code>cores</code>	number of cores for parallel processing
...	other graphical parameters

Details

Algorithm: phase vocoder. In the frequency domain, we separate the complex spectrum of each STFT frame into two parts. The "receiver" is the flattened or smoothed complex spectrum, where smoothing is achieved by obtaining a smoothed magnitude envelope (the amount of smoothing is

controlled by freqWindow) and then dividing the complex spectrum by this envelope. This basically removes the formants from the signal. The second component, "donor", is a scaled and interpolated version of the same smoothed magnitude envelope as above - these are the formants shifted up or down. Warping can be easily implemented instead of simple scaling if nonlinear spectral transformations are required. We then multiply the "receiver" and "donor" spectrograms and reconstruct the audio with iSTFT.

See Also

[shiftPitch transplantFormants](#)

Examples

```
s = soundgen(syllen = 200, ampl = c(0,-10),
            pitch = c(250, 350), rolloff = c(-9, -15),
            noise = -40,
            formants = 'aii', addSilence = 50)
# playme(s)
s1 = shiftFormants(s, samplingRate = 16000, multFormants = 1.25,
                  freqWindow = 200)
# playme(s1)

## Not run:
data(sheep, package = 'seewave') # import a recording from seewave
playme(sheep)
spectrogram(sheep)

# Lower formants by 4 semitones or ~20% = 2 ^ (-4 / 12)
sheep1 = shiftFormants(sheep, multFormants = 2 ^ (-4 / 12), freqWindow = 150)
playme(sheep1, sheep@samp.rate)
spectrogram(sheep1, sheep@samp.rate)

orig = seewave::meanspec(sheep, w1 = 128, plot = FALSE)
shifted = seewave::meanspec(sheep1, w1 = 128, f = sheep@samp.rate, plot = FALSE)
plot(orig[, 1], log(orig[, 2]), type = 'l')
points(shifted[, 1], log(shifted[, 2]), type = 'l', col = 'blue')

# dynamic change: raise formants at the beginning, lower at the end
sheep2 = shiftFormants(sheep, multFormants = c(1.3, .7), freqWindow = 150)
playme(sheep2, sheep@samp.rate)
spectrogram(sheep2, sheep@samp.rate)

## End(Not run)
```

Description

Raises or lowers pitch with or without also shifting the formants (resonance frequencies) and performing a time-stretch. The three operations (pitch shift, formant shift, and time stretch) are independent and can be performed in any combination, statically or dynamically. `shiftPitch` can also be used to shift formants without changing pitch or duration, but the dedicated `shiftFormants` is faster for that task.

Usage

```
shiftPitch(
  x,
  multPitch = 1,
  multFormants = multPitch,
  timeStretch = 1,
  samplingRate = NULL,
  freqWindow = NULL,
  dynamicRange = 80,
  windowLength = 40,
  step = 2,
  overlap = NULL,
  wn = "gaussian",
  interpol = c("approx", "spline")[1],
  propagation = c("time", "adaptive")[1],
  preserveEnv = NULL,
  transplantEnv_pars = list(windowLength = 10),
  normalize = c("max", "orig", "none")[2],
  play = FALSE,
  saveAudio = NULL,
  reportEvery = NULL,
  cores = 1
)
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files <code>c('file1.wav', 'file2.mp3')</code> , Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>multPitch</code>	1 = no change, >1 = raise pitch (eg 1.1 = 10% up, 2 = one octave up), <1 = lower pitch. Anchor format accepted for <code>multPitch</code> / <code>multFormant</code> / <code>timeStretch</code> (see soundgen)
<code>multFormants</code>	1 = no change, >1 = raise formants (eg 1.1 = 10% up, 2 = one octave up), <1 = lower formants
<code>timeStretch</code>	1 = no change, >1 = longer, <1 = shorter
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>freqWindow</code>	the width of spectral smoothing window, Hz. Defaults to detected <code>f0</code> prior to pitch shifting - see shiftFormants for discussion and examples
<code>dynamicRange</code>	dynamic range, dB. All values more than one <code>dynamicRange</code> under maximum are treated as zero

windowLength	length of FFT window, ms
step	you can override overlap by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of 1/samplingRate, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
overlap	overlap between successive FFT frames, %
wn	window type accepted by <code>ftwindow</code> , currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop
interpol	the method for interpolating scaled spectra and anchors
propagation	the method for propagating phase: "time" = horizontal propagation (default), "adaptive" = an experimental implementation of "vocoder done right" (Prusa & Holighaus 2017)
preserveEnv	if TRUE, transplants the amplitude envelope from the original to the modified sound with <code>transplantEnv</code> . Defaults to TRUE if no time stretching is performed and FALSE otherwise
transplantEnv_pars	a list of parameters passed on to <code>transplantEnv</code> if <code>preserveEnv</code> = TRUE
normalize	"orig" = same as input (default), "max" = maximum possible peak amplitude, "none" = no normalization
play	if TRUE, plays the synthesized sound using the default player on your system. If character, passed to <code>play</code> as the name of player to use, eg "aplay", "play", "vlc", etc. In case of errors, try setting another default player for <code>play</code>
saveAudio	full path to the folder in which to save audio files (one per detected syllable)
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)
cores	number of cores for parallel processing

Details

Algorithm: phase vocoder. Pitch shifting is accomplished by performing a time stretch (at present, with horizontal or adaptive phase propagation) followed by resampling. This shifts both pitch and formants; to preserve the original formant frequencies or modify them independently of pitch, a variant of `link{transplantFormants}` is performed to "transplant" the original or scaled formants onto the time-stretched new sound.

See Also

[shiftFormants](#) [transplantFormants](#)

Examples

```
s = soundgen(syllLen = 200, ampl = c(0,-10),
             pitch = c(250, 350), rolloff = c(-9, -15),
             noise = -40,
             formants = 'aii', addSilence = 50)
# playme(s)
```

```

s1 = shiftPitch(s, samplingRate = 16000, freqWindow = 400,
               multPitch = 1.25, multFormants = .8)
# playme(s1)

## Not run:
## Dynamic manipulations
# Add a chevron-shaped pitch contour
s2 = shiftPitch(s, samplingRate = 16000, multPitch = c(1.1, 1.3, .8))
playme(s2)

# Time-stretch only the middle
s3 = shiftPitch(s, samplingRate = 16000, timeStretch = list(
  time = c(0, .25, .31, .5, .55, 1),
  value = c(1, 1, 3, 3, 1, 1))
)
playme(s3)

## Various combinations of 3 manipulations
data(sheep, package = 'seewave') # import a recording from seewave
playme(sheep)
spectrogram(sheep)

# Raise pitch and formants by 3 semitones, shorten by half
sheep1 = shiftPitch(sheep, multPitch = 2 ^ (3 / 12), timeStretch = 0.5)
playme(sheep1, sheep@samp.rate)
spectrogram(sheep1, sheep@samp.rate)

# Just shorten
shiftPitch(sheep, multPitch = 1, timeStretch = 0.25, play = TRUE)

# Raise pitch preserving formants
sheep2 = shiftPitch(sheep, multPitch = 1.2, multFormants = 1, freqWindow = 150)
playme(sheep2, sheep@samp.rate)
spectrogram(sheep2, sheep@samp.rate)

## End(Not run)

```

soundgen

Generate a sound

Description

Generates a bout of one or more syllables with pauses between them. Two basic components are synthesized: the harmonic component (the sum of sine waves with frequencies that are multiples of the fundamental frequency) and the noise component. Both components can be filtered with independently specified formants. Intonation and amplitude contours can be applied both within each syllable and across multiple syllables. Suggested application: synthesis of animal or human non-linguistic vocalizations. For more information, see <http://cogsci.se/soundgen.html> and `vignette('sound_generation', package = 'soundgen')`.

Usage

```
soundgen(  
  repeatBout = 1,  
  nSyl = 1,  
  syllLen = 300,  
  pauseLen = 200,  
  pitch = list(time = c(0, 0.1, 0.9, 1), value = c(100, 150, 135, 100)),  
  pitchGlobal = NA,  
  glottis = 0,  
  temperature = 0.025,  
  tempEffects = list(),  
  maleFemale = 0,  
  creakyBreathy = 0,  
  nonlinBalance = 100,  
  nonlinRandomWalk = NULL,  
  subRatio = 2,  
  subFreq = 0,  
  subDep = 0,  
  subWidth = 10000,  
  shortestEpoch = 300,  
  jitterLen = 1,  
  jitterDep = 0,  
  vibratoFreq = 5,  
  vibratoDep = 0,  
  shimmerDep = 0,  
  shimmerLen = 1,  
  attackLen = 50,  
  rolloff = -9,  
  rolloffOct = 0,  
  rolloffKHz = -3,  
  rolloffParab = 0,  
  rolloffParabHarm = 3,  
  rolloffExact = NULL,  
  lipRad = 6,  
  noseRad = 4,  
  mouthOpenThres = 0,  
  formants = c(860, 1430, 2900),  
  formantDep = 1,  
  formantDepStoch = 1,  
  formantWidth = 1,  
  formantCeiling = 2,  
  formantLocking = 0,  
  vocalTract = NA,  
  amDep = 0,  
  amFreq = 30,  
  amType = c("logistic", "sine")[1],  
  amShape = 0,  
  noise = NULL,
```

```

formantsNoise = NA,
rolloffNoise = -4,
noiseFlatSpec = 1200,
rolloffNoiseExp = 0,
noiseAmpRef = c("f0", "source", "filtered")[3],
mouth = list(time = c(0, 1), value = c(0.5, 0.5)),
ampl = NA,
amplGlobal = NA,
smoothing = list(interpol = c("approx", "spline", "loess")[3], loessSpan = NULL,
  discountThres = 0.05, jumpThres = 0.01),
samplingRate = 16000,
windowLength = 50,
overlap = 75,
addSilence = 100,
pitchFloor = 1,
pitchCeiling = 3500,
pitchSamplingRate = 16000,
dynamicRange = 80,
invalidArgAction = c("adjust", "abort", "ignore")[1],
plot = FALSE,
play = FALSE,
saveAudio = NA,
...
)

```

Arguments

repeatBout	number of times the whole bout should be repeated
nSyl	number of syllables in the bout. 'pitchGlobal', 'amplGlobal', and 'formants' span multiple syllables, but not multiple bouts
syllLen	average duration of each syllable, ms (vectorized)
pauseLen	average duration of pauses between syllables, ms (can be negative between bouts: force with invalidArgAction = 'ignore') (vectorized)
pitch	a numeric vector of f0 values in Hz or a dataframe specifying the time (ms or 0 to 1) and value (Hz) of each anchor, hereafter "anchor format". These anchors are used to create a smooth contour of fundamental frequency f0 (pitch) within one syllable
pitchGlobal	unlike pitch, these anchors are used to create a smooth contour of average f0 across multiple syllables. The values are in semitones relative to the existing pitch, i.e. 0 = no change (anchor format)
glottis	anchors for specifying the proportion of a glottal cycle with closed glottis, % (0 = no modification, 100 = closed phase as long as open phase); numeric vector or dataframe specifying time and value (anchor format)
temperature	hyperparameter for regulating the amount of stochasticity in sound generation
tempEffects	a list of scaling coefficients regulating the effect of temperature on particular parameters. To change, specify just those pars that you want to modify (1 =

default, 0 = no stochastic behavior). `amplDep`, `pitchDep`, `noiseDep`: random fluctuations of user-specified amplitude / pitch / noise anchors; `amplDriftDep`: drift of amplitude mirroring pitch drift; `formDisp`: dispersion of stochastic formants; `formDrift`: formant frequencies; `glottisDep`: proportion of glottal cycle with closed glottis; `pitchDriftDep`: amount of slow random drift of `f0`; `pitchDriftFreq`: frequency of slow random drift of `f0`; `rolloffDriftDep`: drift of rolloff mirroring pitch drift; `specDep`: rolloff, rolloffNoise, nonlinear effects, attack; `subDriftDep`: drift of subharmonic frequency and bandwidth mirroring pitch drift; `syllLenDep`: duration of syllables and pauses

<code>maleFemale</code>	hyperparameter for shifting <code>f0</code> contour, formants, and <code>vocalTract</code> to make the speaker appear more male (-1...0) or more female (0...+1); 0 = no change
<code>creakyBreathy</code>	hyperparameter for a rough adjustment of voice quality from creaky (-1) to breathy (+1); 0 = no change
<code>nonlinBalance</code>	hyperparameter for regulating the (approximate) proportion of sound with different regimes of pitch effects (none / subharmonics only / subharmonics and jitter). 0% = no noise; 100% = the entire sound has jitter + subharmonics. Ignored if <code>temperature</code> = 0
<code>nonlinRandomWalk</code>	a numeric vector specifying the timing of nonlinear regimes: 0 = none, 1 = subharmonics, 2 = subharmonics + jitter + shimmer
<code>subRatio</code>	a positive integer giving the ratio of <code>f0</code> (the main fundamental) to <code>g0</code> (a lower frequency): 1 = no subharmonics, 2 = period doubling regardless of pitch changes, 3 = period tripling, etc; <code>subRatio</code> overrides <code>subFreq</code> (anchor format)
<code>subFreq</code>	instead of a specific number of subharmonics (<code>subRatio</code>), we can specify the approximate <code>g0</code> frequency (Hz), which is used only if <code>subRatio</code> = 1 and is adjusted to <code>f0</code> so <code>f0/g0</code> is always an integer (anchor format)
<code>subDep</code>	the depth of subharmonics relative to the main frequency component (<code>f0</code>), %. 0: no subharmonics; 100: <code>g0</code> harmonics are as strong as the nearest <code>f0</code> harmonic (anchor format)
<code>subWidth</code>	Width of subharmonic sidebands - regulates how rapidly <code>g</code> -harmonics weaken away from <code>f</code> -harmonics: large values like the default 10000 means that all <code>g0</code> harmonics are equally strong (anchor format)
<code>shortestEpoch</code>	minimum duration of each epoch with unchanging subharmonics regime or formant locking, in ms
<code>jitterLen</code>	duration of stable periods between pitch jumps, ms. Use a low value for harsh noise, a high value for irregular vibrato or shaky voice (anchor format)
<code>jitterDep</code>	cycle-to-cycle random pitch variation, semitones (anchor format)
<code>vibratoFreq</code>	the rate of regular pitch modulation, or vibrato, Hz (anchor format)
<code>vibratoDep</code>	the depth of vibrato, semitones (anchor format)
<code>shimmerDep</code>	random variation in amplitude between individual glottal cycles (0 to 100% of original amplitude of each cycle) (anchor format)
<code>shimmerLen</code>	duration of stable periods between amplitude jumps, ms. Use a low value for harsh noise, a high value for shaky voice (anchor format)

attackLen	duration of fade-in / fade-out at each end of syllables and noise (ms): a vector of length 1 (symmetric) or 2 (separately for fade-in and fade-out)
rolloff	basic rolloff from lower to upper harmonics, db/octave (exponential decay). All rolloff parameters are in anchor format. See getRolloff for more details
rolloffOct	basic rolloff changes from lower to upper harmonics (regardless of f0) by rolloffOct dB/oct. For example, we can get steeper rolloff in the upper part of the spectrum
rolloffKHz	rolloff changes linearly with f0 by rolloffKHz dB/kHz. For ex., -6 dB/kHz gives a 6 dB steeper basic rolloff as f0 goes up by 1000 Hz
rolloffParab	an optional quadratic term affecting only the first rolloffParabHarm harmonics. The middle harmonic of the first rolloffParabHarm harmonics is amplified or dampened by rolloffParab dB relative to the basic exponential decay
rolloffParabHarm	the number of harmonics affected by rolloffParab
rolloffExact	user-specified exact strength of harmonics: a vector or matrix with one row per harmonic, scale 0 to 1 (overrides all other rolloff parameters)
lipRad	the effect of lip radiation on source spectrum, dB/oct (the default of +6 dB/oct produces a high-frequency boost when the mouth is open)
noseRad	the effect of radiation through the nose on source spectrum, dB/oct (the alternative to lipRad when the mouth is closed)
mouthOpenThres	open the lips (switch from nose radiation to lip radiation) when the mouth is open >mouthOpenThres, 0 to 1
formants	either a character string like "aui" referring to default presets for speaker "M1" or a list of formant times, frequencies, amplitudes, and bandwidths (see ex. below). formants = NA defaults to schwa. Time stamps for formants and mouthOpening can be specified in ms or an any other arbitrary scale. See getSpectralEnvelope for more details
formantDep	scale factor of formant amplitude (1 = no change relative to amplitudes in formants)
formantDepStoch	the amplitude of additional stochastic formants added above the highest specified formant, dB (only if temperature > 0)
formantWidth	scale factor of formant bandwidth (1 = no change)
formantCeiling	frequency to which stochastic formants are calculated, in multiples of the Nyquist frequency; increase up to ~10 for long vocal tracts to avoid losing energy in the upper part of the spectrum
formantLocking	the approximate proportion of sound in which one of the harmonics is locked to the nearest formant, 0 = none, 1 = the entire sound (anchor format)
vocalTract	the length of vocal tract, cm. Used for calculating formant dispersion (for adding extra formants) and formant transitions as the mouth opens and closes. If NULL or NA, the length is estimated based on specified formant frequencies, if any (anchor format)
amDep	amplitude modulation (AM) depth, %. 0: no change; 100: AM with amplitude range equal to the dynamic range of the sound (anchor format)
amFreq	AM frequency, Hz (anchor format)

amType	"sine" = sinusoidal, "logistic" = logistic (default)
amShape	ignore if amType = "sine", otherwise determines the shape of non-sinusoidal AM: 0 = ~sine, -1 = notches, +1 = clicks (anchor format)
noise	loudness of turbulent noise (0 dB = as loud as voiced component, negative values = quieter) such as aspiration, hissing, etc (anchor format)
formantsNoise	the same as formants, but for unvoiced instead of voiced component. If NA (default), the unvoiced component will be filtered through the same formants as the voiced component, approximating aspiration noise [h]
rolloffNoise, noiseFlatSpec	linear rolloff of the excitation source for the unvoiced component, rolloffNoise dB/kHz (anchor format) applied above noiseFlatSpec Hz
rolloffNoiseExp	exponential rolloff of the excitation source for the unvoiced component, dB/oct (anchor format) applied above 0 Hz
noiseAmpRef	noise amplitude is defined relative to: "f0" = the amplitude of the first partial (fundamental frequency), "source" = the amplitude of the harmonic component prior to applying formants, "filtered" = the amplitude of the harmonic component after applying formants
mouth	mouth opening (0 to 1, 0.5 = neutral, i.e. no modification) (anchor format)
ampl	amplitude envelope (dB, 0 = max amplitude) (anchor format)
amplGlobal	global amplitude envelope spanning multiple syllables (dB, 0 = no change) (anchor format)
smoothing	a list of parameters passed to getSmoothContour to control the interpolation and smoothing of contours: <code>interpol</code> (approx / spline / loess), <code>loessSpan</code> , <code>discontThres</code> , <code>jumpThres</code>
samplingRate	sampling frequency, Hz
windowLength	length of FFT window, ms
overlap	FFT window overlap, %. For allowed values, see istft
addSilence	silence before and after the bout, ms: a vector of length 1 (symmetric) or 2 (different duration of silence before/after the sound)
pitchFloor, pitchCeiling	lower & upper bounds of f0
pitchSamplingRate	sampling frequency of the pitch contour only, Hz. Low values reduce processing time. Set to <code>pitchCeiling</code> for optimal speed or to <code>samplingRate</code> for optimal quality
dynamicRange	dynamic range, dB. Harmonics and noise more than <code>dynamicRange</code> under maximum amplitude are discarded to save computational resources
invalidArgAction	what to do if an argument is invalid or outside the range in <code>permittedValues</code> : 'adjust' = reset to default value, 'abort' = stop execution, 'ignore' = throw a warning and continue (may crash)
plot	if TRUE, plots a spectrogram

play	if TRUE, plays the synthesized sound using the default player on your system. If character, passed to <code>play</code> as the name of player to use, eg "aplay", "play", "vlc", etc. In case of errors, try setting another default player for <code>play</code>
saveAudio	path + filename for saving the output, e.g. '~/Downloads/temp.wav'. If NULL = doesn't save
...	other plotting parameters passed to <code>spectrogram</code>

Value

Returns the synthesized waveform as a numeric vector.

See Also

[generateNoise](#) [beat](#) [fart](#)

Examples

```
# NB: GUI for soundgen is available as a Shiny app.
# Type "soundgen_app()" to open it in default browser

# Set "playback" to TRUE for default system player or the name of preferred
# player (eg "aplay") to play back the audio from examples
playback = FALSE # or TRUE 'aplay', 'vlc', ...

sound = soundgen(play = playback)
# spectrogram(sound, 16000, osc = TRUE)
# playme(sound)

# Control of intonation, amplitude envelope, formants
s0 = soundgen(
  pitch = c(300, 390, 250),
  ampl = data.frame(time = c(0, 50, 300), value = c(-5, -10, 0)),
  attack = c(10, 50),
  formants = c(600, 900, 2200),
  play = playback
)

# Use the in-built collection of presets:
# names(presets) # speakers
# names(presets$Chimpanzee) # calls per speaker
s1 = eval(parse(text = presets$Chimpanzee$Scream_conflict)) # screaming chimp
# playme(s1)
s2 = eval(parse(text = presets$F1$Scream)) # screaming woman
# playme(s2, 18320)
## Not run:
# unless temperature is 0, the sound is different every time
for (i in 1:3) sound = soundgen(play = playback, temperature = .2)

# Bouts versus syllables. Compare:
sound = soundgen(formants = 'uai', repeatBout = 3, play = playback)
sound = soundgen(formants = 'uai', nSyl = 3, play = playback)
```



```
# Intonation contours per syllable and globally:
sound = soundgen(nSyl = 5, syllLen = 200, pauseLen = 140,
  pitch = list(
    time = c(0, 0.65, 1),
    value = c(977, 1540, 826)),
  pitchGlobal = list(time = c(0, .5, 1), value = c(-6, 7, 0)),
  play = playback, plot = TRUE)

# Subharmonics / sidebands (noisy scream)
sound = soundgen(subFreq = 75, subDep = runif(10, 0, 60), subWidth = 130,
  pitch = list(
    time = c(0, .3, .9, 1), value = c(1200, 1547, 1487, 1154)),
  syllLen = 800,
  play = playback, plot = TRUE)

# Jitter and mouth opening (bark, dog-like)
sound = soundgen(repeatBout = 2, syllLen = 160, pauseLen = 100,
  subFreq = 100, subDep = 100, subWidth = 60, jitterDep = 1,
  pitch = c(559, 785, 557),
  mouth = c(0, 0.5, 0),
  vocalTract = 5, formants = NULL,
  play = playback, plot = TRUE)

# See the vignette on sound generation for more examples and in-depth
# explanation of the arguments to soundgen()
# Examples of code for creating human and animal vocalizations are available
# on project's homepage: http://cogsci.se/soundgen.html

## End(Not run)
```

soundgen_app

Interactive sound synthesizer

Description

Starts a shiny app that provides an interactive wrapper to [soundgen](#). Supported browsers: Firefox / Chrome. Note that the browser has to be able to playback WAV audio files, otherwise there will be no sound.

Usage

soundgen_app()

specToMS *Spectrogram to modulation spectrum*

Description

Takes a spectrogram (either complex or magnitude) and returns a MS with proper row and column labels.

Usage

```
specToMS(spec, windowLength = NULL, step = NULL)
```

Arguments

spec	target spectrogram (numeric matrix, frequency in rows, time in columns)
windowLength	length of FFT window, ms
step	you can override overlap by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of 1/samplingRate, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)

Value

Returns a MS - matrix of complex values of the same dimension as spec, with AM in rows and FM in columns.

Examples

```
s = soundgen(syllLen = 500, amFreq = 25, amDep = 50,
             pitch = 250, samplingRate = 16000)
spec = spectrogram(s, samplingRate = 16000, windowLength = 25, step = 5)
ms = specToMS(spec)
image(x = as.numeric(colnames(ms)), y = as.numeric(rownames(ms)),
      z = t(log(abs(ms))), xlab = 'Amplitude modulation, Hz',
      ylab = 'Frequency modulation, cycles/kHz')
abline(h = 0, lty = 3); abline(v = 0, lty = 3)
```

spectrogram *Spectrogram*

Description

Produces the spectrogram of a sound using short-time Fourier transform. Inspired by [spectro](#), this function offers added routines for reassignment, noise reduction, smoothing in time and frequency domains, manual control of contrast and brightness, plotting the oscillogram on a dB scale, grid, etc.

Usage

```
spectrogram(  
  x,  
  samplingRate = NULL,  
  scale = NULL,  
  from = NULL,  
  to = NULL,  
  dynamicRange = 80,  
  windowLength = 50,  
  step = windowLength/2,  
  overlap = NULL,  
  specType = c("spectrum", "reassigned", "spectralDerivative")[1],  
  rasterize = FALSE,  
  wn = "gaussian",  
  zp = 0,  
  normalize = TRUE,  
  smoothFreq = 0,  
  smoothTime = 0,  
  qTime = 0,  
  percentNoise = 10,  
  noiseReduction = 0,  
  output = c("original", "processed", "complex", "all")[1],  
  reportEvery = NULL,  
  cores = 1,  
  plot = TRUE,  
  savePlots = NULL,  
  osc = c("none", "linear", "dB")[2],  
  heights = c(3, 1),  
  ylim = NULL,  
  yScale = c("linear", "log", "bark", "mel", "ERB")[1],  
  contrast = 0.2,  
  brightness = 0,  
  blur = 0,  
  maxPoints = c(1e+05, 5e+05),  
  padWithSilence = TRUE,  
  colorTheme = c("bw", "seewave", "heat.colors", "...")[1],  
  col = NULL,  
  extraContour = NULL,  
  xlab = NULL,  
  ylab = NULL,  
  xaxp = NULL,  
  mar = c(5.1, 4.1, 4.1, 2),  
  main = NULL,  
  grid = NULL,  
  width = 900,  
  height = 500,  
  units = "px",  
  res = NA,
```

```
    ...
  )
```

Arguments

<code>x</code>	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
<code>samplingRate</code>	sampling rate of <code>x</code> (only needed if <code>x</code> is a numeric vector)
<code>scale</code>	maximum possible amplitude of input used for normalization of input vector (only needed if <code>x</code> is a numeric vector)
<code>from, to</code>	if NULL (default), analyzes the whole sound, otherwise from...to (s)
<code>dynamicRange</code>	dynamic range, dB. All values more than one dynamicRange under maximum are treated as zero
<code>windowLength</code>	length of FFT window, ms
<code>step</code>	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
<code>overlap</code>	overlap between successive FFT frames, %
<code>specType</code>	plot the original FFT ('spectrum'), reassigned spectrogram ('reassigned'), or spectral derivative ('spectralDerivative')
<code>rasterize</code>	(only applies if <code>specType</code> = 'reassigned') if TRUE, the reassigned spectrogram is plotted after rasterizing it: that is, showing density per time-frequency bins with the same resolution as an ordinary spectrogram
<code>wn</code>	window type accepted by <code>ftwindow</code> , currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop
<code>zp</code>	window length after zero padding, points
<code>normalize</code>	if TRUE, scales input prior to FFT
<code>smoothFreq, smoothTime</code>	length of the window for median smoothing in frequency and time domains, respectively, points
<code>qTime</code>	the quantile to be subtracted for each frequency bin. For ex., if <code>qTime</code> = 0.5, the median of each frequency bin (over the entire sound duration) will be calculated and subtracted from each frame (see examples)
<code>percentNoise</code>	percentage of frames (0 to 100%) used for calculating noise spectrum
<code>noiseReduction</code>	how much noise to remove (non-negative number, recommended 0 to 2). 0 = no noise reduction, 2 = strong noise reduction: $spectrum - (noiseReduction * noiseSpectrum)$, where <code>noiseSpectrum</code> is the average spectrum of frames with entropy exceeding the quantile set by <code>percentNoise</code>
<code>output</code>	specifies what to return: nothing ('none'), unmodified spectrogram ('original'), denoised and/or smoothed spectrogram ('processed'), or unmodified spectrogram with the imaginary part giving phase ('complex')
<code>reportEvery</code>	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)

cores	number of cores for parallel processing
plot	should a spectrogram be plotted? TRUE / FALSE
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
osc	"none" = no oscillogram; "linear" = on the original scale; "dB" = in decibels
heights	a vector of length two specifying the relative height of the spectrogram and the oscillogram (including time axes labels)
ylim	frequency range to plot, kHz (defaults to 0 to Nyquist frequency). NB: still in kHz, even if yScale = bark, mel, or ERB
yScale	scale of the frequency axis: 'linear' = linear, 'log' = logarithmic (musical), 'bark' = bark with hz2bark , 'mel' = mel with hz2mel , 'ERB' = Equivalent Rectangular Bandwidths with HzToERB
contrast	spectrum is exponentiated by contrast (any real number, recommended -1 to +1). Contrast >0 increases sharpness, <0 decreases sharpness
brightness	how much to "lighten" the image (>0 = lighter, <0 = darker)
blur	apply a Gaussian filter to blur or sharpen the image, two numbers: frequency (Hz), time (ms). A single number is interpreted as frequency, and a square filter is applied. NA / NULL / 0 means to blurring in that dimension. Negative numbers mean un-blurring (sharpening) the image by dividing instead of multiplying by the filter during convolution
maxPoints	the maximum number of "pixels" in the oscillogram (if any) and spectrogram; good for quickly plotting long audio files; defaults to c(1e5, 5e5)
padWithSilence	if TRUE, pads the sound with just enough silence to resolve the edges properly (only the original region is plotted, so the apparent duration doesn't change)
colorTheme	black and white ('bw'), as in seewave package ('seewave'), or any palette from palette such as 'heat.colors', 'cm.colors', etc
col	actual colors, eg rev(rainbow(100)) - see ?hcl.colors for colors in base R (overrides colorTheme)
extraContour	a vector of arbitrary length scaled in Hz (regardless of yScale!) that will be plotted over the spectrogram (eg pitch contour); can also be a list with extra graphical parameters such as lwd, col, etc. (see examples)
xlab, ylab, main, mar, xaxp	graphical parameters for plotting
grid	if numeric, adds n = grid dotted lines per kHz
width, height, units, res	graphical parameters for saving plots passed to png
...	other graphical parameters

Details

Many soundgen functions call `spectrogram`, and you can pass along most of its graphical parameters from functions like [soundgen](#), [analyze](#), etc. However, in some cases this will not work (eg for "units") or may produce unexpected results. If in doubt, omit extra graphical parameters.

Value

Returns nothing (if output = 'none'), absolute - not power! - spectrum (if output = 'original'), denoised and/or smoothed spectrum (if output = 'processed'), or spectral derivatives (if specType = 'spectralDerivative') as a matrix of real numbers.

See Also

[osc modulationSpectrum ssm](#)

Examples

```
# synthesize a sound 500 ms long, with gradually increasing hissing noise
sound = soundgen(syllen = 500, temperature = 0.001, noise = list(
  time = c(0, 650), value = c(-40, 0)), formantsNoise = list(
  f1 = list(freq = 5000, width = 10000)))
# playme(sound, samplingRate = 16000)

# basic spectrogram
spectrogram(sound, samplingRate = 16000, yScale = 'bark')

# add bells and whistles
spectrogram(sound, samplingRate = 16000,
  osc = 'dB', # plot oscillogram in dB
  heights = c(2, 1), # spectro/osc height ratio
  noiseReduction = 1.1, # subtract the spectrum of noisy parts
  brightness = -1, # reduce brightness
  # pick color theme - see ?hcl.colors
  # colorTheme = 'heat.colors',
  # ...or just specify the actual colors
  col = colorRampPalette(c('white', 'yellow', 'red'))(50),
  cex.lab = .75, cex.axis = .75, # text size and other base graphics pars
  grid = 5, # lines per kHz; to customize, add manually with graphics::grid()
  ylim = c(0, 5), # always in kHz
  main = 'My spectrogram' # title
  # + axis labels, etc
)
## Not run:
# save spectrograms of all sounds in a folder
spectrogram('~Downloads/temp', savePlots = '', cores = 2)

# change dynamic range
spectrogram(sound, samplingRate = 16000, dynamicRange = 40)
spectrogram(sound, samplingRate = 16000, dynamicRange = 120)

# remove the oscillogram
spectrogram(sound, samplingRate = 16000, osc = 'none') # or NULL etc

# frequencies on a logarithmic (musical) scale (mel/bark also available)
spectrogram(sound, samplingRate = 16000,
  yScale = 'log', ylim = c(.05, 8))

# broad-band instead of narrow-band
```

```

spectrogram(sound, samplingRate = 16000, windowLength = 5)

# reassigned spectrograms can be plotted without rasterizing, as a
# scatterplot instead of a contour plot
s = soundgen(syllen = 500, pitch = c(100, 1100, 120, 1200, 90, 900, 110, 700),
  samplingRate = 22050, formants = NULL, lipRad = 0, rolloff = -20)
spectrogram(s, 22050, windowLength = 5, step = 1, ylim = c(0, 2))
spectrogram(s, 22050, specType = 'reassigned', windowLength = 5,
  step = 1, ylim = c(0, 2))
# ...or it can be rasterized, but that sacrifices frequency resolution:
sp = spectrogram(s, 22050, specType = 'reassigned', rasterize = TRUE,
  windowLength = 5, step = 1, ylim = c(0, 2), output = 'all')
# The raw reassigned version is saved if output = 'all' for custom plotting
df = sp$reassigned
df$z1 = soundgen::zeroOne(log(df$magn))
plot(df$time, df$freq, col = rgb(df$z1, df$z1, 1 - df$z1, 1),
  pch = 16, cex = 0.25, ylim = c(0, 2))

# focus only on values in the upper 5% for each frequency bin
spectrogram(sound, samplingRate = 16000, qTime = 0.95)

# detect 10% of the noisiest frames based on entropy and remove the pattern
# found in those frames (in this cases, breathing)
spectrogram(sound, samplingRate = 16000, noiseReduction = 1.1,
  brightness = -2) # white noise attenuated

# increase contrast, reduce brightness
spectrogram(sound, samplingRate = 16000, contrast = .7, brightness = -.5)

# apply median smoothing in both time and frequency domains
spectrogram(sound, samplingRate = 16000, smoothFreq = 5,
  smoothTime = 5)

# Gaussian filter to blur or sharpen ("unblur") the image in time and/or
# frequency domains
spectrogram(sound, samplingRate = 16000, blur = c(100, 500))
# TIP: when unblurring, set the first (frequency) parameter to the
# frequency resolution of interest, eg ~500-1000 Hz for human formants
spectrogram(sound, samplingRate = 16000, windowLength = 10, blur = c(-500, 500))

# specify location of tick marks etc - see ?par() for base graphics
spectrogram(sound, samplingRate = 16000,
  ylim = c(0, 3), yaxp = c(0, 3, 5), xaxp = c(0, .8, 10))

# Plot long audio files with reduced resolution
data(sheep, package = 'seewave')
sp = spectrogram(sheep, overlap = 0,
  maxPoints = c(1e4, 5e3), # limit the number of pixels in osc/spec
  output = 'original')
nrow(sp) * ncol(sp) / 5e3 # spec downsampled by a factor of ~2

# Plot some arbitrary contour over the spectrogram (simply calling lines())
# will not work if osc = TRUE b/c the plot layout is modified)

```

```

s = soundgen()
an = analyze(s, 16000, plot = FALSE)
spectrogram(s, 16000, extraContour = an$detailed$dom,
  ylim = c(0, 2), yScale = 'bark')
# For values that are not in Hz, normalize any way you like
spectrogram(s, 16000, ylim = c(0, 2), extraContour = list(
  x = an$detailed$loudness / max(an$detailed$loudness, na.rm = TRUE) * 2000,
  # ylim[2] = 2000 Hz
  type = 'b', pch = 5, lwd = 2, lty = 2, col = 'blue'))

## End(Not run)

```

ssm

Self-similarity matrix

Description

Calculates the self-similarity matrix and novelty vector of a sound.

Usage

```

ssm(
  x,
  samplingRate = NULL,
  from = NULL,
  to = NULL,
  windowLength = 25,
  step = 5,
  overlap = NULL,
  ssmWin = NULL,
  sparse = FALSE,
  maxFreq = NULL,
  nBands = NULL,
  MFCC = 2:13,
  input = c("mfcc", "melspec", "spectrum")[2],
  norm = FALSE,
  simil = c("cosine", "cor")[1],
  kernelLen = 100,
  kernelSD = 0.5,
  padWith = 0,
  summaryFun = c("mean", "sd"),
  reportEvery = NULL,
  cores = 1,
  plot = TRUE,
  savePlots = NULL,
  main = NULL,
  heights = c(2, 1),
  width = 900,

```



```

height = 500,
units = "px",
res = NA,
specPars = list(levels = seq(0, 1, length = 30), colorTheme = c("bw", "seewave",
  "heat.colors", "...")[2], xlab = "Time, s", ylab = "kHz"),
ssmPars = list(levels = seq(0, 1, length = 30), colorTheme = c("bw", "seewave",
  "heat.colors", "...")[2], xlab = "Time, s", ylab = "Time, s"),
noveltyPars = list(type = "b", pch = 16, col = "black", lwd = 3)
)

```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
samplingRate	sampling rate of x (only needed if x is a numeric vector)
from, to	if NULL (default), analyzes the whole sound, otherwise from...to (s)
windowLength	length of FFT window, ms
step	you can override overlap by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of 1/samplingRate, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
overlap	overlap between successive FFT frames, %
ssmWin	window for averaging SSM, ms (has a smoothing effect and speeds up the processing)
sparse	if TRUE, the entire SSM is not calculated, but only the central region needed to extract the novelty contour (speeds up the processing)
maxFreq	highest band edge of mel filters, Hz. Defaults to samplingRate / 2. See melfcc
nBands	number of warped spectral bands to use. Defaults to 100 * windowLength / 20. See melfcc
MFCC	which mel-frequency cepstral coefficients to use; defaults to 2:13
input	the spectral representation used to calculate the SSM
norm	if TRUE, the spectrum of each STFT frame is normalized
simil	method for comparing frames: "cosine" = cosine similarity, "cor" = Pearson's correlation
kernellLen	length of checkerboard kernel for calculating novelty, ms (larger values favor global, slow vs. local, fast novelty)
kernelSD	SD of checkerboard kernel for calculating novelty
padWith	how to treat edges when calculating novelty: NA = treat sound before and after the recording as unknown, 0 = treat it as silence
summaryFun	functions used to summarize each acoustic characteristic, eg "c('mean', 'sd')"; user-defined functions are fine (see examples); NAs are omitted automatically for mean/median/sd/min/max/range/sum, otherwise take care of NAs yourself
reportEvery	when processing multiple inputs, report estimated time left every ... iterations (NULL = default, NA = don't report)

cores	number of cores for parallel processing
plot	if TRUE, plots the SSM
savePlots	full path to the folder in which to save the plots (NULL = don't save, "" = same folder as audio)
main	plot title
heights	relative sizes of the SSM and spectrogram/novelty plot
width, height, units, res	graphical parameters for saving plots passed to png
specPars	graphical parameters passed to <code>filled.contour.mod</code> and affecting the spectrogram
ssmPars	graphical parameters passed to <code>filled.contour.mod</code> and affecting the plot of SSM
noveltyPars	graphical parameters passed to lines and affecting the novelty contour

Value

Returns a list of two components: `$ssm` contains the self-similarity matrix, and `$novelty` contains the novelty vector.

References

- El Badawy, D., Marmaroli, P., & Lissek, H. (2013). Audio Novelty-Based Segmentation of Music Concerts. In *Acoustics 2013* (No. EPFL-CONF-190844)
- Foote, J. (1999, October). Visualizing music and audio using self-similarity. In *Proceedings of the seventh ACM international conference on Multimedia (Part 1)* (pp. 77-80). ACM.
- Foote, J. (2000). Automatic audio segmentation using a measure of audio novelty. In *Multimedia and Expo, 2000. ICME 2000. 2000 IEEE International Conference on* (Vol. 1, pp. 452-455). IEEE.

See Also

[spectrogram](#) [modulationSpectrum](#) [segment](#)

Examples

```

sound = c(soundgen(),
          soundgen(nSyl = 4, sylLen = 50, pauseLen = 70,
                  formants = NA, pitch = c(500, 330)))
# playme(sound)
# detailed, local features (captures each syllable)
s1 = ssm(sound, samplingRate = 16000, kernelLen = 100,
         sparse = TRUE) # much faster with 'sparse'
# more global features (captures the transition b/w the two sounds)
s2 = ssm(sound, samplingRate = 16000, kernelLen = 400, sparse = TRUE)

s2$summary
s2$novelty # novelty contour
## Not run:

```

```

ssm(sound, samplingRate = 16000,
     input = 'mfcc', simil = 'cor', norm = TRUE,
     ssmWin = 25, # speed up the processing
     kernelLen = 300, # global features
     specPars = list(colorTheme = 'seewave'),
     ssmPars = list(col = rainbow(100)),
     noveltyPars = list(type = 'l', lty = 3, lwd = 2))

## End(Not run)

```

timeStretch

Time stretch

Description

Dynamically time-stretches a sound without preserving its pitch or formants, as if gradually changing playback speed. Algorithm: the audio is resampled at time-varying steps. This is about 100 times faster than time-stretching with a phase vocoder in [shiftPitch](#), but pitch and formants cannot be preserved, and large stretch factors may cause artifacts due to aliasing.

Usage

```

timeStretch(
  x,
  stretch = 1,
  samplingRate = NULL,
  precision = 1000,
  play = FALSE,
  saveAudio = NULL,
  reportEvery = NULL,
  cores = 1
)

```

Arguments

x	path to a folder, one or more wav or mp3 files c('file1.wav', 'file2.mp3'), Wave object, numeric vector, or a list of Wave objects or numeric vectors
stretch	1 = no change, >1 = longer, <1 = shorter. Single value, vector, or anchor format (see soundgen)
samplingRate	sampling rate of x (only needed if x is a numeric vector)
precision	the number of points used for estimating the duration of output (more = better, but slower)
play	if TRUE, plays the synthesized sound using the default player on your system. If character, passed to play as the name of player to use, eg "aplay", "play", "vlc", etc. In case of errors, try setting another default player for play
saveAudio	full path to the folder in which to save audio files (one per detected syllable)

reportEvery when processing multiple inputs, report estimated time left every ... iterations
 (NULL = default, NA = don't report)

cores number of cores for parallel processing

See Also

[shiftPitch](#)

Examples

```
data(sheep, package = 'seewave') # import a recording from seewave
# playme(sheep)
# spectrogram(sheep)
s1 = timeStretch(sheep, stretch = c(1, 3))
# playme(s1, sheep@samp.rate)
# spectrogram(s1, sheep@samp.rate)

# compare to a similar effect achieved with a phase vocoder in pitchShift():
s2 = shiftPitch(
  sheep,
  timeStretch = c(1, 3), # from 1 (original) to mult
  multPitch = c(1, 1/3), # also drop pitch
  multFormants = c(1, 1/3) # also drop formants (by the same proportion)
)
# playme(s2, sheep@samp.rate)
# spectrogram(s2, sheep@samp.rate)
# NB: because the two algorithms calculate transitions between stretch
# factors in different ways, the duration is not identical, even though the
# range of pitch change is the same
```

transplantEnv

Transplant envelope

Description

Extracts a smoothed amplitude envelope of the donor sound and applies it to the recipient sound. Both sounds are provided as numeric vectors; they can differ in length and sampling rate. Note that the result depends on the amount of smoothing (controlled by `windowLength`) and the chosen method of calculating the envelope. Very similar to [setenv](#), but with a different smoothing algorithm and with a choice of several types of envelope: `hil`, `rms`, or `peak`.

Usage

```
transplantEnv(
  donor,
  samplingRateD = NULL,
  recipient,
  samplingRateR = samplingRateD,
  windowLength = 50,
```

```

method = c("hil", "rms", "peak")[3],
killDC = FALSE,
dynamicRange = 80,
plot = FALSE
)

```

Arguments

donor	the sound that "donates" the amplitude envelope
samplingRateD, samplingRateR	sampling rate of the donor and recipient, respectively (only needed for vectors, not files)
recipient	the sound that needs to have its amplitude envelope adjusted
windowLength	the length of smoothing window, ms
method	hil = Hilbert envelope, rms = root mean square amplitude, peak = peak amplitude per window
killDC	if TRUE, dynamically removes DC offset or similar deviations of average waveform from zero (see examples)
dynamicRange	parts of sound quieter than $-\text{dynamicRange}$ dB will not be amplified
plot	if TRUE, plots the original sound, the smoothed envelope, and the compressed sound

Value

Returns the recipient sound with the donor's amplitude envelope - a numeric vector with the same sampling rate as the recipient

See Also

[flatEnv](#) [setenv](#)

Examples

```

donor = rnorm(500) * seq(1, 0, length.out = 500)
recipient = soundgen(syllLen = 600, addSilence = 50)
transplantEnv(donor, samplingRateD = 200,
              recipient, samplingRateR = 16000,
              windowLength = 50, method = 'hil', plot = TRUE)
transplantEnv(donor, samplingRateD = 200,
              recipient, samplingRateR = 16000,
              windowLength = 10, method = 'peak', plot = TRUE)

```

 transplantFormants *Transplant formants*

Description

Takes the general spectral envelope of one sound (donor) and "transplants" it onto another sound (recipient). For biological sounds like speech or animal vocalizations, this has the effect of replacing the formants in the recipient sound while preserving the original intonation and (to some extent) voice quality. Note that `freqWindow` is a crucial parameter: too narrow, and noise between harmonics will be amplified, creating artifacts; too wide, and formants may be missed. The default is to set `freqWindow` to the estimated median pitch, but this is time-consuming and error-prone, so set it to a reasonable value manually if possible. Also ensure that both sounds have the same sampling rate.

Usage

```
transplantFormants(
    donor,
    recipient,
    samplingRate = NULL,
    freqWindow = NULL,
    dynamicRange = 80,
    windowLength = 50,
    step = NULL,
    overlap = 90,
    wn = "gaussian",
    zp = 0
)
```

Arguments

<code>donor</code>	the sound that provides the formants (vector, Wave, or file) or the desired spectral filter (matrix) as returned by getSpectralEnvelope
<code>recipient</code>	the sound that receives the formants (vector, Wave, or file)
<code>samplingRate</code>	sampling rate of x (only needed if x is a numeric vector)
<code>freqWindow</code>	the width of smoothing window. Defaults to median pitch of the donor (or of the recipient if donor is a filter matrix)
<code>dynamicRange</code>	dynamic range, dB. All values more than one <code>dynamicRange</code> under maximum are treated as zero
<code>windowLength</code>	length of FFT window, ms
<code>step</code>	you can override <code>overlap</code> by specifying FFT step, ms (NB: because digital audio is sampled at discrete time intervals of $1/\text{samplingRate}$, the actual step and thus the time stamps of STFT frames may be slightly different, eg 24.98866 instead of 25.0 ms)
<code>overlap</code>	overlap between successive FFT frames, %

wn window type accepted by `ftwindow`, currently gaussian, hanning, hamming, bartlett, rectangular, blackman, flattop

zp window length after zero padding, points

Details

Algorithm: makes spectrograms of both sounds, interpolates and smoothes the donor spectrogram, flattens the recipient spectrogram, multiplies the spectrograms, and transforms back into time domain with inverse STFT.

See Also

[transplantEnv](#) [getSpectralEnvelope](#) [addFormants](#) [soundgen](#)

Examples

```
## Not run:
# Objective: take formants from the bleating of a sheep and apply them to a
# synthetic sound with any arbitrary duration, intonation, nonlinearities etc
data(sheep, package = 'seewave') # import a recording from seewave
playme(sheep)
spectrogram(sheep, osc = TRUE)

recipient = soundgen(
  sylLen = 1200,
  pitch = c(100, 300, 250, 200),
  vibratoFreq = 9, vibratoDep = 1,
  addSilence = 180,
  samplingRate = sheep@samp.rate, # same as donor
  invalidArgAction = 'ignore') # force to keep the low samplingRate
playme(recipient, sheep@samp.rate)
spectrogram(recipient, sheep@samp.rate, osc = TRUE)

s1 = transplantFormants(
  donor = sheep,
  recipient = recipient,
  samplingRate = sheep@samp.rate)
playme(s1, sheep@samp.rate)
spectrogram(s1, sheep@samp.rate, osc = TRUE)

# The spectral envelope of s1 will be similar to sheep's on a frequency scale
# determined by freqWindow. Compare the spectra:
par(mfrow = c(1, 2))
seewave::meanspec(sheep, dB = 'max0', alim = c(-50, 20), main = 'Donor')
seewave::meanspec(s1, f = sheep@samp.rate, dB = 'max0',
  alim = c(-50, 20), main = 'Processed recipient')
par(mfrow = c(1, 1))

# if needed, transplant amplitude envelopes as well:
s2 = transplantEnv(donor = sheep, samplingRateD = sheep@samp.rate,
  recipient = s1, windowLength = 10)
playme(s2, sheep@samp.rate)
```

```
spectrogram(s2, sheep@samp.rate, osc = TRUE)

# Now we use human formants on sheep source: the sheep asks "why?"
s3 = transplantFormants(
  donor = getSpectralEnvelope(
    nr = 512, nc = 100, # fairly arbitrary dimensions
    formants = 'uaaai',
    samplingRate = sheep@samp.rate),
  recipient = sheep,
  samplingRate = sheep@samp.rate)
playme(s3, sheep@samp.rate)
spectrogram(s3, sheep@samp.rate, osc = TRUE)

## End(Not run)
```


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