Using Audacity for bioacoustic analysis

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1 Opening files in Audacity

Download the file C_vagans.wav from http://audioblast.org/keys and open it in Audacity (File: Open). Depending on your computer's operating system and browser you may have to right click on the link to the file to save it to your computer.

When opening files in Audacity you will be asked whether you want to work on the original file, or make a copy. As this file forms part of a training course it is safe to use the original file, but if working on your own files it is always safer to make a copy (Figure 1).

● ○ ○ Warning	
When importing uncompressed audio files you can either copy them into the project, or read them directly from their current location (without copying).	
Your current preference is set to read directly.	
Reading the files directly allows you to play or edit them almost immediately. This is less safe than copying in, because you must retain the files with their original names in their original location. File > Check Dependencies will show the original names and location of any files that you are reading directly. How do you want to import the current file(s)? Choose an import method	
Make a convict the files before editing (actor)	
 Read the files directly from the original (faster) Don't warn again and always use my choice above 	
Cancel	ОК

Figure 1:

Figure 2 shows the file C_vagans.wav opened in Audacity. The default view in Audacity is the *time domain* or *waveform* view. This shows how the amplitude ("volume") of the sound varies with time.



Figure 2:

2 Normalisation

The sound file we have opened does not make full use of the range in amplitude. In order to make individual features of the sound more obvious when analysing them later we can 'normalise' the recording to make full use of the amplitude range available (Effect: Normalize...). The options for the normalisation are shown in Figure 3.

00	Normalize
Remove DC offset (center on 0.0 vertically)	
🗹 Normalize r	naximum amplitude to -1.0 dB
Normalize stereo channels independently	
Preview	Cancel OK

Figure 3:

The effect of the normalisation is illustrated in Figure 4.



Figure 4:

3 The time domain

The time domain view can be used to measure the duration of features in the time domain. Accurate measurement of durations will require us to zoom in on the features of interest. There are several zoom tools located in the toolbar (Figure 5).





Zoom in on a region of the recording and identify a feature to measure the duration of. Use the mouse cursor to select the region of the waveform you wish to measure. Ensure that the selection option is set to Length rather than End (Figure 6). If you are trying to measure features of short duration you will need to ensure that the units selector is set to the option with milliseconds (Figure 7). The duration of the selection, in the units you selected, are shown in the text box under the End/Length selectors.



Figure 6:



Figure 7:

Many insect sounds are produced by stridulation: rubbing a series or raised areas or protrusions (the file) against a plectrum (often an enlarged wing vein). Each back-and-forth motion of the file against the plectrum results in a syllable, syllables may be grouped into groups ("echemes") or form a continuous song. More complex songs may be composed of echeme sequences. There may therefore be many different features to measure the duration of, often with very different durations.

4 The frequency domain

While the time domain view allows for easy measurement of the duration of features, it does not allow for the measurement of the frequency of the song. In order to do examine this we must switch the view of Audacity from 'waveform' to 'spectrogram'. This is illustrated in Figure 8.





A comparison of the waveform (time domain) and spectrogram (frequency domain)

views is given in Figure 9. The spectrogram shows frequency on the vertical axis, regions of brighter colouration show a greater amplitude for that frequency.





The spectrogram view is good for getting an overview of the frequencies present in a song, but it is hard to analyse directly. To analyse the frequency, select the region to analyse, then click Analyse: Plot Spectrum... Figure 10. The Plot Spectrum plots frequency (x-axis) against the amplitude of that frequency (y-axis). It can be seen that for this species the distribution is bimodal, and that this can be related to the spectrogram view. Moving your cursor over the plot allows individual values to be read.



Figure 10:

You can see that for this species the song covers a wide range of frequencies, this known as a *broadband* signal and is typical in grasshoppers and bushcrickets. The true crickets have a much narrower frequency spectrum (narrowband signal) as their song is modified by resonators (special regions of the wing). For comparison the song of the mole cricket *Gryllotalpa vineae* is given in Figure 11.



Figure 11: