A short introduction to acoustic template matching with monitoR

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1 Getting started

The motivation behind the development of the monitoR package was automated detection and identification of animal vocalizations. Here, we describe how to use monitoR functions to detect and identify songs from two bird species from a survey recording. We'll start by loading the package.

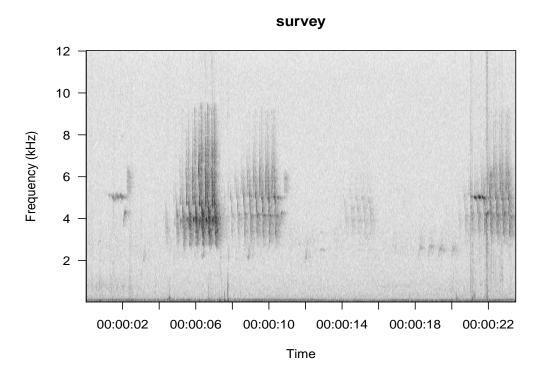
library(monitoR)

The monitoR package includes a 23 second recording of songbirds as a Wave object (defined in the tuneR package) named survey.

data(survey) survey # # Wave Object # Number of Samples: 564000 # Duration (seconds): 23.5 # Samplingrate (Hertz): 24000 # Channels (Mono/Stereo): Mono PCM (integer format): # TRUE Bit (8/16/24/32/64): # 16

A spectrogram of the recordings can be generated using the viewSpec function¹.

```
viewSpec(survey)
```



We could also play the recording using the play function from the tuneR package².

```
setWavPlayer("play")
play(survey)
```

You might be see two different types of songs here. The songs around 2, 10, and maybe 22 seconds are from a black-throated green warbler *Setophaga virens*, and there are ovenbird *Seiurus aurocapilla* songs around 6, 9, 15, and 23 seconds. In the following, we show how monitoR can be used to detect and identify these songs.

The monitoR package also includes short recordings of songs from the same two species in survey: btnw, which contains a single song from a black-throated green warbler, and oven which contains a single song from an ovenbird. As with survey, these objects are Wave objects.

```
data(btnw)
data(oven)
btnw
```

#

 $^{^{1}}$ By default, this function just transforms the data and generates a spectrogram. Try setting the interactive argument to TRUE for more viewing options that are useful for longer recordings.

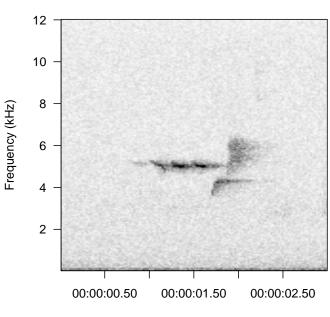
² This requires that a command-line wav player is installed. Here, we are using SoX.

```
# Wave Object
# Number of Samples: 72001
# Duration (seconds): 3
# Samplingrate (Hertz): 24000
# Channels (Mono/Stereo): Mono
# PCM (integer format): TRUE
# Bit (8/16/24/32/64): 16
```

oven

Ŧ		
#	Wave Object	
#	Number of Samples:	120001
#	Duration (seconds):	5
#	Samplingrate (Hertz):	24000
#	Channels (Mono/Stereo):	Mono
#	PCM (integer format):	TRUE
#	Bit (8/16/24/32/64):	16

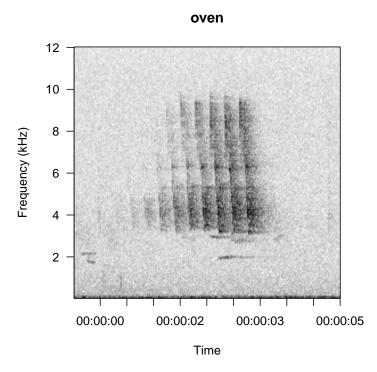
viewSpec(btnw)



btnw

Time

viewSpec(oven)



As Wave objects, all of these recordings can be used directly in the template matching functions³. Users will typically work with wav files, however, as opposed to mp3 files, or Wave objects. To be consistent with this typical approach, we will save the two clips and the survey as wav files⁴. We'll include a made-up date in the name for the survey file, since that is one way to determine "absolute" times for detections⁵.

```
btnw.fp <- file.path(tempdir(), "btnw.wav")
oven.fp <- file.path(tempdir(), "oven.wav")
survey.fp <- file.path(tempdir(), "survey2010-12-31_120000_EST.wav")
writeWave(btnw, btnw.fp)
writeWave(oven, oven.fp)
writeWave(survey, survey.fp)</pre>
```

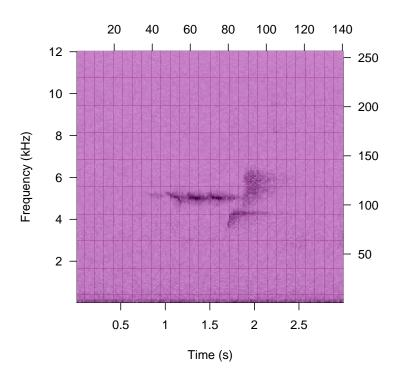
2 Spectrogram cross correlation

The monitoR package provides functions for two kinds of template detection-spectrogram cross correlation and binary point matching. We'll start with the correlation approach, which is slightly simpler. The easiest way to create a correlation template is to call up makeCorTemplate with only the clip argument specified. This function carries out a Fourier transform, and, by default, saves amplitude (volume) data for each cell within the resulting frequency-by-time matrix.

 $^{^3}$ But you must set the write.wav argument to TRUE to do this.

⁴ Here, we will use a temporary directory, but for your own recordings a more permanent location is better.

 $^{^{5}}$ The other option is to use the modification date of the survey file, which clearly would be wrong here, since we just created it.



```
#
# Automatic point selection.
#
# Done.
```

The makeCorTemplate function displays a spectrogram of the recording⁶, and shows the cells included in the template (all the cells here) in transparent purple⁷. Let's look at a summary of the template we just created⁸.

wct1

```
#
# Object of class "corTemplateList"
#
   containing 1 templates
#
          original.recording sample.rate lower.frequency
#
 A /tmp/RtmpCXW6Mo/btnw.wav
                                    24000
                                                     0.047
#
    upper.frequency lower.amp upper.amp duration n.points
# A
                 12
                        -97.52
                                       0
                                            2.965
                                                      35840
#
    score.cutoff
# A
             0.4
```

 $^{^{6}}$ This plot is more important when templates are produced interactively, as described below.

⁷ This color can be selected with the argument sel.col.

 $^{^{8}}$ Just entering the template list name will call up the appropriate show method, which just displays a simple summary.

The w1 object is a template list, of class corTemplateList. Here the idea of a list doesn't make a lot of sense, since we only have one template, but in monitoR, templates only exist as part of template lists. Below, we'll create template lists with multiple templates.

The name associated with our single template in the list is A, which is the default, and is not very descriptive. The name of a template can be set when it is created, using the name argument⁹¹⁰.

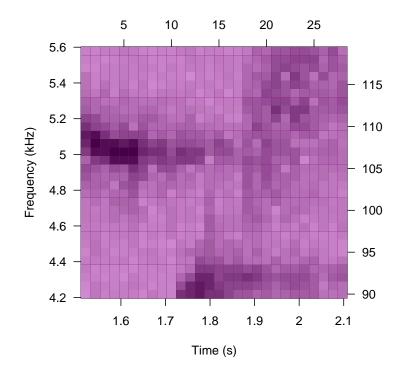
```
wct1 <- makeCorTemplate(btnw.fp, name = "w1")</pre>
#
# Automatic point selection.
#
# Done.
wct1
#
# Object of class "corTemplateList"
#
  containing 1 templates
           original.recording sample.rate lower.frequency
#
# w1 /tmp/RtmpCXW6Mo/btnw.wav 24000
                                                    0.047
     upper.frequency lower.amp upper.amp duration n.points
#
                  12
# w1
                        -97.52
                                       0
                                            2.965
                                                     35840
#
     score.cutoff
# w1
       0.4
```

This template is functional, but it does not take advantage of all the options available in makeCorTemplate. Setting time and frequency limits is one of simplest, and may be necessary when making a template from a long recording. We'll make a new template that uses the t.lim and frq.lim arguments to focus on a particular section within the song.

wct2 <- makeCorTemplate(btnw.fp, t.lim = c(1.5, 2.1), frq.lim = c(4.2, 5.6), name = "w2")</pre>

 $^{^9}$ The templateNames function can be used to change template names in an existing template list. For more detailed information, the templateComment function can be used to check and add comments to templates.

 $^{^{10}}$ In the interest of keeping this file from being too big, the figures produced by the calls in most of the following examples will not be shown. For a complete version of this vignette, visit the monitoR website.



```
#
# Automatic point selection.
#
# Done.
wct2
#
# Object of class "corTemplateList"
#
  containing 1 templates
#
          original.recording sample.rate lower.frequency
# w2 /tmp/RtmpCXW6Mo/btnw.wav 24000
                                                    4.219
#
     upper.frequency lower.amp upper.amp duration n.points
# w2
               5.578
                        -93.52
                                       0
                                            0.576
                                                       840
#
     score.cutoff
              0.4
# w2
```

Let's make two templates with the ovenbird recording as well.

```
oct1 <- makeCorTemplate(oven.fp, t.lim = c(1, 4), frq.lim = c(1, 11), name = "o1")
#
# Automatic point selection.
#
# Done.</pre>
```

For the next template, we'll use the **dens** argument to reduce the size of the template–that is, the number of points included.

```
oct2 <- makeCorTemplate(oven.fp, t.lim = c(1, 4), frq.lim = c(1, 11), dens = 0.1, name = "o2")
#
# Automatic point selection.
#
# Done.</pre>
```

Here, our template will include *approximately* one-tenth of all the spectrogram points.

Template lists with just a single template, can be used for detection. However, monitoR is designed to use template lists that contain multiple templates. Templates can be combined together in a single list with combineCorTemplates.

```
ctemps <- combineCorTemplates(wct1, wct2, oct1, oct2)</pre>
ctemps
#
# Object of class "corTemplateList"
  containing 4 templates
#
#
           original.recording sample.rate lower.frequency
# w1 /tmp/RtmpCXW6Mo/btnw.wav 24000
                                                       0.047
# w2 /tmp/RtmpCXW6Mo/btnw.wav 24000
# o1 /tmp/RtmpCXW6Mo/oven.wav 24000
# o2 /tmp/RtmpCXW6Mo/oven.wav 24000
                                                       4.219
                                                       1.031
                                                       1.031
#
     upper.frequency lower.amp upper.amp duration n.points
# w1
             12.000
                         -97.52 0.00
                                              2.965
                                                      35840
                         -93.52
-85.94
                                      0.00
# w2
               5.578
                                              0.576
                                                          840
# o1
              10.969
                                      0.00 2.965
                                                        29820
                         -71.79 -3.32 2.965
# o2
              10.969
                                                         2955
#
     score.cutoff
# w1
               0.4
               0.4
# w2
# o1
               0.4
               0.4
# o2
```

The output from combineCorTemplates is a template list, just like the output from makeCorTemplate. This function can be used to combine templates present in any number of correlation template lists (and each of the lists can contain any number of templates).

We can view all the templates in **ctemps** with the **plot** function.

plot(ctemps)

We can now use our templates to look for black-throated green warbler and ovenbird songs within the **survey** recording. This is a three-step process: we first calculate correlation score between the templates and each time bin in the survey, then identify "peaks" in the scores, and finally determine which, if any, score peaks exceed the score cutoff. Correlation scores are calculated with **corMatch**.

```
cscores <- corMatch(survey.fp, ctemps)</pre>
#
# Starting w1 . . .
# Fourier transform on survey . . .
#
  Continuing. . .
#
#
  Done.
#
# Starting w2 . . .
# Done.
#
# Starting o1 . . .
# Done.
#
# Starting o2 . . .
# Done.
```

This particular example should run in a few seconds, but corMatch can take much longer for long surveys ¹¹. We can get a summary of the scores object by entering its name, but it requires additional processing in order to be used for detecting songs.

```
cscores
#
# A "templateScores" object
#
# Based on the survey file: /tmp/RtmpCXW6Mo/survey2010-12-31_120000_EST.wav
#
# And 4 templates
# Score information
#
   min.score max.score n.scores
# w1 0.13 0.46 961
      -0.22 0.73 1073
-0.07 0.63 961
# w2
# o1
     -0.10 0.63 961
# o2
```

The corMatch function returns templateScores objects, which contain the correlation scores for each template, but also the complete original templates, along with other information, like the name of the survey file used. To make detections, we need to look within the correlation scores returned by corMatch for peaks, and then identify those peaks with values above the score cutoff as detections. These two steps are carried out with findPeaks.

```
cdetects <- findPeaks(cscores)
#
# Done with w1
# Done with w2</pre>
```

 11 Set the parallel argument to TRUE to speed this step up if you are have a multi-core processor and are not using Windows.

Done with o1
Done with o2
Done

The findPeaks function returns detectionList objects, which contain all the information originally present in templateScores objects, plus detection results. To see a summary of the detections, just enter the name of the object.

```
cdetects
#
# A "detectionList" object
#
# Based on survey file: /tmp/RtmpCXW6Mo/survey2010-12-31_120000_EST.wav
#
# and 4 templates
#
# Detection information
#
    n.peaks n.detections min.peak.score max.peak.score
# w1
        6 1 0.297370075 0.4615955
         38
                     3 0.009121254
# w2
                                            0.7291718
                           0.154263730
0.182259544
# o1
         5
                      3
                                            0.6322787
# o2
                                            0.6267112
         5
                     3
#
    min.detection.score max.detection.score
# w1
             0.4615955
                                 0.4615955
              0.4057619
                                 0.7291718
# w2
              0.4562568
                                 0.6322787
# o1
              0.4523693
# o2
                                 0.6267112
```

From the **n.peaks** column, we can see there are from five to 38 peaks per template, and from the **n.detections** column, we can see that the templates resulted in from one to three detections. The detections can be extracted with the getDetections function.

getDetections(cdetects)

template date.time time score # 1 w1 2010-12-31 11:59:38 1.834667 0.4615955 # 2 w2 2010-12-31 11:59:38 2.133333 0.7291718 w2 2010-12-31 11:59:47 10.709333 0.4057619 # 3 # 4 w2 2010-12-31 11:59:58 21.717333 0.4691309 o1 2010-12-31 11:59:42 6.186667 0.6322787 # 5 # 6 o1 2010-12-31 11:59:45 9.472000 0.5078371 # 7 o1 2010-12-31 11:59:58 21.973333 0.4562568 # 8 o2 2010-12-31 11:59:42 6.144000 0.6267112 # 9 02 2010-12-31 11:59:45 9.450667 0.5158251 # 10 o2 2010-12-31 11:59:58 21.888000 0.4523693

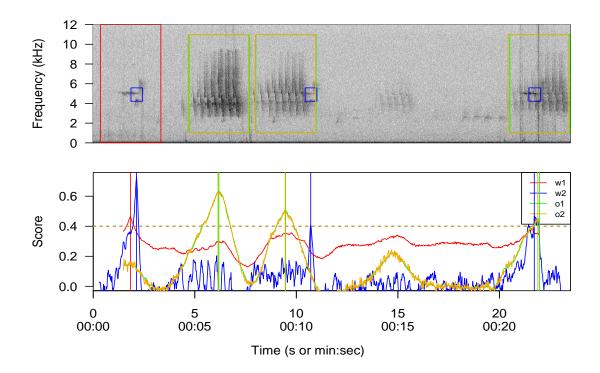
Whether or not a peak qualifies as a detection depends a parameter called the score cutoff, which can be set separately for each template. If we look at our template list again, we can see the score cutoffs given in the last column of the summary. ctemps

#						
#	Obj	ject of class "co	orTemplate	List"		
#	С	ontaining 4 ter	nplates			
#		original.	recording :	sample.rate	e lower.fi	requency
#	w1	/tmp/RtmpCXW6Mo,	/btnw.wav	24000)	0.047
#	w2	/tmp/RtmpCXW6Mo,	/btnw.wav	24000)	4.219
#	o1	/tmp/RtmpCXW6Mo,	/oven.wav	24000)	1.031
#	o2	/tmp/RtmpCXW6Mo,	/oven.wav	24000)	1.031
#		upper.frequency	lower.amp	upper.amp	duration	n.points
#	w1	12.000	-97.52	0.00	2.965	35840
#	w2	5.578	-93.52	0.00	0.576	840
#	o1	10.969	-85.94	0.00	2.965	29820
#	o2	10.969	-71.79	-3.32	2.965	2955
#		score.cutoff				
#	w1	0.4				
#	w2	0.4				
#	o1	0.4				
#	о2	0.4				

They are currently set to the default value, but there is no reason to think that this value should work for all or even any templates. Instead, score cutoffs need to be determined based on template performance, typically with a subset of the survey or surveys that will ultimately be searched for matching sounds. Here we just have one, very short, survey, so we'll use the entire survey to determine what the score cutoff should be set to for each template. We can see a graphical summary of the results with the generic **plot** function.

plot(cdetects)

0 to 23.5 seconds



Starting with the warbler templates, it is easy to see that template w2 is much better than template w1-its scores are high for the three apparent songs, and relatively low elsewhere, including during ovenbird songs. Changing the score cutoff cannot much improve template w1. For template w2, the default value works well enough for this survey. But to incorporate a safety factor, we might want to lower it, perhaps to 0.30.

Looking at the ovenbird results, both templates perform similarly, but if we want to detect songs like the faint one around 15 sec, the score cutoff needs to be reduced, perhaps to 0.20.

We could recreate this template list by repeating the steps above, but set the score.cutoff argument for w2 to 0.30 and 0.20 for o1 and o2. Fortunately, using templateCutoff is a more efficient option. It can be used as an extractor or replacement function.

templateCutoff(ctemps)

w1 w2 o1 o2 # 0.4 0.4 0.4 0.4

To change the cutoff for templates w2, o1, and o2 in the the template list cdetects, we can use:

templateCutoff(ctemps)[2:4] <- c(0.3, 0.2, 0.2)</pre>

Or, this call does the same thing (and is perhaps less prone to mistakes):

templateCutoff(ctemps) <- c(w2 = 0.3, o1 = 0.2, o2 = 0.2)
ctemps</pre>

#

```
# Object of class "corTemplateList"
#
   containing 4 templates
#
             original.recording sample.rate lower.frequency
                                     24000
# w1 /tmp/RtmpCXW6Mo/btnw.wav
                                                               0.047
# w2 /tmp/RtmpCXW6Mo/btnw.wav
                                           24000
                                                               4.219
# o1 /tmp/RtmpCXW6Mo/oven.wav
                                          24000
                                                               1.031
# o2 /tmp/RtmpCXW6Mo/oven.wav
                                           24000
                                                               1.031
#
      upper.frequency lower.amp upper.amp duration n.points
# w1
         12.000 -97.52 0.00 2.965 35840
                           -93.52
                                                                 840
# w2
                5.578
                                         0.00 0.576

        -93.52
        0.00
        0.576
        840

        -85.94
        0.00
        2.965
        29820

        -71.79
        -3.32
        2.965
        2955

# o1
                10.969
# o2
                10.969
#
      score.cutoff
# w1
               0.4
# w2
                 0.3
# o1
                 0.2
# o2
                 0.2
```

Or, we could even use the special name default to specify a default value.

```
templateCutoff(ctemps) <- c(w2 = 0.3, default = 0.2)</pre>
ctemps
#
# Object of class "corTemplateList"
# containing 4 templates
#
           original.recording sample.rate lower.frequency
# w1 /tmp/RtmpCXW6Mo/btnw.wav 24000
                                                      0.047
# w2 /tmp/RtmpCXW6Mo/btnw.wav
                                      24000
                                                       4.219
# o1 /tmp/RtmpCXW6Mo/oven.wav
# o2 /tmp/RtmpCXW6Mo/oven.wav
                                    24000
                                                       1.031
                                    24000
                                                       1.031
#
     upper.frequency lower.amp upper.amp duration n.points
# w1
       12.000
                       -97.52 0.00 2.965 35840
             5.578-93.520.000.57684010.969-85.940.002.9652982010.969-71.79-3.322.9652955
# w2
# o1
# o2
#
     score.cutoff
# w1
         0.2
# w2
              0.3
              0.2
# o1
# 02
              0.2
```

And now we could call corMatch and findPeaks again. This approach would work fine, but takes more time and effort than is needed, because it isn't necessary to recalculate the scores, since they are independent of the cutoffs. Instead, we can just change the score cutoffs in our existing detectionList object cdetects, with the same function templateCutoff. The detections are automatically updated when score cutoffs are changed in a detectionList object.

templateCutoff(cdetects) <- c(w2 = 0.3, default = 0.2)</pre>

So it isn't even necessary to change ctemps¹². Now let's take a look at cdetects.

cdetects

```
#
# A "detectionList" object
#
# Based on survey file: /tmp/RtmpCXW6Mo/survey2010-12-31_120000_EST.wav
#
# and 4 templates
#
# Detection information
#
    n.peaks n.detections min.peak.score max.peak.score
                      6 0.297370075
# w1
         6
                                             0.4615955
# w2
         38
                      3
                            0.009121254
                                             0.7291718
          5
                       4
# o1
                            0.154263730
                                             0.6322787
          5
                       4
                            0.182259544
# 02
                                             0.6267112
#
    min.detection.score max.detection.score
# w1
              0.2973701
                                  0.4615955
# w2
                                  0.7291718
              0.4057619
# o1
              0.2261725
                                  0.6322787
# o2
              0.2437855
                                  0.6267112
```

Notice how the number of detections has changed for o1 and $o2^{13}$.

plot(cdetects)

0 to 23.5 seconds

Since the template w1 is nearly useless, and template o1 and o2 nearly identical, we might want to drop w1 and either o1 or o2 from our results. This can be done using indexing¹⁴.

```
cdetects <- cdetects[c("w2", "o2")]</pre>
cdetects
#
# A "detectionList" object
#
# Based on survey file: /tmp/RtmpCXW6Mo/survey2010-12-31_120000_EST.wav
#
# and 2 templates
#
# Detection information
#
     n.peaks n.detections min.peak.score max.peak.score
# w2
          38
                        3
                             0.009121254
                                               0.7291718
           5
# o2
                        4
                             0.182259544
                                               0.6267112
#
    min.detection.score max.detection.score
```

 $^{^{12}}$ Of course, you may want to reuse or save the template list with optimized cutoffs. The getTemplates function can be helpful here–it can be used to extract the templates from a detectionList object.

 $^{^{13}}$ The plot produced by the call below is omitted, but it would now show a different number of detections.

 $^{^{14}}$ Here, the omitted plot would only show results for w2 and o2.

# w2	0.4057619	0.7291718
# o2	0.2437855	0.6267112

plot(cdetects)

0 to 23.5 seconds

In this short example, it is easy to verify results in the results using the above plot. For longer surveys, for which these methods make more sense, the **showPeaks** function is more efficient. It can be used to view all detections or even all peaks, one by one, and by setting the **verify** argument to TRUE, it can be used to save results of the verification process.

This simple example doesn't show all the available options for creating templates. In particular, with the **select** argument set to "**rectangle**", or "**cell**", it is possible to select areas or even individual cells to be included in a template.

To summarize, for template matching using spectrogram cross-correlation one would typically use the following functions in the order given:

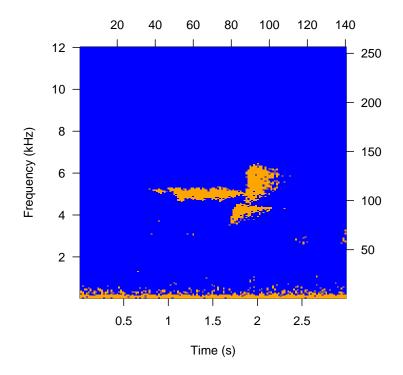
- 1. makeCorTemplate to make the template(s)
- 2. combineCorTemplates to combine templates together in a single template list
- 3. corMatch to calculate scores
- 4. findPeaks to find peaks and identify detections
- 5. plot to see the scores and detections
- 6. getDetections to get the (numeric) detection results
- 7. templateCutoff to change template cutoffs in the detection list object (iteratively with plot and getDetections)

3 Binary point matching

We can use binary point matching to carry out a procedure similar to the one described above in Section 2. Because the type of data needed to calculate scores differ, binary templates differ from correlation templates, and the corresponding functions used for creating templates also differ in some ways. The makeBinTemplate function converts the time-domain data into a binary format by default-i.e., cells are identified as either "high" or "low" (we'll refer to these as "on" and "off") depending on whether they are greater than or less than the user-set amp.cutoff, respectively. The value of amp.cutoff is set interactively by default ¹⁵. With the default option, the spectrogram is updated each time the value of amp.cutoff changes. Here, we'll set the cutoff directly in the function call, based on earlier trials.

wbt1 <- makeBinTemplate(btnw.fp, amp.cutoff = -40, name = "w1")</pre>

¹⁵ This can be changed with the amp.cutoff argument.



```
#
# Automatic point selection.
#
# Done.
```

Time and frequency limits can be set as with makeCorTemplate. We'll also include a buffer around the "on" cells-cells in the buffer are excluded from the template¹⁶.

The amp.cutoff and buffer values used above are based on trial-and-error not shown here. Let's also make two ovenbird templates, with and without a buffer.

```
#
```

 $^{^{16}}$ Although the resulting plot isn't shown here, again, to keep this file size down.

Binary templates are combined with combineBinTemplates.

```
btemps <- combineBinTemplates(wbt1, wbt2, obt1, obt2)</pre>
btemps
#
# Object of class "binTemplateList"
#
#
  containing 4 templates
#
             original.recording sample.rate lower.frequency
# w1 /tmp/RtmpCXW6Mo/btnw.wav 24000 0.046875

        # w2 /tmp/RtmpCXW6Mo/btnw.wav
        24000

        # o1 /tmp/RtmpCXW6Mo/oven.wav
        24000

        # o2 /tmp/RtmpCXW6Mo/oven.wav
        24000

                                                             4.218750
                                                             1.031250
                                                             1.031250
#
      upper.frequency duration on.points off.points score.cutoff
# w1
           12.000000
                              2.97 1845
                                                       33995
                                                                            12
              5.578125
                               0.58
                                            223
                                                         310
                                                                             12
# w2
             10.968750
# o1
                               2.97
                                            2676
                                                        27144
                                                                            12
# o2
             10.968750
                           2.97
                                           1607
                                                        23833
                                                                             12
```

Notice how binary templates have "on" points and "off" points, while correlation templates only have one type of point. Scores are calculated using binMatch, which is analogous to corMatch. As with corMatch, the output from this function is a templateScores object.

```
bscores <- binMatch(survey.fp, btemps)</pre>
#
# Starting w1 . . .
#
#
  Fourier transform on survey . . .
  Continuing. . .
#
#
#
  Done.
#
# Starting w2 . . .
#
#
  Done.
#
```

```
# Starting o1 . . .
#
# Done.
#
# Starting o2 . . .
#
# Done.
```

From this point on, objects created by binary point matching are identical to those made with spectrogram cross correlation. To find detections, use the same findPeaks function that we used above.

```
bdetects <- findPeaks(bscores)
#
# Done with w1
# Done with w2
# Done with o1
# Done with o2
# Done</pre>
```

And, as above, we can use the plot method for detectionList objects to view detections.

plot(bdetects)

0 to 23.5 seconds

Based on these results, it looks like all templates except w1 could be useful with the right score cutoff. We might want to use a cutoff of about 7 for w2, and perhaps 4 for o1 and o2 if the song around 15 seconds should be detected. First, let's drop $w1^{17}$.

```
bdetects <- bdetects[-1]</pre>
```

templateCutoff(bdetects) <- c(w2 = 7, default = 4)</pre>

plot(bdetects)

```
# 0 to 23.5 seconds
```

Finally, the detections are:

getDetections(bdetects)

#		template	(date.time	time	score
#	1	w2	2010-12-31	11:59:38	2.133333	17.583758
#	2	w2	2010-12-31	11:59:47	10.730667	8.669633
#	3	w2	2010-12-31	11:59:58	21.717333	10.490479

 $^{^{17}}$ The plot that is omitted below would no longer show any results for w1.

#	4	o1	2010-12-31	11:59:42	6.208000	18.757236
#	5	o1	2010-12-31	11:59:45	9.450667	11.775177
#	6	o1	2010-12-31	11:59:51	14.698667	4.199115
#	7	o1	2010-12-31	11:59:58	21.952000	10.884734
#	8	o2	2010-12-31	11:59:38	1.536000	4.019856
#	9	o2	2010-12-31	11:59:42	6.208000	22.261260
#	10	o2	2010-12-31	11:59:45	9.472000	14.209354
#	11	о2	2010-12-31	11:59:51	14.720000	5.351632
#	12	o2	2010-12-31	11:59:58	21.952000	13.189866

As with correlation templates, there is much more flexibility in making binary templates than this example suggests. In particular, options for the **select** should be explored. The most flexible is "cell", but it is certainly not the quickest and because individual cells are selected manually, may not be readily repeatable.

To summarize, for template matching using binary point matching one would typically use the following functions in the order given:

- 1. makeBinTemplate to make the template(s)
- 2. combineBinTemplates to combine templates together in a single template list
- 3. binMatch to calculate scores
- 4. findPeaks to find peaks and identify detections
- 5. plot to see the scores and detections
- 6. getDetections to get the (numeric) detection results
- 7. templateCutoff to change template cutoffs in the detection list object (iteratively with plot and getDetections)

4 Other functions

The monitoR package includes several functions not described in this short introduction.

- For manipulating recordings directly, there are the cutWave¹⁸, changeSampRate, and mp3Subsamp functions.
- With the plot methods for template list and detection list objects (used in sections 2 and 3 above), color palettes and other characteristics of the plots can be modified.
- The viewSpec function is capable of much more than the above examples suggest. It can be used interactively to view different parts of a spectrogram, or even annotate it.
- Comments can be added to templates within a template list using the templateComment functions.
- Binary and correlation templates were designed to be portable, and can be saved to text files using writeCorTemplates or writeBinTemplates. Existing template files can be read in with readCorTemplates and writeCorTemplates.

 $^{^{18} \}rm Originally named cutw, but recently changed.$

- For verification of detections, there is the **showPeaks** function, which, shows (or plays) individual detections (or peaks), and, in the interactive mode, allows the user to add verification data to a detection list object.
- The functions collapseClips and bindEvents can be used to extract and bind together detections from a longer recording, and could also be useful for verification.
- The function eventEval provides a way to compare results from template detection to a manually annotated spectrogram, and the function timeAlign removes redundant detections from a survey (e.g. those from multiple templates).
- The compareTemplates function can be used to compare the performance of multiple templates and evaluate similarity between templates.
- An alternative to writing templateLists and detectionLists locally is to store templates, survey metadata, and detection results in a MySQL database. MonitoR has a variety of MySQL queries using package RODBC to push and pull data to an acoustics database: uploads are accomplished with dbUploadSurvey, dbUploadResult, and dbUploadTemplate. Templates can be downloaded to templateList objects with dbDownloadTemplate, and results can be downloaded to detectionList objects with dbDownloadResult. Survey metadata and media card/recorder metadata are downloaded with dbDownloadSurvey and dbDownloadCardRecorderID. An acoustics database schema is available for download at the monitoR website, which can be loaded on an active database instance using dbSchema or incorporated into an existing MySQL database.

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