Animal

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1 Introduction

The Animal package is a collection of functions for analyzing animal behavior data originating from a variety of sources. The package was originally created to analyze data files from CowLog (open source software for coding behaviors from digital video), but the functionality has been extended to cover also other data sources. This document is still very much under development, but the aim is to describe some the key features of the package via simple examples. Animal has been created in the Research Centre of Animal Welfare ¹.

2 CowLog data

The package has basic analysis functions for analyzing time coded behavioral data coded with $CowLog [1]^2$. The main function in the package is cowAnalyze. The function takes the data file name, labels of the codes and the type (event or state) of the codes as inputs, and gives a summary table and plot of the results as output. The function also removes double state (duplicated) errors for state codes.

¹http://http://www.vetmed.helsinki.fi/hyvinvointikeskus/english/index.htm

²http://mpastell.com/CowLog

3 RIC data 2

Here is a short example on how to use the function in R: Analyze CowLog datafile named calf1.bh1: Define codes 1-3 and give them names walking, standing and lying. The descriptive statistics will appear on screen and they are also saved to variable analyzed.

> library(Animal)

> analyzed <- cowAnalyze(file='calf1.bh1',states=c(1,2,3),state.names=c('walking','standing

Event results

Frequency 0 times

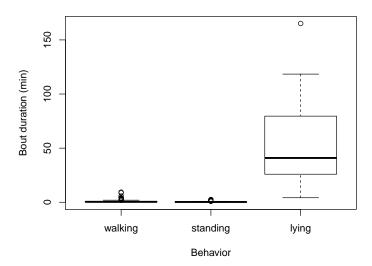
State results walking standing lying

Frequency 270 269 21 times

Bout 0.8 0.37 54.98 minutes

Total 214.81 99.35 1154.62 minutes

Bout Duration of Different Behaviours



3 RIC data

The Animal package has several functions for working with the data produced by Insentec RIC feed measurement system. The basic function read.RIC can be used to read in RIC log files and they can be processed using clean.RIC and bouts.RIC functions.

Example data has been read in by using read.RIC with option clean=F It is included in the package as dataset RIC.

> data(RIC)

3 RIC data

First we clean the data from zero rows and negative feed in takes, then we merge the feeding bouts that are less than 5 minutes apart.

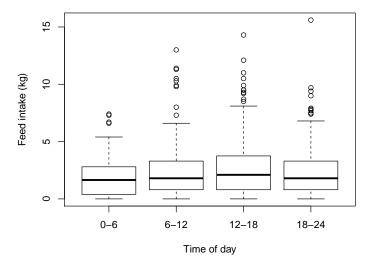
```
> RIC2 <- clean.RIC(RIC)
> bouts <- bouts.RIC(RIC2)
> head(bouts,5)
```

	cowID t	rough		begin		end	intake		
1	320	11	2009-03-18	00:47:02	2009-03-18	01:12:03	4.8		
2	320	11	2009-03-18	01:40:57	2009-03-18	01:51:53	2.6		
3	320	11	2009-03-18	04:02:20	2009-03-18	04:22:30	2.9		
4	320	11	2009-03-18	06:10:23	2009-03-18	07:24:27	9.8		
5	320	11	2009-03-18	08:51:23	2009-03-18	08:56:11	1.0		
	bout.duration intake.duration								
1	25.0166	7 mins	24	.51667					
2	10.9333	3 mins	10	. 93333					
3	20.1666	7 mins	15	.81667					
4	74.0666	7 mins	63	. 13333					
5	4.8000	0 mins	4	.80000					

We can plot the feed intake distribution during different hours of the day with the help of partOfDay function.

```
> boxplot(intake~partOfDay(begin),data=bouts,ylab='Feed intake (kg)',
+ xlab='Time of day',main='Default settings: start =1, nsplit=4')
```

Default settings: start =1, nsplit=4



4 Summarizing time series

We are frequently interested in summarising events by an hour, day, week or a month. Say how many times a cow has visited the feeding trough during the day or what is the hourly sum of the feed intake for each animal. Animal provides functions to calculate these things easily with functions hourly, daily, weekly and monthly.³

The basic syntax for all functions is similar: We need to specify the data vector we want to analyze, the time stamps for the data in POSIXct format, the summarizing function and optionally the subject.

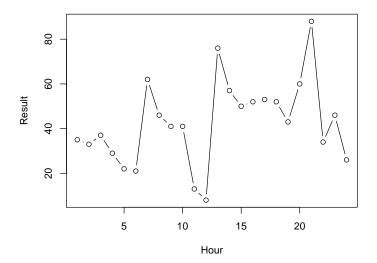
For instance, we are intersted in what is the hourly (summarized according to the start time of feeding event) number of visits in the dataset bouts (created in RIC data chapter).

```
> attach(bouts)
> hourly.visits <- hourly(intake,begin,fun=length)
> head(hourly.visits)
  Hour Result
     1
           35
2
     2
           33
3
     3
           37
4
     4
           29
5
     5
           22
     6
           21
```

> plot(Result~Hour, data=hourly.visits, type='b')

member and more convinient in frequent use

³These are just simple wrapper functions to aggregate, but I feel these are easier to re-



Similarly we can summarize the hourly intake for each cow:

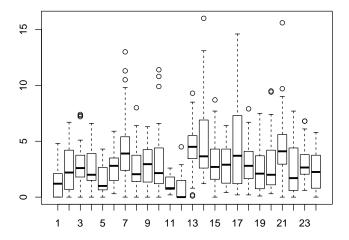
> cow.intake <- hourly(intake,begin,fun=sum,subject=cowID)

> head(cow.intake)

	Hour	Subject	Result
1	1	320	4.8
2	2	320	2.6
3	5	320	2.9
4	7	320	9.8
5	9	320	1.0
6	10	320	3.8

- > boxplot(Result~Hour,data=cow.intake)
- > detach(bouts)

REFERENCES 6



References

[1] Hänninen, L. & Pastell, M. 2009. CowLog: Open source software for coding behaviors from digital video. Behavior Research Methods. 41(2), 472-476.