Updates to R Programming in Foundations and Applications of Statistics

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Since the publication of Foundations and Applications of Statistics, I have been working with colleagues from the NSF-funded Project MOSAIC to create and improve the mosaic package. Many functions originally in the fastR package have been moved to the mosaic package; some of these have subsequently been improved. Additional functionality has been added to the mosaic package over time that I would have used in Foundations and Applications of Statistics, had they existed at the time the book was written. This vignette points out some of these features for students and intructors who might prefer these alternative approaches.

1 Chapter 1: Summarizing Data

1.1 Access to Data

CRAN has requested that we separate the data that were previously in the mosaic package into a separate package (mosaicData) and that this not be a dependency of mosaic of fastR. So some of the data sets used in the book will require loading the new mosaicData package.

```
require(mosaicData)
```

1.2 Taking Advantage of Formulas

One of the big changes in mosaic is the wider support for formula interfaces. Several instances of this approach could be used in Chapter 1. The use of a formula interface has several advantages, the chief among them being a systematization of numerical summaries, graphical summaries, and linear models into a common syntactic template:

```
goal( formula, data=mydata, ...)
```

Common formula shapes include the following

```
goal( ~ x, data=mydata) # for single variable summaries
goal( y ~ x, data=mydata) # for two-variable summaries and linear models
goal( y ~ x | z, data=mydata) # for multi-variable summaries and faceting in plots
```

The function name typically names the goal for the computation (e.g., histogram(), mean(), tally(), etc.). The formula is described using variables in data frame mydata (and removing the need for the \$ operator or with() constructions).

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1.2.1 tally()

The tally() function provides a formula interface for constructing tables.

```
require(fastR)
trellis.par.set(theme=col.mosaic())  # change default colors, etc.
table( iris $ Species )

##
## setosa versicolor virginica
## 50 50 50

tally( ~ Species, data=iris )

##
## setosa versicolor virginica
## 50 50 50
```

By default, tally() adds marginal totals, but these can be turned off, if desired:

```
tally( ~ Species, data=iris, margins=FALSE )

##

## setosa versicolor virginica
## 50 50 50
```

Tallies can be presented as counts, proportions, or percents:

```
tally( ~ Species, data=iris, format="count")
##
##
       setosa versicolor virginica
##
           50
                      50
tally( ~ Species, data=iris, format="percent")
##
##
       setosa versicolor virginica
                   33.33
                              33.33
tally( ~ Species, data=iris, format="proportion")
##
##
       setosa versicolor virginica
       0.3333
                  0.3333
                             0.3333
##
```

The default format is chosen based on the shape of the formula.

1.2.2 Numerical Summaries

The mosaic package provides a formula interface for a number of numerical summary functions.

```
mean( ~ Sepal.Length, data=iris )
## [1] 5.843

median( ~ Sepal.Length, data=iris )
## [1] 5.8

sd( ~ Sepal.Length, data=iris )
## [1] 0.8281

iqr( ~ Sepal.Length, data=iris )
## [1] 1.3

favstats( ~ Sepal.Length, data=iris )
## min Q1 median Q3 max mean sd n missing
## 4.3 5.1 5.8 6.4 7.9 5.843 0.8281 150 0
```

Furthermore, the use of a formula with left and right sides allows us to summarize within groups without using the summary() function:

```
mean (Sepal.Length ~ Species, data=iris )
##
      setosa versicolor virginica
##
       5.006
                  5.936
                             6.588
favstats( Sepal.Length ~ Species, data=iris )
##
                      Q1 median Q3 max mean sd n missing
         .group min
## 1
        setosa 4.3 4.800 5.0 5.2 5.8 5.006 0.3525 50
                                                              0
## 2 versicolor 4.9 5.600
                            5.9 6.3 7.0 5.936 0.5162 50
                                                              0
## 3 virginica 4.9 6.225
                            6.5 6.9 7.9 6.588 0.6359 50
```

Use

?mean

to get a list of additional functions that take advantage of the formula interface.

1.3 Treating data like distributions

In analogy to functions like pnorm() and qnorm(), the mosaic package provides pdata() and qdata().

```
qdata( .5, Sepal.Length, data=iris )

## p quantile
## 0.5 5.8
```

```
median( ~ Sepal.Length, data=iris )

## [1] 5.8

pdata( 5, Sepal.Length, data=iris )

## [1] 0.2133

tally( ~ (Sepal.Length <= 5), data=iris, format="proportion")

##

## TRUE FALSE

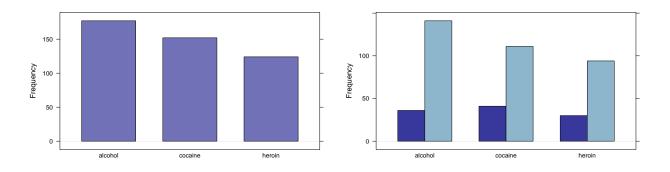
## 0.2133 0.7867</pre>
```

1.4 More plots

1.4.1 bargraph()

The mosaic function barchart() requires the user to first tally the data to be plotted. The bargraph() function makes it easy to create bar graphs in the same way other lattice plots are created.

```
bargraph( ~substance, data=HELPrct)
bargraph( ~substance, data=HELPrct, groups=sex )
```

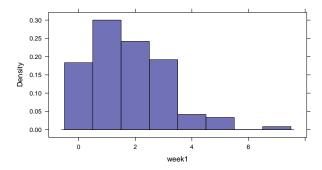


1.4.2 Augmented histogram()

The mosaic package adds several features to the histogram() function (taking advantage of some new features in the lattice package to change the default panel and prepanel functions used). With these changes, xhistogram() has been deprecated and histogram() has all the functionality of xhistogram().

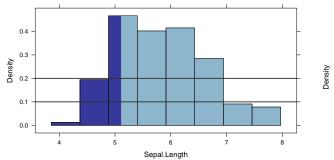
For example, one can choose the bins used for a histogram by setting values for center (defaults to 0) and width. Setting width to 1 is often useful for histograms of integer data with relatively few possible values.

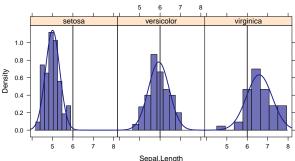
```
histogram( ~ week1, data=fumbles, width=1 )
```



Here are some additional features:

```
histogram( ~ Sepal.Length, data=iris, groups = Sepal.Length > 5, h=c(.1,.2) )
histogram( ~ Sepal.Length | Species, data=iris, fit="normal", v=6 )
```





1.4.3 mPlot()

For RStudio users, the mosaic package provides an interactive interface for creating a wide variety of lattice and ggplot2 graphics using the mPlot() function. The code used to create these plots can subsequently be exported to the console and copied and pasted into other documents. mPlot() requires a data frame and a default plot to produce (scatter plot if none is specified) and allows the user to select variables and several other properties of the plots.

```
mPlot(iris)
mPlot(HELPrct, "density")
```

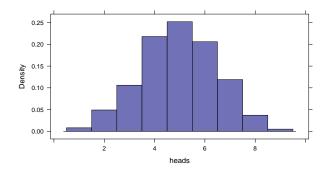
2 Chapter 2: Probability and Random Variables

2.1 The Lady Tasting Tea, rflip(), and do()

For those who want to introduce randomization methods early, the rflip() function provides a natural way to simulate coin tosses, and the do() function does things repeatedly and stores the results in a useful format. For example, the Lady Tasting Tea example can be handled using the following commands.

```
rflip(10)
##
## Flipping 10 coins [ Prob(Heads) = 0.5 ] ...
```

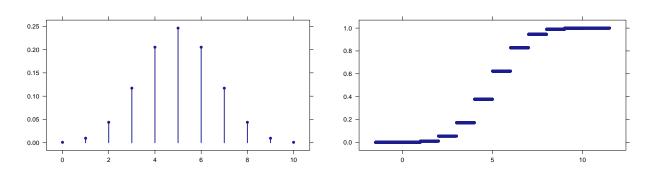
```
##
## H H T T H H T H H T
##
## Number of Heads: 6 [Proportion Heads: 0.6]
do(3) * rflip(10)
## Loading required package: parallel
      n heads tails prop
            4
## 1 10
                  6
                     0.4
## 2 10
            6
                  4
                     0.6
## 3 10
            4
                    0.4
Flips <- do(1000) * rflip(10)
tally( ~heads, data=Flips )
##
##
     1
         2
             3
                 4
                     5
                          6
                                  8
##
     8 49 106 218 252 206 119
                                 37
histogram( ~ heads, data=Flips, width=1 )
```



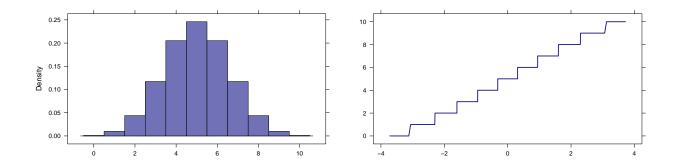
2.2 Plotting Distributions

We can use plotDist() to plot discrete and continuous distributions in a number of ways.

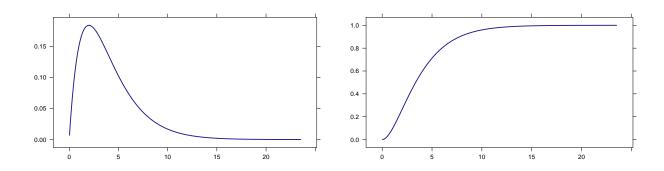
```
plotDist("binom", params=list(size=10,prob=.5))
plotDist("binom", params=list(size=10,prob=.5), kind='cdf')
```



```
plotDist("binom", params=list(size=10,prob=.5), kind='hist')
plotDist("binom", params=list(size=10,prob=.5), kind='qq')
```



```
plotDist("chisq", params=list(df=4))
plotDist("chisq", params=list(df=4), kind='cdf')
```



2.3 Formulas for binom.test()

```
binom.test( ~ sex, data=HELPrct )

##

## Exact binomial test

##

## data: HELPrct$sex

## number of successes = 107, number of trials = 453, p-value <

## 2.2e-16

## alternative hypothesis: true probability of success is not equal to 0.5

## 95 percent confidence interval:

## 0.1978 0.2781

## sample estimates:

## probability of success

## 0.2362</pre>
```

Also, if you only want to extract the p-value or a confidence interval from a hypothesis test object, the pval() and confint() functions will do this for you.

```
pval( binom.test( ~ sex, data=HELPrct ) )
     p.value
##
## 1.932e-30
confint( binom.test( ~ sex, data=HELPrct ) )
## probability of success
                                             lower
                                                                     upper
##
                                            0.1978
                                                                     0.2781
                    0.2362
##
                     level
##
                    0.9500
```

3 Chapter 3: Continuous Distributions

3.1 makeFun()

For functions that are essentially algebraic in nature, the mosaic package provides a simplified method of defining functions via makeFun().

```
f <- makeFun( x^2 ~ x )
f(3)

## [1] 9

g <- makeFun( A*x^2 + B*x + C ~ x, A=1, B=2, C=3 )
g(2)

## [1] 11

g(2, A=3, B=2, C=1)

## [1] 17</pre>
```

3.2 Calculus with D() and antiD()

The mosaic package provides functions for computing derivatives and antiderivates. Each of these functions returns a *function*, which can then be evaluated as needed. This is often easier than working with, for example, integrate() which returns an object from which the value of the integral must be extracted.

```
fprime <- D(f(x) ~ x)
fprime(2)

## [1] 4

fprime

## function (x)

## 2 * (x)

gprime <- D(g(x) ~ x)</pre>
```

```
## Warning: Implicit variables without default values (dangerous!): A, B, C

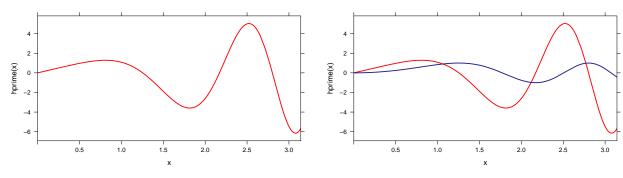
gprime(3)

## [1] 8

gprime(3, A=3, B=2, C=1)

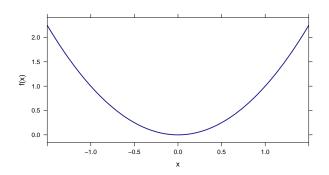
## [1] 20

h <- makeFun( sin(x^2) ~ x )
hprime <- D( h(x) ~ x )
plotFun(hprime(x) ~ x, col="red", x.lim=c(0,pi))
plotFun( h(x) ~ x, x.lim=c(0,pi), add=TRUE )</pre>
```



Antiderivatives work similarly.

```
plotFun(f(x) ~ x, type="h")
F <- antiD( f(x) ~ x )
F(1) - F(0)
## [1] 0.3333</pre>
```



4 Chapter 4: Parameter Estimation and Testing

4.1 t.test()

As was the case for binom.test(), we can now use formulas for the 1-sample t-test:

```
t.test(~age, data=HELPrct)

##

## One Sample t-test

##

## data: data$age

## t = 98.42, df = 452, p-value < 2.2e-16

## alternative hypothesis: true mean is not equal to 0

## 95 percent confidence interval:

## 34.94 36.37

## sample estimates:

## mean of x

## 35.65</pre>
```

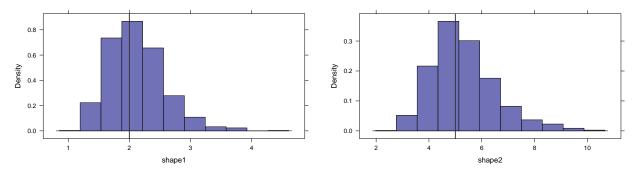
4.2 Simulations with do()

The simulations done using replicate() can be done with do() instead. do() is slower because it does more packaging up of the results, but the format of the data returned is often easier to work with. Here's some code that could replace the code in Example 4.3.3.

```
snippet("mom-beta01") # to define beta.mom
##
##
##
    snippet(mom-beta01)
##
##
## > beta.mom <- function(x,lower=0.01,upper=100) {</pre>
         x.bar <- mean (x)
## +
        n <- length(x)
         v \leftarrow var(x) * (n-1) / n
## +
         R < -1/x.bar - 1
## +
## +
## +
         f <- function(a){</pre>
                                         # note: undefined when a=0
              R * a^2 / ((a/x.bar)^2 * (a/x.bar + 1)) - v
## +
         }
## +
## +
## +
         u <- uniroot(f,c(lower,upper))</pre>
## +
         return( c(shape1=u$root, shape2=u$root * R) )
## +
## + }
##
## > x <- rbeta(50,2,5); beta.mom(x)
## shape1 shape2
## 1.758 4.077
results <- do(1000) * beta.mom( rbeta(50,2,5) )
head(results, 2)
```

```
## shape1 shape2
## 1 1.777 4.620
## 2 2.472 6.097

histogram( ~shape1, data=results, type='density', v=2 )
histogram( ~shape2, data=results, type='density', v=5 )
```



The advantages of using do() are even more pronounced when working with lm(). See the vignettes in the mosaic package for more examples using do().

5 Chapter 5: Likelihood-Based Statistics

5.1 Zermelo's Algorithm

Section 5.6 focuses on the main ideas of the Bradley-Terry model and uses software to do the fitting. But it is not difficult to simplify the (large) system of partial differential equations involved in the maximum likelihood estimation into a form that leads to both a natural characterization of the MLE and an iterative algorithm for approximating the MLE that go back to Zermelo.

6 Chapter 6: Introduction to Linear Models

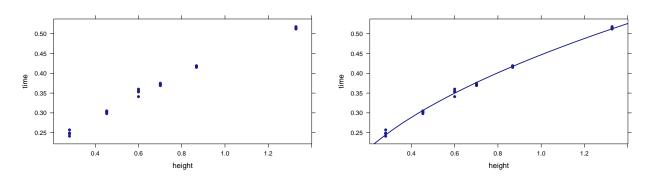
6.1 Converting models to functions with makeFun()

makeFun() can convert models made with lm() and glm() into functions. In both cases the functions produced is a wrapper around predict(). These functions take care of any transformations of the explanatory variables but not transformations of the response variable. In the case of glm() models, the default type is "response" rather than "link" since this is more natural for beginners.

6.2 And adding fitted functions to plots with plotFun()

We can add the model fit function to our scatter plot using plotFun().

```
xyplot( time ~ height, data=balldrop )
plotFun( time(height) ~ height, add=TRUE )
```



7 Chapter 7: More Linear Models

7.1 TukeyHSD() no longer requires use of aov()

```
# TukeyHSD() can take a model created by lm()
model <- lm( pollution ~ location, data=airpollution)
TukeyHSD(model)
     Tukey multiple comparisons of means
##
##
       95% family-wise confidence level
## Fit: aov(formula = x)
##
## $location
##
                             diff
                                     lwr
                                            upr p adj
## Plains Suburb-Hill Suburb
                               -6 -40.29 28.29 0.7643
## Urban City-Hill Suburb
                               15 -19.29 49.29 0.3019
## Urban City-Plains Suburb
                               21 -13.29 55.29 0.1601
# we can even let TukeyHSD build the model for us
TukeyHSD( pollution ~ location, data=airpollution)
##
     Tukey multiple comparisons of means
##
       95% family-wise confidence level
## Fit: aov(formula = x)
## $location
##
                             diff
                                     lwr
                                            upr p adj
## Plains Suburb-Hill Suburb
                               -6 -40.29 28.29 0.7643
## Urban City-Hill Suburb
                               15 -19.29 49.29 0.3019
## Urban City-Plains Suburb
                               21 -13.29 55.29 0.1601
```