

## R Introduction/Review Assignment

The `RuffeBio.csv` file contains a variety of biological measurements from Ruffe (*Gymnocephalus cernuus*) captured in the St. Louis River Harbor. Use these data to answer the following questions.

```
> setwd("C:/aaaWork/Web/fishR/Courses/MNAFS2013/CourseMaterial")
> ruf <- read.csv("RuffeBio.csv",header=TRUE)
> str(ruf)

'data.frame': 40 obs. of 10 variables:
 $ fishID : int 60 61 62 63 64 65 66 67 68 69 ...
 $ locShort: Factor w/ 1 level "St. Louis R. (2007)": 1 1 1 1 1 1 1 1 1 1 ...
 $ year : int 2007 2007 2007 2007 2007 2007 2007 2007 2007 2007 ...
 $ month : int 9 9 9 9 9 9 9 9 9 9 ...
 $ day : int 20 20 20 20 20 20 20 20 20 20 ...
 $ date : Factor w/ 1 level "9/20/2007": 1 1 1 1 1 1 1 1 1 1 ...
 $ tl : int 134 111 110 115 92 88 95 90 99 107 ...
 $ wt : num 24.6 14.7 12.3 16 8.3 7.8 9.7 8.2 11.7 13 ...
 $ sex : Factor w/ 3 levels "female","male",...: 1 1 1 1 1 1 1 1 1 1 ...
 $ maturity: Factor w/ 3 levels "", "immature",...: 3 3 2 3 3 3 3 3 3 3 ...
```

1. How many variables are in this data frame?  
There are 10 variables in this data frame.
2. Data was recorded for how many Ruffe?  
Data was recorded on 40 Ruffe in this data frame.
3. Show the vector of `tl` measures?

```
> ruf$tl
 [1] 134 111 110 115 92 88 95 90 99 107 NA 99 102 105 90 102 114 NA 56 90
[21] 101 109 110 111 101 95 84 105 120 104 102 99 84 87 81 81 65 42 NA 115
```

All of the total length measurements are shown above.

4. What is the `tl` for the 17th Ruffe?

```
> ruf$tl[17]
[1] 114
```

The `tl` for the 17th measured individual is 114 mm.

5. Show all information for the 20th Ruffe.

```
> ruf[20,]
  fishID      locShort year month day      date tl  wt  sex maturity
20    79 St. Louis R. (2007) 2007    9  20 9/20/2007 90 7.6 female  mature
```

All of the information for the 20th Ruffe is shown above.

6. Create a new data frame of just female Ruffe. How many fish are in this data frame?

```
> df1 <- Subset(ruf,sex=="female")
```

There are 31 Ruffe in this data frame.

7. Create a new data frame excluding Ruffe of unknown sex. How many fish are in this data frame?

```
> df2 <- Subset(ruf,sex!="unknown")
```

There are 39 Ruffe in this data frame.

8. Create a new data frame of Ruffe greater than 110 mm. How many fish are in this data frame?

```
> df3 <- Subset(ruf,tl>110)
```

There are 7 Ruffe in this data frame.

9. Create a new data frame of Ruffe between 80 and 120 mm. How many fish are in this data frame?

```
> df4 <- Subset(ruf,tl>80 & tl<120)
```

There are 32 Ruffe in this data frame.

10. Create a new data frame of male Ruffe less than 100 mm. How many fish are in this data frame?

```
> df5 <- Subset(ruf,sex=="male" & tl<100)
```

There are 5 Ruffe in this data frame.

11. Create a new variable in the original data frame that is Fulton's condition factor (i.e., the weight of the fish divided by the cubed length of the fish multiplied by 100000).

```
> ruf$fult <- ruf$wt/(ruf$tl^3)*100000
```

12. Compute the mean Fulton's condition factor.

```
> Summarize(~fult,data=ruf,digits=3)
```

	n	mean	sd	min	Q1	median	Q3	max	percZero
	37.000	1.061	0.105	0.837	1.010	1.070	1.120	1.310	0.000

The mean Fulton's condition factor is 1.061 Ruffe in this data frame.

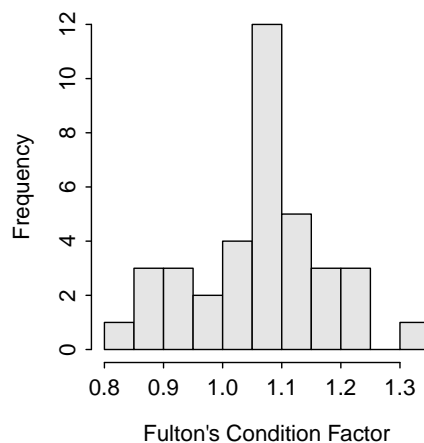
13. Compute the mean Fulton's condition factor separately for males and females.

```
> Summarize(fult~sex,data=ruf,digits=3)
```

	sex	n	mean	sd	min	Q1	median	Q3	max	percZero
1	female	30	1.070	0.109	0.837	1.040	1.070	1.13	1.31	0
2	male	6	1.017	0.088	0.938	0.949	0.986	1.07	1.15	0
3	unknown	1	1.080	NA	1.080	1.080	1.080	1.08	1.08	0

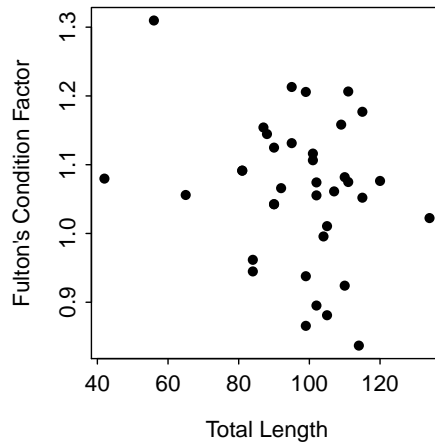
14. Create a histogram of Fulton's condition factor.

```
> hist(~fult,data=ruf,xlab="Fulton's Condition Factor",col="gray90")
```



15. Create a scatterplot of the relationship between Fulton's condition factor and total length.

```
> plot(fult~tl,data=ruf,xlab="Total Length",ylab="Fulton's Condition Factor",pch=16)
```



16. Fit a model that can be used to predict Fulton's condition factor from total length. Find the model coefficients and  $r^2$  value and construct a plot that shows the fitted model.

```
> lm1 <- lm(fult~tl,data=ruf)
> summary(lm1)
```

Call:  
lm(formula = fult ~ tl, data = ruf)

Residuals:

Min	1Q	Median	3Q	Max
-0.2005	-0.0559	0.0134	0.0603	0.1899

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	1.199293	0.097239	12.33	2.7e-14
tl	-0.001420	0.000985	-1.44	0.16

Residual standard error: 0.104 on 35 degrees of freedom  
(3 observations deleted due to missingness)  
Multiple R-squared: 0.056, Adjusted R-squared: 0.029  
F-statistic: 2.08 on 1 and 35 DF, p-value: 0.158

```
> plot(fult~tl,data=ruf,xlab="Total Length",ylab="Fulton's Condition Factor",pch=16)
> b <- coef(lm1)[1]
> m <- coef(lm1)[2]
> curve(m*x+b,from=40,to=135,add=TRUE,col="red",lwd=2)
```

