

Discussion 1

Randy Johnson

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BIFX 553 Discussion 1: January 19, 2017

Pretest

- 1) What is a p-value?
- 2) Write a short R script to add all numbers from 1 to 100 using a for loop.

```
s <- 0
for(i in 1:100)
  s <- s + i
```

- 3) Repeat problem 2 using only one line of code (i.e. without using a for loop).

```
sum(1:100)
```

```
## [1] 5050
```

- 4) Given the data.frame, *dat*, with the variables $\{x1, x2, x3, y\}$, write the R command to create a linear model for *y* using *x1*, *x2* and *x3* as predictors.

```
lm(y ~ x1 + x2 + x3, data = dat)
```

- 5) Write an R command to plot the relationship between *x1* and *y*.

```
plot(x1, y, bty = 'l')
```

- 6) Given the relationship between *x1* and *y* in this figure, update the R command in question 4 with a more appropriate model.

```
dat$x1sqr <- dat$x1^2 lm(y ~ x1 + x1sqr + x2 + x3)
```

- 7) A collaborator presents a data analysis to you. The p-value they give you is $p = 0.012$. Is this statistically significant?

Yes

- 8) After discussing the results further, you discover that this p-value represents just one of 20 tests performed in the analysis. What is the Bonferroni threshold for significance if you want the family-wise error rate to be $\alpha = 0.05$? Is the result statistically significant?

```
(thresh <- 0.05 / 20)
```

```
## [1] 0.0025
```

No

- 9) What assumptions are made in a simple linear regression? What R command(s) can you use to get a quick look at whether those assumptions have been violated?

- Linear relationship
- Multivariate Normality
- No/little multicollinearity
- No autocorrelation
- Homoscedasticity

- General lm diagnostics
 - plot()
 - broom package
 - car package
 - gvlma package
- 10) Explain the relationship between confidence intervals and p-values. Why are confidence intervals more useful than p-values?

Discussion of R programming basics

- What data types does R have?
 - numeric
 - integer
 - double
 - character
 - factor
- What are the standard data structures in R? How are they indexed?
 - scalar
 - vector
 - matrix
 - array
 - lists
 - data.frames
 - (tibble)

Lecture notes are located in the course repository

- Examples of vector calculations

```
x <- 1:10
y <- rnorm(10)
x + y

## [1] 1.714201 2.473387 3.287186 3.739490 4.599493 5.885009 7.878700
## [8] 8.414016 9.447221 9.516530

mat <- cbind(x, y)
apply(mat, 1, sum)

## [1] 1.714201 2.473387 3.287186 3.739490 4.599493 5.885009 7.878700
## [8] 8.414016 9.447221 9.516530

lst <- list(rnorm(100),
            rnorm(100, 5),
            rnorm(100, 5, 10))
sapply(lst, mean)

## [1] 0.197392 4.993251 5.322037

lapply(lst, mean)

## [[1]]
## [1] 0.197392
##
## [[2]]
```

```
## [1] 4.993251
##
## [[3]]
## [1] 5.322037
```

```
ifelse(y > 0, 1, -1)
```

```
## [1] 1 1 1 1 -1 -1 -1 1 1 1 -1
```

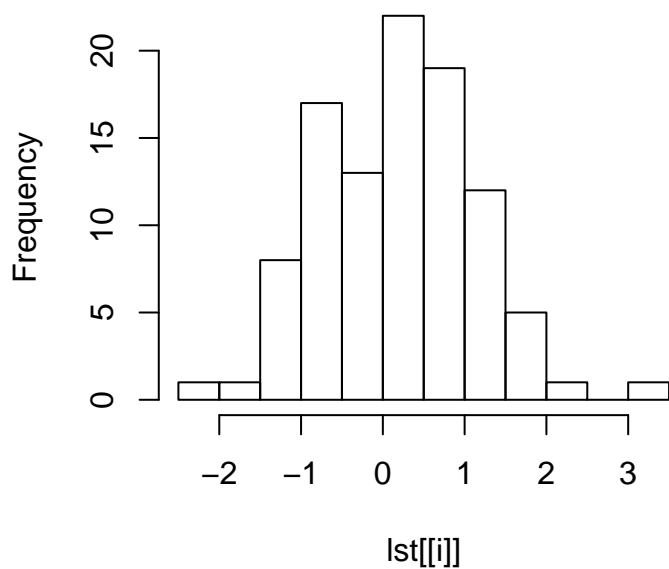
- Examples of control structures

```
test <- TRUE
if(test)
{
  print("This is executed when test is true.")
}else{
  print("This is executed when test is false.")
}
```

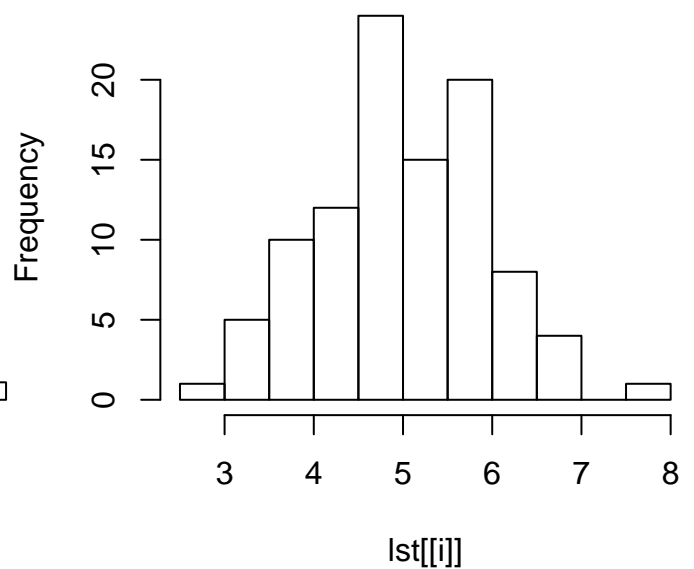
```
## [1] "This is executed when test is true."
```

```
for(i in 1:2)
{
  hist(lst[[i]])
}
```

Histogram of lst[[i]]



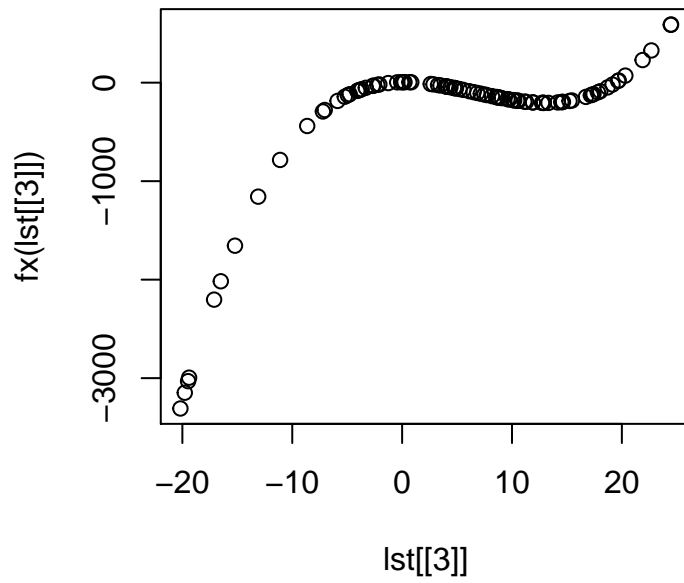
Histogram of lst[[i]]



- There are other control structures, but you shouldn't need them very often (if ever).
- Functions

```
fx <- function(x)
{
  # do something here and return a value
  return(5 + 2*x - 4*x^2 + x^3/5)
}
```

```
plot(lst[[3]], fx(lst[[3]]))
```



Questions?