DSSH 6301 - HW 02 Solutions

Here are suggested solutions to the homework assignment. Note that many problems have multiple solutions, so we present here merely one example. If you did not have points taken off for a question, we considered your solution correct. If you have questions about any of these solutions or your own graded homework, please direct questions to Matt Harrigan m.harrigan@neu.edu .

Problem 1

Part a

Note how (1) simple functions and if-then statements can be defined without the explicit use of brackets; (2) the function eliminates NA elements; and (3) it returns "NaN" (Not a Number) if the input is non-numeric. Note also how how all functions return the last quantity without needing an explicit use of "return".

```
mean_fn <- function(x) if (is.numeric(x)) sum(x, na.rm=T)/length(x[!is.na(x)]) else NaN
mean_fn(1:4)

## [1] 2.5
mean_fn(c("cat", "dog", "frog"))</pre>
```

[1] NaN

A simpler version of the function without the error handling, and using brackets and an explicit "return" might be:

```
mean_fn <- function(x){
  mn <- sum(x)/length(x)
  return(mn)
}</pre>
```

Part b

```
fn <- function(x) if(sum(x[1:2]) > sum(x[3:4])) x else 0
fn(1:4)

## [1] 0

fn(4:1)

## [1] 4 3 2 1
```

Part c

The two examples given here show the calculation of the Fibonacci sequence by appending a single element to the end of a vector.

```
fib_seq_concat1 <- function(n) {</pre>
  if (n < 1)
    stop("n must be a positive integer") # Stops on invalid input
  if (n == 1)
    return(1)
  if (n == 2)
    return(c(1, 1))
  fib <- c(1, 1)
  for (i in 3:n)
    fib <- c(fib, fib[i-2] + fib[i-1])
  return(fib)
fib_seq_concat2 <- function(n) {</pre>
  if (n < 1)
    stop("n must be a positive integer") # Stops on invalid input
  if (n == 1)
    return(1)
  if (n == 2)
    return(c(1, 1))
  fib <- c(1, 1)
  for (i in 3:n)
    fib[i] \leftarrow fib[i-2] + fib[i-1]
  return(fib)
fib_seq_concat1(10)
```

```
## [1] 1 1 2 3 5 8 13 21 34 55
```

```
fib_seq_concat2(10)
```

```
## [1] 1 1 2 3 5 8 13 21 34 55
```

Part d

```
m1 <- matrix(1:16, nrow=4, ncol=4)
apply(m1, 1, mean_fn)
## [1] 7 8 9 10</pre>
```

Problem 2

Part a

Only measurements where both Wind and Ozone are not NA are aggregated when using the aggregate function like this. Thus the discrepancy in means for month 5 that you see below.

```
aggregate(cbind(Wind, Ozone) ~ Month, data=airquality, mean)
##
    Month
               Wind
                       Ozone
## 1
       5 11.457692 23.61538
       6 12.177778 29.44444
       7 8.523077 59.11538
## 3
## 4
       8 8.565385 59.96154
## 5
     9 10.075862 31.44828
mean(airquality$Wind[airquality$Month == 5])
## [1] 11.62258
mean(airquality$Wind[airquality$Month == 5 & !is.na(airquality$Ozone)])
## [1] 11.45769
```

Part b

```
##
      surname nationality
                                                  title
## 1
      McNeil
              Australia
                              Interactive Data Analysis
## 2
      Ripley
                                     Spatial Statistics
                       UK
                                  Stochastic Simulation
## 3
      Ripley
## 4
     Tierney
                       US
                                              LISP-STAT
## 5
                       US
                              Exploratory Data Analysis
       Tukey
## 6 Venables
               Australia Modern Applied Statistics ...
       R Core
                     <NA>
                                   An Introduction to R
## 7
```

Part c

Note that using cat rather than print preserves the carriage returns in the output.

```
str <- "To be, or not to be- that is the question:
Whether 'tis nobler in the mind to suffer
The slings and arrows of outrageous fortune
Or to take arms against a sea of troubles,
And by opposing end them. To die - to sleep -
No more..."

str <- gsub("to", 2, str, ignore.case = T)
cat(str)

## 2 be, or not 2 be- that is the question:
## Whether 'tis nobler in the mind 2 suffer
## The slings and arrows of outrageous fortune
## Or 2 take arms against a sea of troubles,
## And by opposing end them. 2 die - 2 sleep -</pre>
```

Problem 3

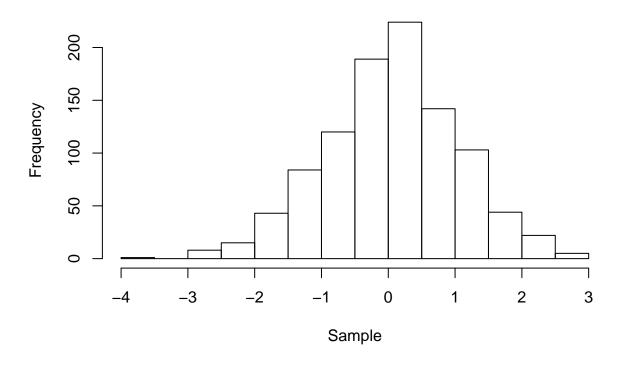
No more...

Part a

We create 1000 samples from a normal distribution here. More information on this function and its uses will be given later in the course.

```
data <- rnorm(1000)
hist(data, main="Standard Normal Distribution", xlab="Sample")</pre>
```

Standard Normal Distribution



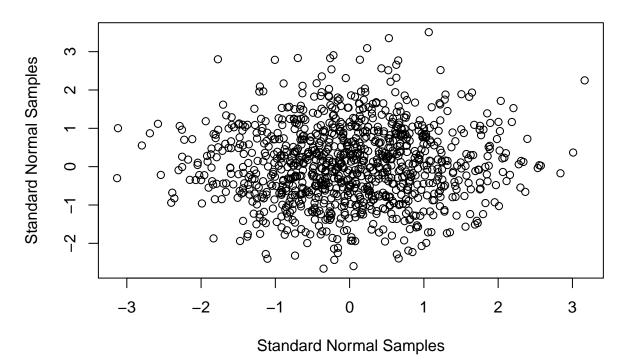
Part b

```
x <- rnorm(1000)
y <- rnorm(1000)

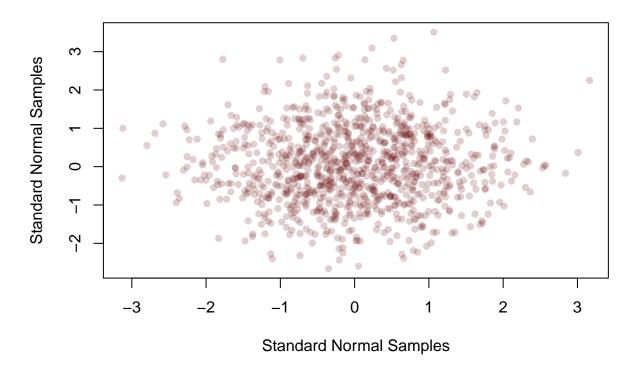
xlab <- "Standard Normal Samples"
ylab <- "Standard Normal Samples"

plot(x, y, main="Two Independent Gaussians", xlab=xlab, ylab=ylab)</pre>
```

Two Independent Gaussians



Two Independent Gaussians



Part c

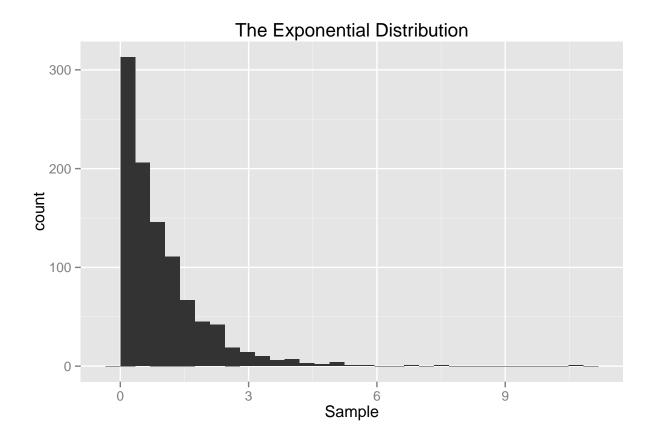
```
require(ggplot2)

## Loading required package: ggplot2

data <- data.frame(Sample=rexp(1000))

main <- "The Exponential Distribution"

ggplot(data, aes(x=Sample)) + geom_histogram() + ggtitle(main)</pre>
```

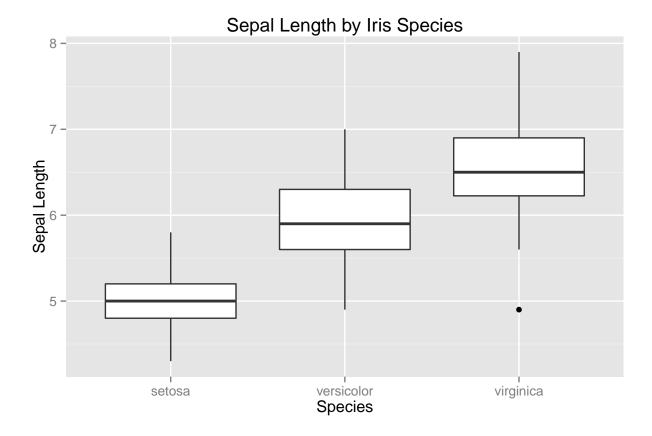


Part d

Be sure that if you choose to use a dataset that's built into a package (such as ggplot2), you must be sure to load the package earlier in your .Rmd file. In this case iris is part of the base R and is always available.

```
main <- "Sepal Length by Iris Species"

ggplot(iris, aes(x=Species, y=Sepal.Length)) + geom_boxplot() +
    ggtitle(main) + ylab("Sepal Length")</pre>
```



Part e

```
require(car)

## Loading required package: car

main <- "1992 US Statistical Abstract, Census Bureau"
xlab <- "Mean State SAT Verbal Scores of Graduating HS Students"
ylab <- "Mean State SAT Math Scores of Graduating HS Students"

ggplot(States, aes(x=SATV, y=SATM)) + geom_point() + ggtitle(main) +
    xlab(xlab) + ylab(ylab)</pre>
```

