# "COMPSCIX 415.2 Homework 6"

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## Github location

My homework assignments can be found at https://github.com/santumagic/compscix-415-2assignments.git

## Exercise 1

#### Question: 1

```
# Load the required packages
library(tidyverse)
library(mdsr)
library(mosaicData)
# glimpse the given dataset
glimpse(Whickham)

## Observations: 1,314
## Variables: 3
## $ outcome <fct> Alive, Alive, Dead, Alive, Alive, Alive, Alive, Dead, ...
## $ smoker <fct> Yes, Yes, Yes, No, No, Yes, Yes, No, No, No, Yes, ...
## $ age <int> 23, 18, 71, 67, 64, 38, 45, 76, 28, 27, 28, 34, 20, 72...
```

#### Answer:

Below are the three variables from the Whickham dataset.

- outcome
- smoker
- age

### Question: 2

#### Answer:

There are 1314 observations. Each observation represents a person. The data set indicates if the individual is a smoker or not, current age an individual, and if the individual is alive or dead.

## Question: 3

```
library(mosaicData)
library(tidyverse)
Whickham %>% count( smoker , outcome )
## # A tibble: 4 x 3
##
     smoker outcome
                        n
##
     <fct> <fct>
                    <int>
## 1 No
            Alive
                      502
## 2 No
            Dead
                       230
## 3 Yes
                      443
            Alive
## 4 Yes
            Dead
                      139
```

#### ${f Answer:}$

By looking at the above table, it is difficult to conclude anything, so I calculate the proportions first by using the below code.

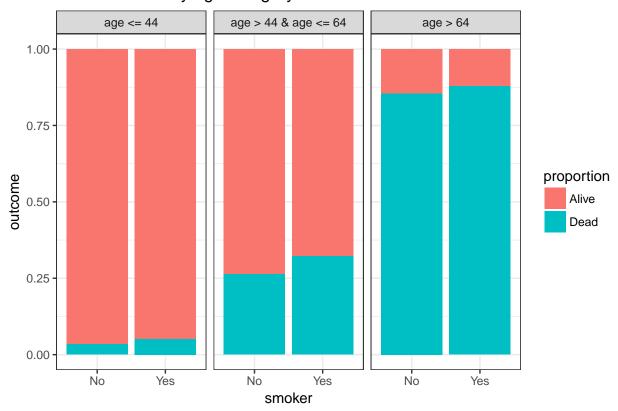
```
Whickham_proportions <- Whickham %>% group_by(smoker,outcome) %>%
  summarize(n = n()) \%
 mutate (prop = n/sum(n))
Whickham_proportions
## # A tibble: 4 x 4
## # Groups:
              smoker [2]
##
     smoker outcome
                        n prop
##
     <fct> <fct> <int> <dbl>
## 1 No
           Alive
                     502 0.686
## 2 No
           Dead
                     230 0.314
## 3 Yes
           Alive
                      443 0.761
## 4 Yes
                     139 0.239
           Dead
```

By looking at the above proportions it is observed that, 31.4 % of non smokers are dead and 23.8 % of smokers are dead which means more healthy people are dead. So there must be other reasons for deaths or data might be wrong. So the data doesn't make any sense.

#### Question: 4

```
# creating the age groups column
Whickham_factor <- Whickham %>% mutate (category =
                                          factor (
case_when(age <= 44 ~ "age <= 44", age > 44 & age <= 64 ~ "age > 44 & age <= 64",
          age > 64 ~ "age > 64")))
head(Whickham factor) # display the top rows of the result dataset
##
     outcome smoker age
                                    category
## 1
      Alive
               Yes 23
                                   age <= 44
## 2
      Alive
               Yes 18
                                   age <= 44
## 3
       Dead
               Yes 71
                                    age > 64
## 4
      Alive
                No 67
                                    age > 64
## 5
      Alive
                No 64 age > 44 & age <= 64
## 6
      Alive
               Yes 38
                                   age <= 44
# reorganize the data by grouping, summarise the data and finding the proportions
Whickham_cat <- Whickham_factor %>%
group_by(category,smoker,outcome) %>% summarise( n = n()) %>%
mutate (proportion = n/sum(n))
head(Whickham_cat) # display the top rows of the result dataset
## # A tibble: 6 x 5
## # Groups: category, smoker [3]
##
                         smoker outcome
                                             n proportion
     category
     <fct>
                          <fct> <fct>
                                         <int>
##
                                                    <dbl>
## 1 age <= 44
                                 Alive
                                           327
                                                   0.965
                          No
## 2 age <= 44
                         No
                                 Dead
                                           12
                                                   0.0354
## 3 age <= 44
                          Yes
                                 Alive
                                           270
                                                   0.947
## 4 age <= 44
                          Yes
                                 Dead
                                           15
                                                   0.0526
## 5 age > 44 & age <= 64 No
                                           147
                                                   0.735
                                 Alive
## 6 age > 44 & age <= 64 No
                                                   0.265
                                 Dead
                                            53
```

# Smoker status by age category



#### Answer:

From the above dataset it is observed that untill the age of 44, non-smokers have only a 1% advantage compared to smokers, but this gap increases dramatically between the ages 44 & 64 where 6% more non-smokers are alive than smokers. Beyond 64 year of age, the difference drops to 3% with non-smokers still being alive more often than smokers.

## Exercise 2

### Question: 1

```
# given sample code
library(tidyverse)
n <- 10000
# look at ?rgamma to read about this function
gamma_samp <- tibble(x = rgamma(n, shape = 1, scale = 2))
gamma_samp</pre>
```

```
## # A tibble: 10,000 x 1
##
         х
##
      <dbl>
##
   1 9.89
   2 1.31
##
##
  3 1.16
## 4 2.22
##
  5 1.51
##
  6 1.59
## 7 0.169
## 8 1.76
## 9 0.463
## 10 4.09
## # ... with 9,990 more rows
\# histogram for the above sample gamma
ggplot(data = gamma_samp) +
  geom_histogram(aes(x=x), bins=100) +
  theme_bw()
   900
   600
   300
```

# Question: 2

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5

```
# finding mean
sprintf ("Mean = %f", sapply(gamma_samp, mean, na.rm = TRUE))
```

Х

15

20

25

10

```
## [1] "Mean = 1.988671"
# finding standard deviation
sprintf ("Standard deviation = %f", sapply(gamma_samp, sd, na.rm = TRUE))
## [1] "Standard deviation = 1.994718"

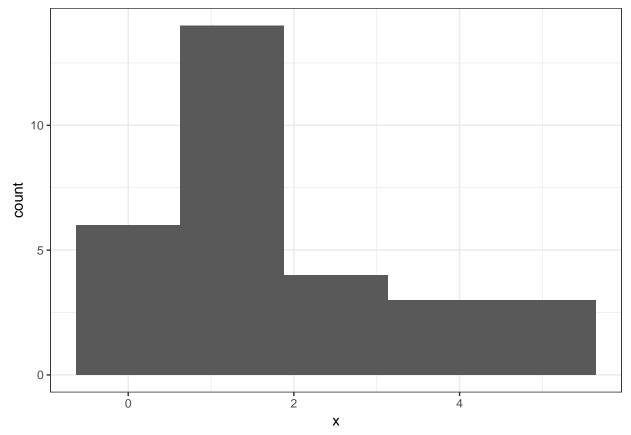
Question: 3
```

```
# sample of size n = 30
sample_30 <- gamma_samp %>% sample_n(30, replace = TRUE)
# finding mean
sprintf("Mean = %f", sapply(sample_30, mean, na.rm = TRUE))
```

```
## [1] "Mean = 1.680049"
# finding standard deviation
sprintf ("Standard deviation = %f", sapply(sample_30, sd, na.rm = TRUE))
```

## [1] "Standard deviation = 1.495609"

```
# Plot the histogram
ggplot(data =sample_30, mapping = aes(x=x)) +
  geom_histogram(bins=5) +
  theme_bw()
```



Question: 4

Question: 5

Question: 6

Question: 7

Question: 8